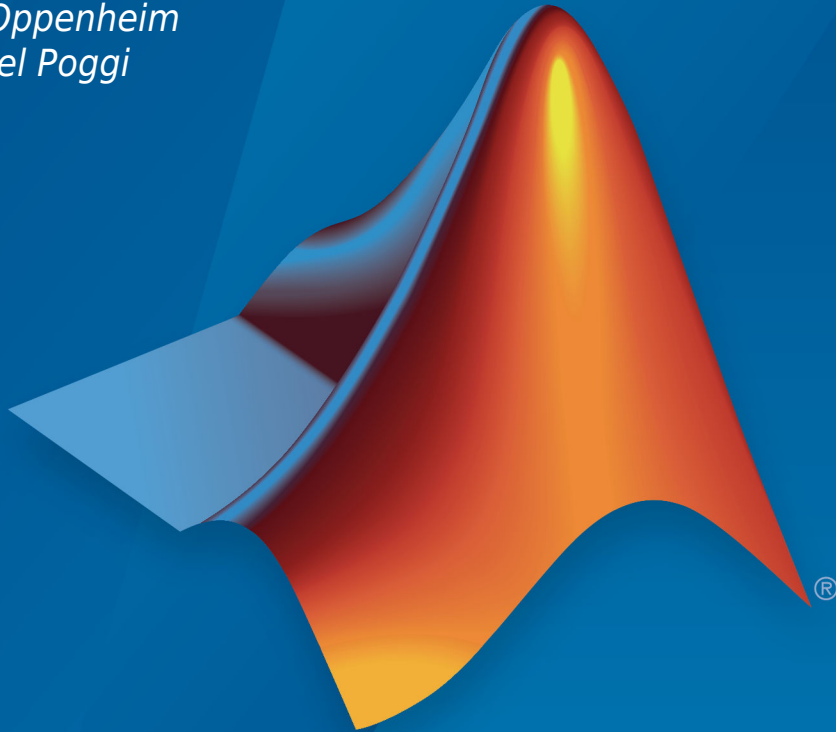


Wavelet Toolbox™

Reference

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Wavelet Toolbox™ Reference

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March 2019	Online only	Revised for Version 5.2 (Release 2019a)

1	<u>Function Reference</u>
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Function Reference

addLabelDefinitions

Add label definitions to labeled signal set

Syntax

```
addLabelDefinitions(lss, lbldefs)
addLabelDefinitions(lss, lbldefs, lblname)
```

Description

`addLabelDefinitions(lss, lbldefs)` adds the labels defined in the vector of signal label definitions `lbldefs` to the labeled signal set `lss`.

`addLabelDefinitions(lss, lbldefs, lblname)` adds the labels defined in `lbldefs` as sublabels of the label `lblname`.

Examples

Add Label Definition

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss

lss =
    labeledSignalSet with properties:
        Source: {2x1 cell}
        NumMembers: 2
        TimeInformation: "sampleRate"
        SampleRate: 4000
        Labels: [2x3 table]
        Description: "Characterize wave song regions"
```


Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Create a label definition that specifies whether a signal corresponds to a calf or to an adult whale.

```
calf = signalLabelDefinition('Calf','LabelDataType','logical','DefaultValue',false, ..
    'Description','Is the specimen a calf, or an adult?')
```

```
calf =
    signalLabelDefinition with properties:
        Name: "Calf"
        LabelType: "attribute"
        LabelDataType: "logical"
        ValidationFunction: []
        DefaultValue: 0
        Sublabels: [0x0 signalLabelDefinition]
        Tag: ""
        Description: "Is the specimen a calf, or an adult?"
```

Use `labeledSignalSet` to create a labeled signal set.

Add the definition to the labeled signal set. Retrieve the names of the labels.

```
addLabelDefinitions(lss,calf)
```

```
getLabelNames(lss)
```

```
ans = 4x1 string array
    "WhaleType"
    "MoanRegions"
    "TrillRegions"
    "Calf"
```

Create a label definition that specifies the sex of the whale. Add the label to the set as a sublabel of 'WhaleType'.

```
sx = signalLabelDefinition('Sex','LabelDataType','categorical', ...
    'Categories',["male" "female"]);
addLabelDefinitions(lss,sx,'WhaleType')
```

```
labelDefinitionsHierarchy(lss)
```

```
ans =  
    'WhaleType  
      Sublabels: Sex  
    MoanRegions  
      Sublabels: []  
    TrillRegions  
      Sublabels: TrillPeaks  
    Calf  
      Sublabels: []  
,
```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: labeledSignalSet({randn(100,1)
randn(10,1)}, signalLabelDefinition('female')) specifies a two-member set of random signals containing the attribute 'female'.

lbldefs — Signal label definitions

signalLabelDefinition object | vector of signalLabelDefinition objects

Signal label definitions, specified as a signalLabelDefinition object or a vector of signalLabelDefinition objects.

Example:

signalLabelDefinition("Asleep", 'LabelType', 'roi', 'LabelDataType', 'logical') can label a region of a signal in which a patient is asleep.

lblname — Label name

character vector | string scalar

Label name, specified as a character vector or a string scalar.

Example: signalLabelDefinition("Asleep", 'LabelType', 'roi') specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

addlift

Add lifting steps to lifting scheme

Syntax

```
LSN = addlift(LS,ELS)  
LSN = addlift(LS,ELS, 'begin')  
LSN = addlift(LS,ELS, 'end')  
addfilt(LS,ELS)
```

Description

LSN = addlift(*LS*,*ELS*) returns the new lifting scheme *LSN* obtained by appending the elementary lifting step *ELS* to the lifting scheme *LS*.

LSN = addlift(*LS*,*ELS*, 'begin') prepends the specified elementary lifting step.

ELS is either a cell array (see `lsinfo`)

```
{TYPEVAL, COEFS, MAX_DEG}
```

or a structure (see `liftfilt`)

```
struct('type',TYPEVAL,'value',LPVAL)
```

with

```
LPVAL = laurpoly(COEFS, MAX_DEG)
```

LSN = addlift(*LS*,*ELS*, 'end') is equivalent to `addfilt(LS,ELS)`.

If *ELS* is a sequence of elementary lifting steps, stored in a cell array or an array of structures, then each of the elementary lifting steps is added to *LS*.

For more information about lifting schemes, see `lsinfo`.

Examples

Add Primal Lifting Step

This example shows how to start with the Haar lifting scheme and add a primal lifting step.

```
LSbegin = liftwave('haar');
```

Display the lifting scheme.

```
displs(LSbegin);
```

```
LSbegin = {...
'd'          [ -1.00000000] [0]
'p'          [  0.50000000] [0]
[  1.41421356] [  0.70710678] []
};
```

Create a primal lifting step.

```
pstep = { 'p', [-1 2 -1]/4 , 1 };
```

Add the primal lifting step.

```
LSend = addlift(LSbegin,pstep);
```

Display the final lifting scheme.

```
displs(LSend);
```

```
LSend = {...
'd'          [ -1.00000000] [0]
'p'          [  0.50000000] [0]
'p'          [ -0.25000000  0.50000000 -0.25000000] [1]
[  1.41421356] [  0.70710678] []
};
```

See Also

liftfilt

Introduced before R2006a

addMembers

Add members to labeled signal set

Syntax

```
addMembers(lss,src)
addMembers(lss,src,tinfo)
addMembers(lss,src,tinfo,mnames)
```

Description

`addMembers(lss,src)` adds members to the labeled signal set `lss` from the input data source `src`.

`addMembers(lss,src,tinfo)` sets the time information for the new members to `tinfo`.

`addMembers(lss,src,tinfo,mnames)` sets the names of the new members to `mnames`. The length of `mnames` must be equal to the number of new members.

Examples

Add Member to Labeled Signal Set

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss

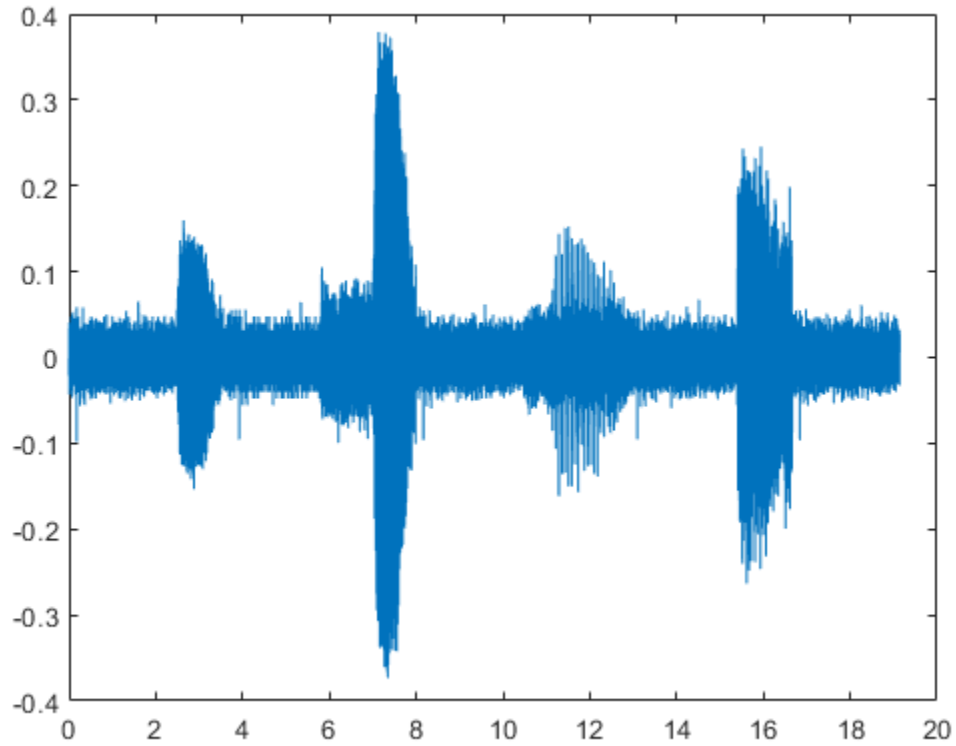
lss =
  labeledSignalSet with properties:
      Source: {2x1 cell}
      NumMembers: 2
```

```
TimeInformation: "sampleRate"  
  SampleRate: 4000  
  Labels: [2x3 table]  
  Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

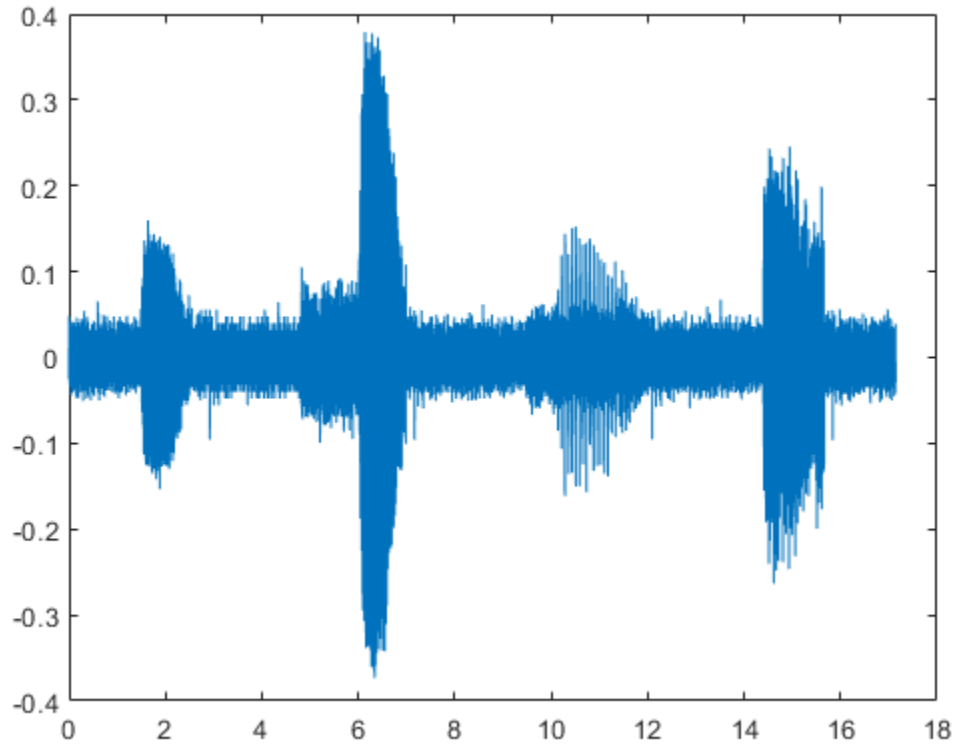
Retrieve the second member of the set and plot it.

```
[song,tinfo] = getSignal(lss,2);  
t = (0:length(song)-1)/tinfo.SampleRate;  
plot(t,song)
```

Remove the first and last seconds of the retrieved signal.

```
song2 = song(t>1 & t<t(end)-1);  
t2 = (0:length(song2)-1)/tinfo.SampleRate;  
  
plot(t2,song2)
```



Add the shorter signal as a new member of the labeled set.

```
addMembers(lss, song2)  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {3x1 cell}  
      NumMembers: 3  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [3x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Flip the shorter signal upside-down and add it as a new member of the labeled set.
Specify that the new member is sampled at 1 kHz.

```
addMembers(lss, flipud(song2), 1000)
lss.SampleRate
```

```
ans = 4×1
      4000
      4000
      4000
      1000
```

Input Arguments

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1), randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

src — Input data source

`matrix` | `timetable` | `audio datastore`

Input data source, specified as a matrix, a timetable, or an audio datastore. The particular form of `src` depends on the “Source” on page 1-0 property of `lss`.

- If “Source” on page 1-0 is a cell array of matrices:
 - Specify `src` as a matrix to add one member to the set.
 - Specify `src` as a cell array of matrices to add multiple members to the set.

- If “Source” on page 1-0 is a cell array containing cell arrays of vectors:
 - Specify `src` as a cell array of vectors to add one member to the set.
 - Specify `src` as a cell array containing cell arrays of vectors to add multiple members to the set.
- If “Source” on page 1-0 is a cell array of timetables:
 - Specify `src` as a timetable to add one member to the set.
 - Specify `src` as a cell array of timetables to add multiple members to the set.
- If “Source” on page 1-0 is an audio datastore, then add members by setting `src` as another audio datastore that points to new files.

Example: `{randn(10,3),randn(17,9)}` specifies two members. The first member contains three 10-sample signals. The second member contains nine 17-sample signals.

Example: `{{randn(10,1)},{randn(17,1),randn(27,1)}}` specifies two members. The first member contains one 10-sample signal. The second member contains a 17-sample signal and a 27-sample signal.

Example:

`{{timetable(seconds(1:10)',randn(10,3)),timetable(seconds(1:7)',randn(7,2))},{timetable(seconds(1:3)',randn(3,1))}}` specifies two members. The first member contains three signals sampled at 1 Hz for 10 seconds and two signals sampled at 1 Hz for 7 seconds. The second member contains one signal sampled at 1 Hz for 3 seconds.

tinfo — Time information for new members

scalar | vector | matrix | duration scalar | duration vector

Time information for new members, specified as a scalar, a vector, a matrix, a duration scalar, or a duration vector. This argument is valid only if the “TimeInformation” on page 1-0 property of `lss` is `'sampleRate'`, `'sampleTime'`, or `'timeValues'`.

- If “TimeInformation” on page 1-0 is `'sampleRate'`, then `tinfo` specifies sample rate values.
- If “TimeInformation” on page 1-0 is `'sampleTime'`, then `tinfo` specifies sample time values.
- If “TimeInformation” on page 1-0 is `'timeValues'`, then `tinfo` specifies time values.

If you add multiple members to a set, then specifying only one value of `tinfo` sets the same value for all members. If you want to specify a different value for each new member, then set `tinfo` to have multiple values.

When no source has been specified, or when the labeled signal set source is empty, you can change the “TimeInformation” on page 1-0 property to `'sampleRate'`, `'sampleTime'`, or `'timeValues'` to make `lss` interpret `tinfo` correctly.

Example: `addMembers(ks, {randn(10,5), randn(10,3)}, seconds([1 2]))` adds two new members with different time information to `ks = labeledSignalSet(randn(10,3), 'SampleTime', seconds(1))`.

Example: `addMembers(ks, {randn(10,5), randn(10,3)}, [1:10;2:2:20]')` adds two new members with different time information to `ks = labeledSignalSet(randn(10,3), 'TimeValues', 1:10)`.

mnames — Member names

character vector | string scalar | cell array of character vectors | string array

Member names, specified as a character vector, a string scalar, a cell array of character vectors, or a string array.

Example: `labeledSignalSet({randn(100,1) randn(10,1)}, 'MemberNames', {'llama' 'alpaca'})` specifies a set of random signals with two members, 'llama' and 'alpaca'.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

allnodes

Tree nodes

Syntax

```
N = allnodes(T)
N = allnodes(T, 'deppos')
```

Description

`allnodes` is a tree management utility that returns one of two node descriptions: either indices, or depths and positions.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

`N = allnodes(T)` returns the indices of all the nodes of the tree T in column vector N .

`N = allnodes(T, 'deppos')` returns the depths and positions of all the nodes in matrix N .

$N(i, 1)$ is the depth and $N(i, 2)$ the position of the node i .

Examples

Return Nodes of Wavelet Packet Tree

This example shows how to obtain the depth-position and linear indices of a wavelet packet tree.

Load the noisy Doppler signal and obtain the wavelet packet decomposition down to the level 4 using the 'db2' wavelet.

```
load noisdopp;
T = wptdec(noisdopp, 4, 'db2');
```

Obtain the depth-position indices.

```
DepthPosition = allnodes(T, 'deppos');
```

Obtain the corresponding linear indices.

```
LinearIndices = allnodes(T);
```

Display the correspondence in a table.

```
table(DepthPosition, LinearIndices)
```

```
ans=31x2 table
   DepthPosition   LinearIndices
   _____   _____
         0         0
         1         1
         1         2
         2         3
         2         4
         2         5
         2         6
         3         7
         3         8
         3         9
         3        10
         3        11
         3        12
         3        13
         3        14
         4        15
         :
```

Introduced before R2006a

appcoef

1-D approximation coefficients

Syntax

```
A = appcoef(C,L,wname)
A = appcoef(C,L,LoR,HiR)
A = appcoef( ___,N)
```

Description

`A = appcoef(C,L,wname)` returns the approximation coefficients at the coarsest scale using the wavelet decomposition structure `[C,L]` of a 1-D signal and the wavelet specified by `wname`. (See `wavedec` for more information.)

`A = appcoef(C,L,LoR,HiR)` uses the lowpass reconstruction filter `LoR` and highpass reconstruction filter `HiR`. (See `wfilters` for more information.)

`A = appcoef(___,N)` returns the approximation coefficients at level `N`. If `[C,L]` is the `M`-level wavelet decomposition structure of a 1-D signal, then $0 \leq N \leq M$.

Examples

Level 3 Approximation Coefficients

This example shows how to extract the level 3 approximation coefficients.

Load the signal consisting of electricity usage data.

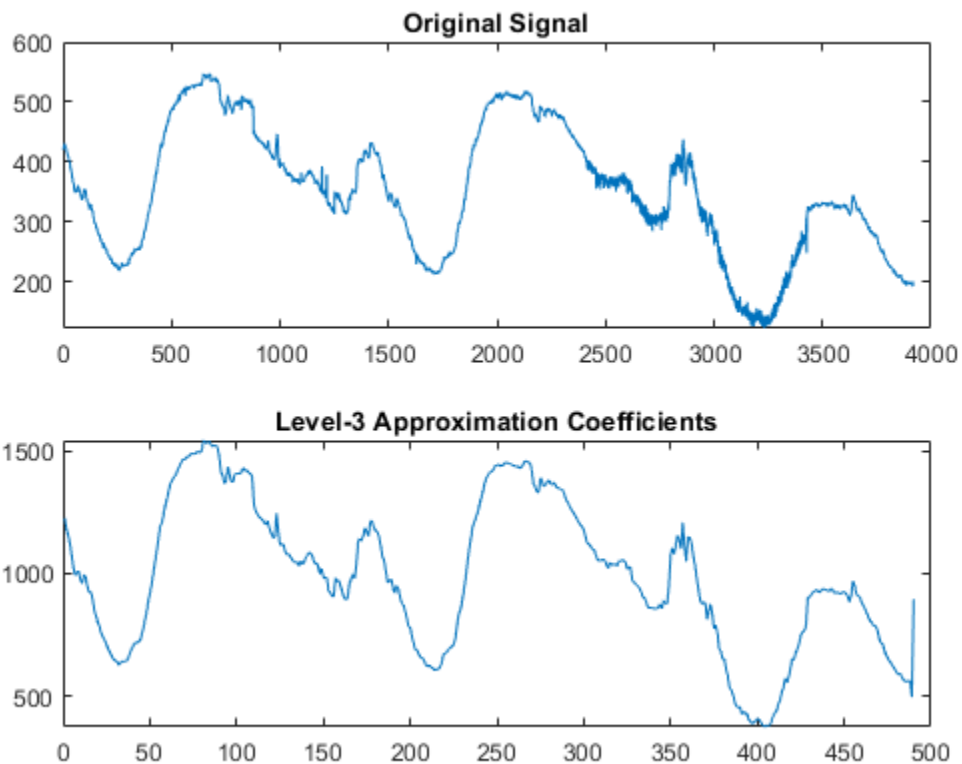
```
load leleccum;
sig = leleccum(1:3920);
```

Obtain the DWT down to level 5 with the 'sym4' wavelet.

```
[C,L] = wavedec(sig,5,'sym4');
```


Extract the level-3 approximation coefficients. Plot the original signal and the approximation coefficients.

```
Lev = 3;  
a3 = appcoef(C,L,'sym4',Lev);  
subplot(2,1,1)  
plot(sig); title('Original Signal');  
subplot(2,1,2)  
plot(a3); title('Level-3 Approximation Coefficients');
```



You can substitute any value from 1 to 5 for Lev to obtain the approximation coefficients for the corresponding level.

Input Arguments

C — Wavelet decomposition vector

real-valued vector

Wavelet decomposition vector of a 1-D signal, specified as a real-valued vector. C is the output of `wavedec`. The bookkeeping vector L is used to parse the coefficients in the wavelet decomposition vector by level.

Example: `[C,L] = wavedec(randn(1,256),4,'coif1')` returns the 4-level wavelet decomposition of a vector.

Data Types: `double`

L — Bookkeeping vector

vector of positive integers

Bookkeeping vector of the wavelet decomposition of a 1-D signal, specified as a vector of positive integers. The bookkeeping vector is used to parse the coefficients in the wavelet decomposition vector C by level.

Example: `[C,L] = wavedec(randn(1,256),4,'coif1')` returns the 4-level wavelet decomposition of a vector.

Data Types: `double`

wname — Wavelet

character vector | string scalar

Wavelet used to generate the wavelet decomposition of a 1-D signal, specified as a character vector or string scalar. The wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wavemngr` for the wavelets available in each family.

Example: `'db4'`

LoR — Wavelet lowpass reconstruction filter

even-length real-valued vector

Wavelet lowpass reconstruction filter, specified as an even-length real-valued vector. LoR must be the same length as HiR. LoR must be the lowpass reconstruction filter associated with the wavelet used to create the wavelet decomposition structure [C,L]. (See `wfilters` for more information.)

HiR — Wavelet highpass reconstruction filter

even-length real-valued vector

Wavelet highpass reconstruction filter, specified as an even-length real-valued vector. HiR must be the same length as LoR. HiR must be the highpass reconstruction filter associated with the wavelet used to create the wavelet decomposition structure [C,L]. (See `wfilters` for more information.)

N — Approximation coefficients level

positive integer

Approximation coefficients level, specified as a positive integer. If [C,L] is the M-level wavelet decomposition structure of a 1-D signal, then $0 \leq N \leq M$.

Output Arguments

A — Approximation coefficients

real-valued vector

Approximation coefficients at level N, returned as a real-valued vector.

Algorithms

The input vectors C and L contain all the information about the signal decomposition.

Let $NMAX = \text{length}(L) - 2$; then $C = [A(NMAX) \ D(NMAX) \ \dots \ D(1)]$ where A and the D are vectors. If $N = NMAX$, then a simple extraction is done; otherwise, `appcoef` computes iteratively the approximation coefficients using the inverse wavelet transform.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

detcoef | wavedec

Introduced before R2006a

appcoef2

2-D approximation coefficients

Syntax

```
A = appcoef2(C,S,wname)
A = appcoef2(C,S,LoR,HiR)
A = appcoef2( ____,N)
```

Description

`A = appcoef2(C,S,wname)` returns the approximation coefficients at the coarsest scale using the wavelet decomposition structure `[C,S]` of a 2-D signal and the wavelet specified by `wname`. (See `wavedec2` for more information.)

`A = appcoef2(C,S,LoR,HiR)` uses the lowpass reconstruction filter `LoR` and highpass reconstruction filter `HiR`. (See `wfilters` for more information.)

`A = appcoef2(____,N)` returns the approximation coefficients at level `N`. If `[C,S]` is the `M`-level wavelet decomposition structure of a 2-D signal, then $0 \leq N \leq M$.

Examples

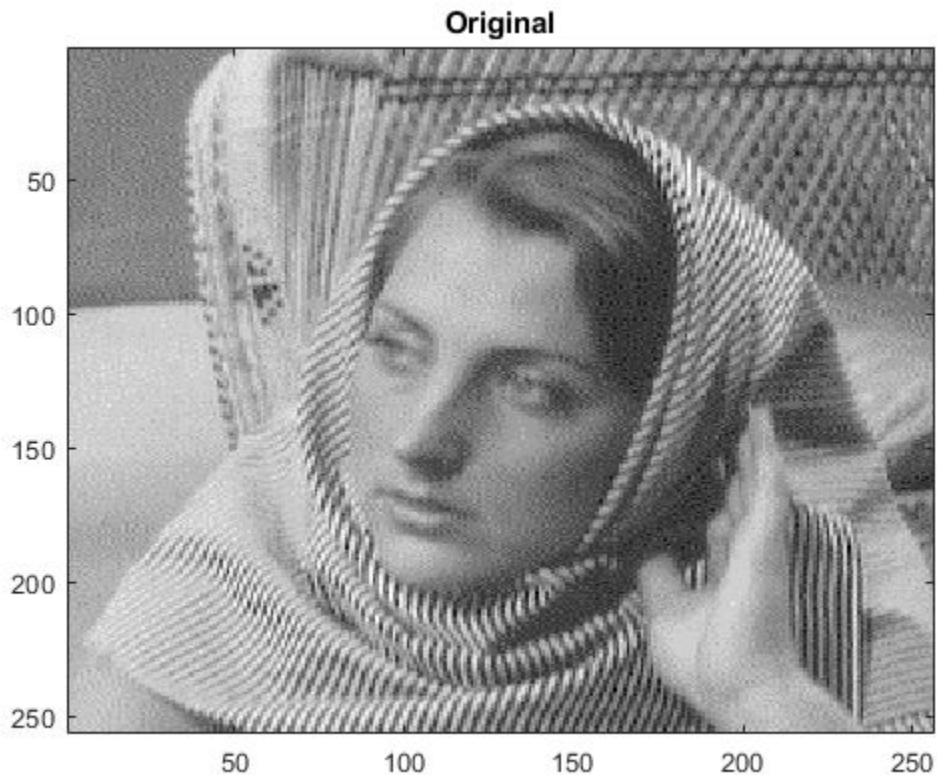
Reconstruct Approximation Coefficients of an Image

This example shows how to reconstruct approximation coefficients from a multilevel wavelet decomposition of an image.

Set the DWT extension mode to zero-padding. Load and display an image.

```
dwtmode('zpd','nodisp')
load woman
image(X)
```

```
colormap(map)
title('Original')
```



```
size(X)
ans = 1x2
    256    256
```

Perform a three-level wavelet decomposition of the image using the `db1` wavelet. Display the number of elements in the coefficients array `cfs`, and the contents of the bookkeeping matrix `inds`. Note that `cfs` has the same number of elements as `X`.

```
wv = 'db1';  
[cfs,inds] = wavedec2(X,3,wv);  
numel(X)  
  
ans = 65536  
  
numel(cfs)  
  
ans = 65536  
  
inds  
  
inds = 5×2  
  
    32    32  
    32    32  
    64    64  
   128   128  
   256   256
```

Extract and display the approximation coefficients at level 2.

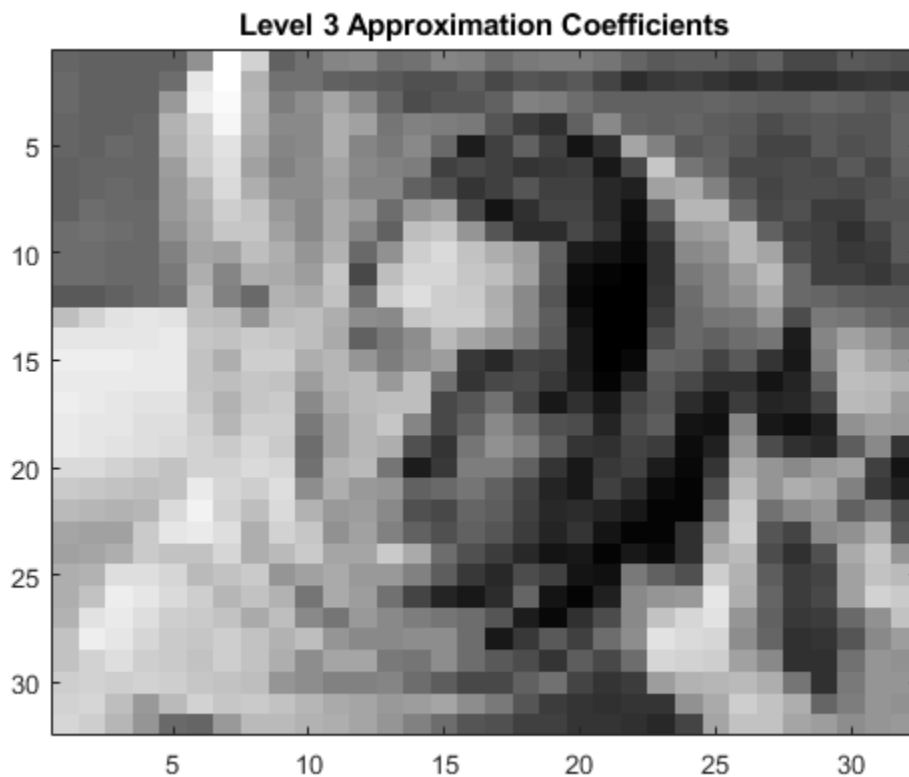
```
cfs2 = appcoef2(cfs,inds,wv,2);  
figure;  
imagesc(cfs2)  
colormap('gray')  
title('Level 2 Approximation Coefficients')
```



```
size(cfs2)
ans = 1x2
      64    64
```

Extract and display the approximation coefficients at level 3.

```
cfs3 = appcoef2(cfs,inds,wv,3);
figure;
imagesc(cfs3)
colormap('gray')
title('Level 3 Approximation Coefficients')
```

```
size(cfs3)
ans = 1x2
      32    32
```

Input Arguments

C — Wavelet decomposition vector
real-valued vector

Wavelet decomposition vector of a 2-D signal, specified as a real-valued vector. **C** is the output of `wavedec2`. The bookkeeping matrix **S** contains the dimensions of the coefficients by level.

Example: `[C,S] = wavedec2(randn(256,256),4,'db4')` returns the 4-level wavelet decomposition of a matrix.

Data Types: `double`

S — Bookkeeping matrix

matrix of positive integers

Bookkeeping matrix of the wavelet decomposition of a 2-D signal, specified as a matrix of positive integers. The bookkeeping matrix is used to parse the coefficients in the wavelet decomposition vector **C** by level.

Example: `[C,S] = wavedec2(randn(256,256),4,'db4')` returns the 4-level wavelet decomposition of a matrix.

Data Types: `double`

wname — Wavelet

character vector | string scalar

Wavelet used to generate the wavelet decomposition of a 2-D signal, specified as a character vector or string scalar. The wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wavemngr` for the wavelets available in each family.

Example: `'db4'`

LoR — Wavelet lowpass reconstruction filter

even-length real-valued vector

Wavelet lowpass reconstruction filter, specified as an even-length real-valued vector. **LoR** must be the same length as **HiR**. **LoR** must be the lowpass reconstruction filter associated with the wavelet used to create the wavelet decomposition structure `[C,S]`. (See `wfilters` for more information.)

HiR — Wavelet highpass reconstruction filter

even-length real-valued vector

Wavelet highpass reconstruction filter, specified as an even-length real-valued vector. **HiR** must be the same length as **LoR**. **HiR** must be the highpass reconstruction filter

associated with the wavelet used to create the wavelet decomposition structure [C,S]. (See `wfilters` for more information.)

N — Approximation coefficients level

positive integer

Approximation coefficients level, specified as a positive integer. If [C,S] is the M-level wavelet decomposition structure of a 2-D signal, then $0 \leq N \leq M$.

Output Arguments

A — Approximation coefficients

real-valued matrix | real-valued 3-D array

Approximation coefficients at level N, returned as a real-valued matrix or 3-D real-valued array. If C and S are obtained from an indexed image analysis or a truecolor image analysis, A is an m-by-n matrix or an m-by-n-by-3 array, respectively.

For more information on image formats, see `image` and `imfinfo`.

Algorithms

The input vector C and bookkeeping matrix S contain all the information about the 2-D signal decomposition.

Let `NMAX = size(S,1)-2`; then `C = [A(NMAX) H(NMAX) V(NMAX) D(NMAX) ... H(1) V(1) D(1)]` where A, H, V, and D are vectors. If `N = NMAX`, then a simple extraction is done; otherwise, `appcoef2` computes iteratively the approximation coefficients using the inverse wavelet transform.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

`detcoef2` | `wavedec2`

Introduced before R2006a

bestlevt

Best level tree wavelet packet analysis

Syntax

```
T = bestlevt(T)
[T,E] = bestlevt(T)
```

Description

`bestlevt` is a one- or two-dimensional wavelet packet analysis function.

`bestlevt` computes the optimal complete subtree of an initial tree with respect to an entropy type criterion. The resulting complete tree may be of smaller depth than the initial one.

`T = bestlevt(T)` computes the modified wavelet packet tree T corresponding to the best level tree decomposition.

`[T,E] = bestlevt(T)` computes the best level tree T , and in addition, the best entropy value E .

The optimal entropy of the node, whose index is $j - 1$, is $E(j)$.

Examples

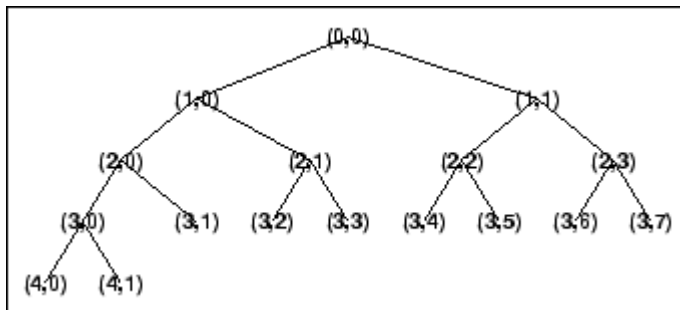
```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load signal.
load noisdopp;
x = noisdopp;
```

```
% Decompose x at depth 3 with db1 wavelet, using default
% entropy (shannon).
wpt = wptdec(x,3,'db1');
```

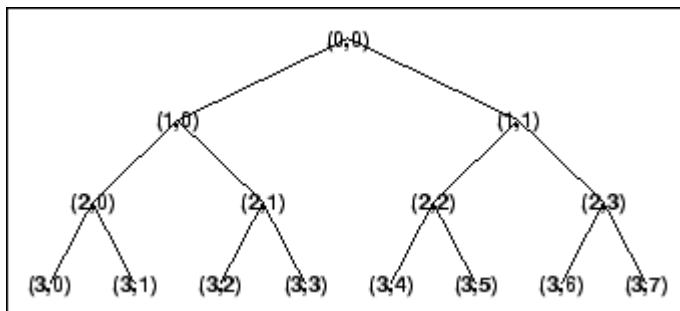
```
% Decompose the packet [3 0].
wpt = wpsplt(wpt,[3 0]);

% Plot wavelet packet tree wpt.
plot(wpt)
```



```
% Compute best level tree.
blt = bestlevt(wpt);

% Plot best level tree blt.
plot(blt)
```



Algorithms

See `besttree` algorithm section. The only difference is that the optimal tree is searched among the complete subtrees of the initial tree, instead of among all the binary subtrees.

See Also

besttree | wenergy | wpdec | wpdec2

Introduced before R2006a

besttree

Best tree wavelet packet analysis

Syntax

```
T = besttree(T)
[T,E] = besttree(T)
[T,E,N] = besttree(T)
```

Description

`besttree` is a one- or two-dimensional wavelet packet analysis function that computes the optimal subtree of an initial tree with respect to an entropy type criterion. The resulting tree may be much smaller than the initial one.

Following the organization of the wavelet packets library, it is natural to count the decompositions issued from a given orthogonal wavelet.

A signal of length $N = 2^L$ can be expanded in α different ways, where α is the number of binary subtrees of a complete binary tree of depth L .

As a result, we can conclude that $\alpha \geq 2^{N/2}$ (for more information, see the Mallat's book given in References at page 323).

This number may be very large, and since explicit enumeration is generally intractable, it is interesting to find an optimal decomposition with respect to a convenient criterion, computable by an efficient algorithm. We are looking for a minimum of the criterion.

`T = besttree(T)` computes the best tree T corresponding to the best entropy value.

`[T,E] = besttree(T)` computes the best tree T and, in addition, the best entropy value E .

The optimal entropy of the node, whose index is $j - 1$, is $E(j)$.

`[T,E,N] = besttree(T)` computes the best tree T , the best entropy value E and, in addition, the vector N containing the indices of the merged nodes.

Examples

Best Wavelet Packet Tree

This example shows to obtain the optimal wavelet packet tree based on an entropy criterion.

Load the noisy Doppler signal. Obtain the wavelet packet tree down to level 4 with the 'sym4' wavelet. Use the periodic extension mode.

```
dwtmode('per');
```

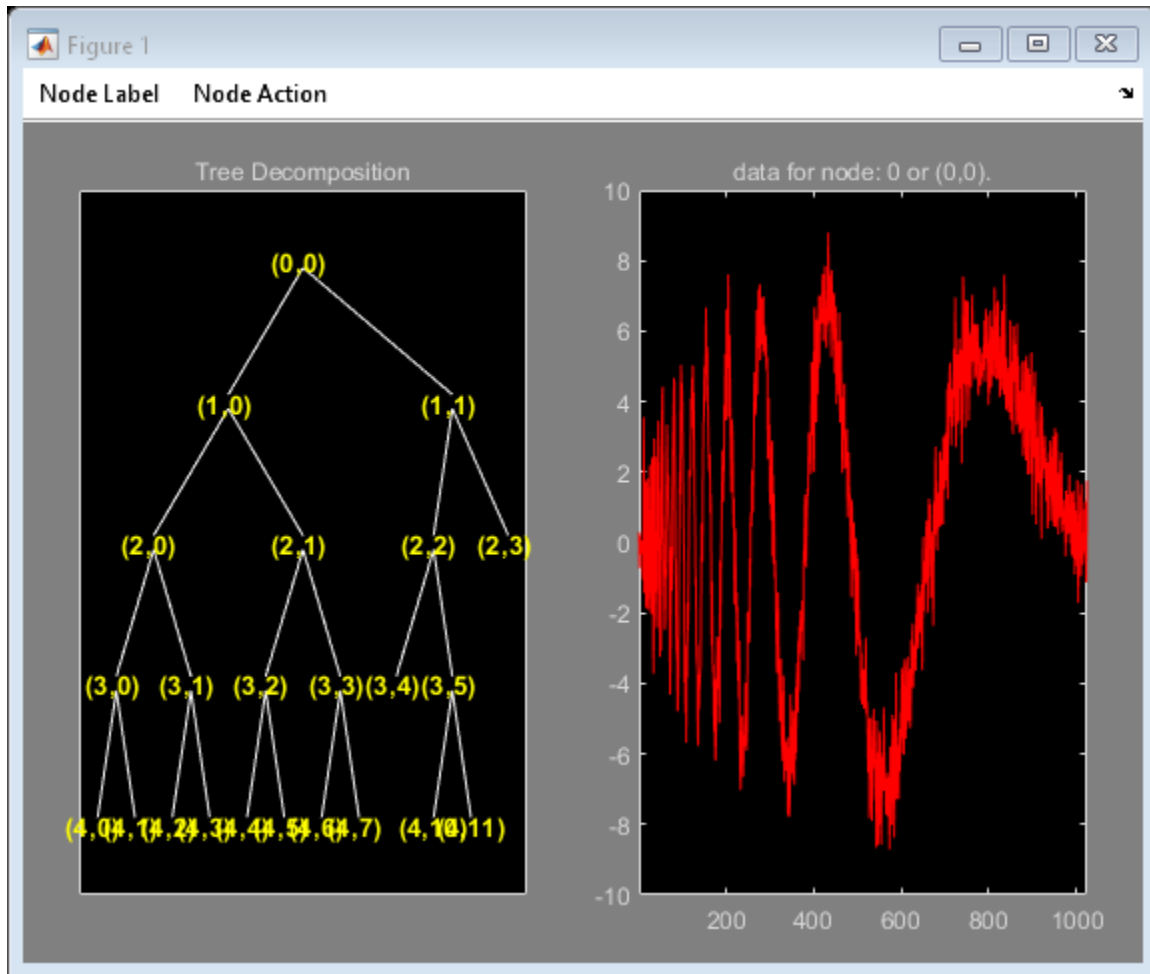
```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!  WARNING: Change DWT Extension Mode  !
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
*****
**  DWT Extension Mode: Periodization  **
*****
```

```
load noisdopp;
T = wpdec(noisdopp,4,'sym4');
```

Obtain the best wavelet packet tree and plot the result.

```
BstTree = besttree(T);
plot(BstTree)
```



Return the DWT extension mode to the default value.

```
dwtmode('sym');
```

```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
! WARNING: Change DWT Extension Mode !
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
*****
```

```
** DWT Extension Mode: Symmetrization (half-point) **  
*****
```

Algorithms

Consider the one-dimensional case. Starting with the root node, the best tree is calculated using the following scheme. A node N is split into two nodes $N1$ and $N2$ if and only if the sum of the entropy of $N1$ and $N2$ is lower than the entropy of N . This is a local criterion based only on the information available at the node N .

Several entropy type criteria can be used (see `wenergy` for more information). If the entropy function is an additive function along the wavelet packet coefficients, this algorithm leads to the best tree.

Starting from an initial tree T and using the merging side of this algorithm, we obtain the best tree among all the binary subtrees of T .

References

Coifman, R.R.; M.V. Wickerhauser (1992), "Entropy-based algorithms for best basis selection," *IEEE Trans. on Inf. Theory*, vol. 38, 2, pp. 713-718.

Mallat, S. (1998), *A wavelet tour of signal processing*, Academic Press.

See Also

`bestlevt` | `wenergy` | `wpcoef` | `wpdec` | `wpdec2` | `wprcoef`

Topics

"Reconstructing a Signal Approximation from a Node"

Introduced before R2006a

biorfilt

Biorthogonal wavelet filter set

Syntax

```
[Lo_D,Hi_D,Lo_R,Hi_R] = biorfilt(DF,RF)
[Lo_D1,Hi_D1,Lo_R1,Hi_R1,Lo_D2,Hi_D2,Lo_R2,Hi_R2] =
biorfilt(DF,RF,'8')
```

Description

The `biorfilt` command returns either four or eight filters associated with biorthogonal wavelets.

`[Lo_D,Hi_D,Lo_R,Hi_R] = biorfilt(DF,RF)` computes four filters associated with the biorthogonal wavelet specified by decomposition filter *DF* and reconstruction filter *RF*. These filters are

Lo_D	Decomposition low-pass filter
Hi_D	Decomposition high-pass filter
Lo_R	Reconstruction low-pass filter
Hi_R	Reconstruction high-pass filter

`[Lo_D1,Hi_D1,Lo_R1,Hi_R1,Lo_D2,Hi_D2,Lo_R2,Hi_R2] = biorfilt(DF,RF,'8')` returns eight filters, the first four associated with the decomposition wavelet, and the last four associated with the reconstruction wavelet.

It is well known in the subband filtering community that if the same FIR filters are used for reconstruction and decomposition, then symmetry and exact reconstruction are incompatible (except with the Haar wavelet). Therefore, with biorthogonal filters, two wavelets are introduced instead of just one:

One wavelet, $\tilde{\psi}$, is used in the analysis, and the coefficients of a signal *s* are

$$\tilde{c}_{j,k} = \int s(x)\tilde{\psi}_{j,k}(x)dx$$

The other wavelet, ψ , is used in the synthesis:

$$s = \sum_{j,k} \tilde{c}_{j,k} \psi_{j,k}$$

Furthermore, the two wavelets are related by duality in the following sense:

$$\int \tilde{\psi}_{j,k}(x) \psi_{j',k'}(x) dx = 0 \text{ as soon as } j \neq j' \text{ or } k \neq k' \text{ and}$$

$$\int \tilde{\phi}_{0,k}(x) \phi_{0,k'}(x) dx = 0 \text{ as soon as } k \neq k'.$$

It becomes apparent, as A. Cohen pointed out in his thesis (p. 110), that “the useful properties for analysis (e.g., oscillations, null moments) can be concentrated in the $\tilde{\psi}$ function; whereas, the interesting properties for synthesis (regularity) are assigned to the ψ function. The separation of these two tasks proves very useful.”

$\tilde{\psi}$ and ψ can have very different regularity properties, ψ being more regular than $\tilde{\psi}$.

The $\tilde{\psi}$, ψ , $\tilde{\phi}$ and ϕ functions are zero outside a segment.

Examples

Biorthogonal Filters and Transfer Functions

This example shows how to obtain the decomposition (analysis) and reconstruction (synthesis) filters for the 'bior3.5' wavelet.

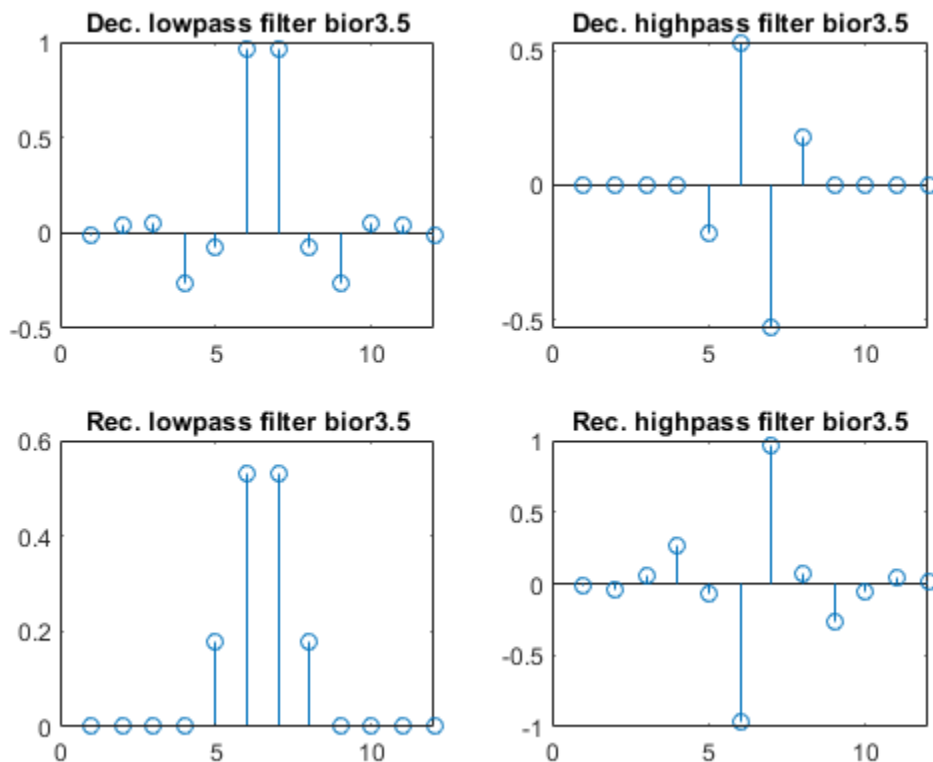
Determine the two scaling and wavelet filters associated with the 'bior3.5' wavelet.

```
wv = 'bior3.5';
[Rf,Df] = biorwavf(wv);
[LoD,HiD,LoR,HiR] = biorfilt(Df,Rf);
```

Plot the filter impulse responses.

```
subplot(2,2,1)
stem(LoD)
title(['Dec. lowpass filter ',wv])
subplot(2,2,2)
stem(HiD)
title(['Dec. highpass filter ',wv])
```

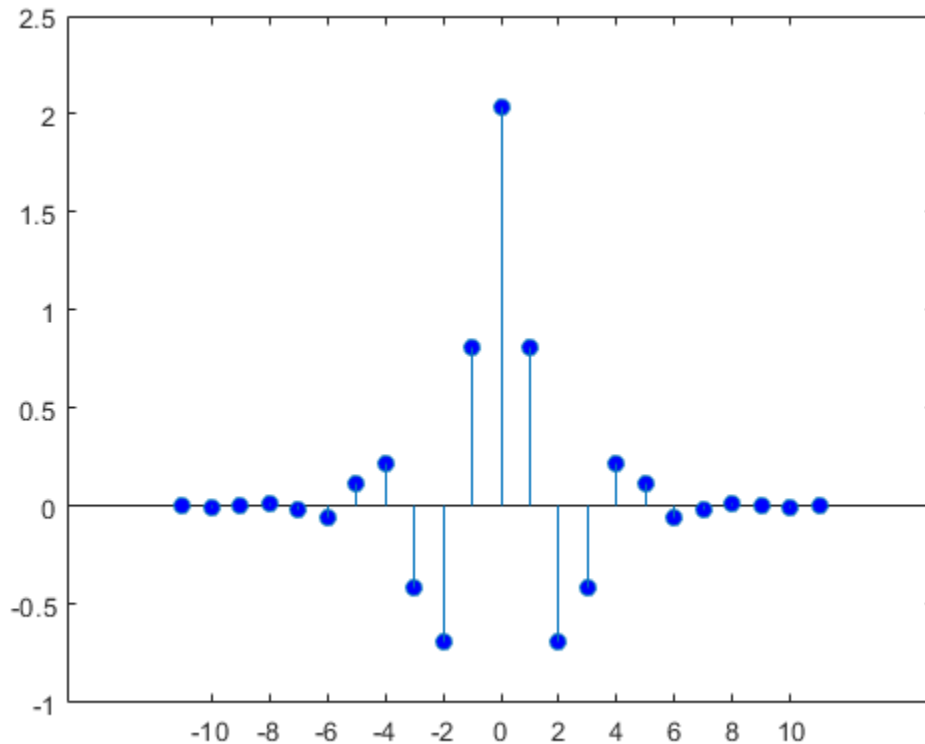
```
subplot(2,2,3)
stem(LoR)
title(['Rec. lowpass filter ',wv])
subplot(2,2,4)
stem(HiR)
title(['Rec. highpass filter ',wv])
```



Demonstrate that autocorrelations at even lags are only zero for dual pairs of filters. Examine the autocorrelation sequence for the lowpass decomposition filter.

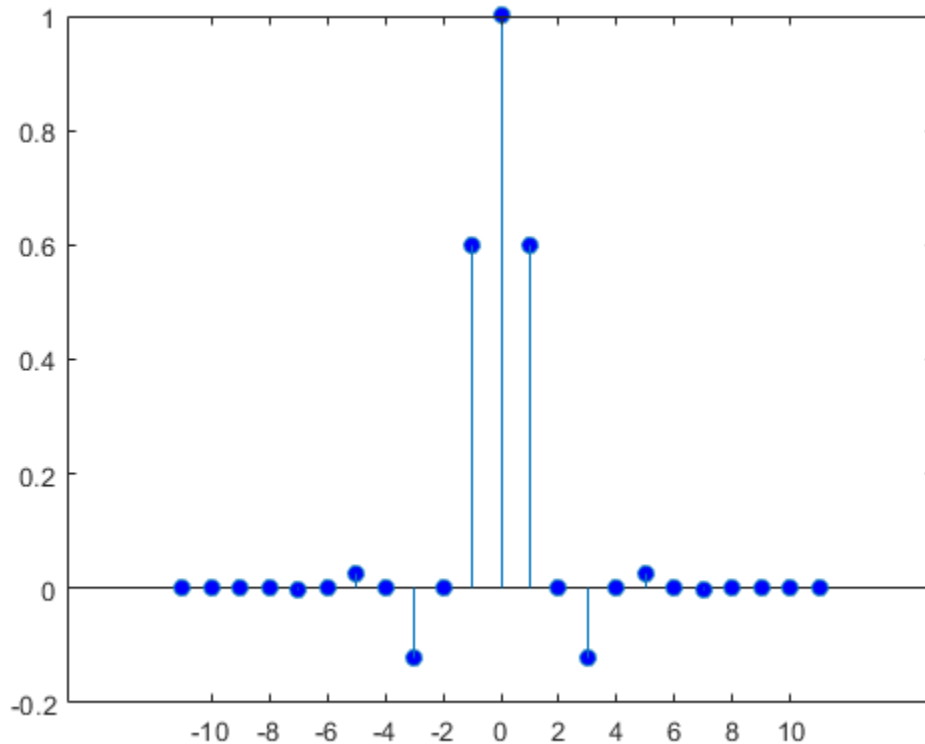
```
npad = 2*length(LoD)-1;
LoDxcr = fftshift(iff(abs(fft(LoD,npad)).^2));
lags = -floor(npad/2):floor(npad/2);
figure
```

```
stem(lags,LoDxcr,'markerfacecolor',[0 0 1])
set(gca,'xtick',-10:2:10)
```



Examine the cross correlation sequence for the lowpass decomposition and synthesis filters. Compare the result with the preceding figure.

```
npad = 2*length(LoD)-1;
xcr = fftshift(iffshift(fft(LoD,npad).*conj(fft(LoR,npad))));
lags = -floor(npad/2):floor(npad/2);
stem(lags,xcr,'markerfacecolor',[0 0 1])
set(gca,'xtick',-10:2:10)
```



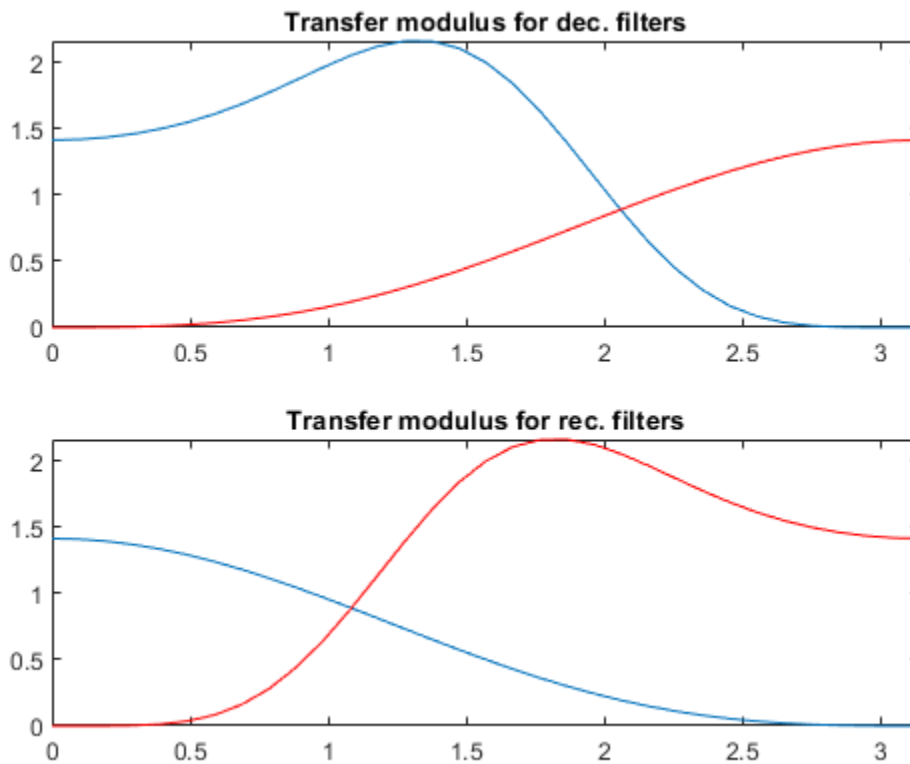
Compare the transfer functions of the analysis and synthesis scaling and wavelet filters

```
dftLoD = fft(LoD,64);
dftLoD = dftLoD(1:length(dftLoD)/2+1);
dftHiD= fft(HiD,64);
dftHiD = dftHiD(1:length(dftHiD)/2+1);
dftLoR = fft(LoR,64);
dftLoR = dftLoR(1:length(dftLoR)/2+1);
dftHiR = fft(HiR,64);
dftHiR = dftHiR(1:length(dftHiR)/2+1);
df = (2*pi)/64;
freqvec = 0:df:pi;

subplot(2,1,1)
```



```
plot(freqvec,abs(dftLoD),freqvec,abs(dftHiD),'r')
axis tight
title('Transfer modulus for dec. filters')
subplot(2,1,2)
plot(freqvec,abs(dftLoR),freqvec,abs(dftHiR),'r')
axis tight
title('Transfer modulus for rec. filters')
```



References

Cohen, A. (1992), "Ondelettes, analyses multirésolution et traitement numérique du signal," *Ph. D. Thesis*, University of Paris IX, DAUPHINE.

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed.

See Also

`biorwavf` | `orthfilt`

Introduced before R2006a

biorwavf

Biorthogonal spline wavelet filter

Syntax

```
[RF,DF] = biorwavf(wname)
```

Description

`[RF,DF] = biorwavf(wname)` returns the reconstruction (synthesis) and decomposition (analysis) scaling filters, RF and DF, respectively, associated with the biorthogonal wavelet specified by `wname`.

Examples

Biorthogonal Spline Wavelet Filter

Return the biorthogonal spline wavelet scaling filters with two vanishing moments.

```
wname = 'bior2.2';  
[RF,DF] = biorwavf(wname)
```

```
RF = 1×3
```

```
    0.2500    0.5000    0.2500
```

```
DF = 1×5
```

```
   -0.1250    0.2500    0.7500    0.2500   -0.1250
```

Input Arguments

wname — Name of biorthogonal wavelet

character vector | string scalar

Name of biorthogonal wavelet, specified as 'biorNr.Nd' where possible values for Nr and Nd are as follows:

Nr = 1	Nd = 1 , 3 or 5
Nr = 2	Nd = 2 , 4 , 6 or 8
Nr = 3	Nd = 1 , 3 , 5 , 7 or 9
Nr = 4	Nd = 4
Nr = 5	Nd = 5
Nr = 6	Nd = 8

Nr and Nd are the numbers of vanishing moments for the reconstruction and decomposition filters, respectively.

Example: 'biorwavf3.7'

Output Arguments

RF — Reconstruction filter

real-valued vector

Reconstruction filter associated with the biorthogonal wavelet wname, returned as a real-valued vector.

DF — Decomposition filter

real-valued vector

Decomposition filter associated with the biorthogonal wavelet wname, returned as a real-valued vector.

See Also

biorfilt | waveinfo

Introduced before R2006a

BPfrequencies

CWT filter bank bandpass center frequencies

Note BPfrequencies is not recommended and may be removed in a future release. Use centerFrequencies instead.

Syntax

```
bpcf = BPfrequencies(fb)
```

Description

`bpcf = BPfrequencies(fb)` returns the wavelet bandpass center frequencies `bpcf` for the CWT filter bank `fb`. Frequencies are ordered from high to low. Frequencies are in cycles/sample if a sampling frequency or sampling period is not specified. If a sampling frequency is specified, `bpcf` has units of hertz. If a sampling period is specified, `bpcf` has units cycles/unit time where the time unit is the same as the duration `SamplingPeriod`.

Examples

Wavelet Bandpass Center Frequencies

Create a CWT filter bank.

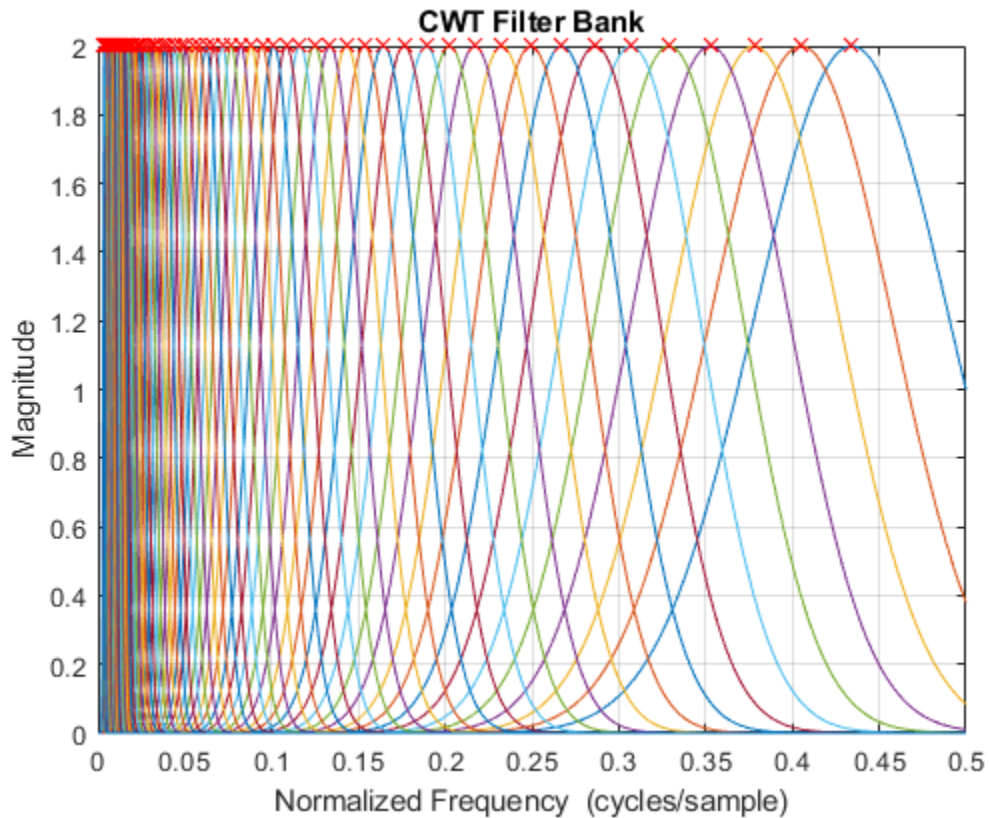
```
fb = cwtfilterbank;
```

Return the bandpass center frequencies.

```
bpcf = centerFrequencies(fb);
```

Plot the frequency responses of the filter bank and the bandpass center frequencies. Observe that the bandpass center frequencies correspond to the peaks of the frequency response of each wavelet in the filter bank.

```
freqz(fb)  
hold on  
plot(bpcf,2*ones(size(bpcf)), 'rx')
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

bpcf — Wavelet bandpass center frequencies

real-valued vector

Wavelet bandpass center frequencies, returned as a real-valued vector of length N_s where N_s is the number of scales in the filter bank. Frequencies are ordered from high to low. Frequencies are in cycles/sample if a sampling frequency or sampling period is not specified. If a sampling frequency is specified, `bpcf` has units of hertz. If a sampling period is specified, `bpcf` has units cycles/unit time where the time unit is the same as the duration `SamplingPeriod`.

Compatibility Considerations

BPfrequencies will be removed

Not recommended starting in R2018b

The `BPfrequencies` object function of `cwtfilterbank` has been renamed `centerFrequencies`. The functionality remains unchanged. `BPfrequencies` will be removed in a future release.

Functionality	What Happens When You Use This Functionality?	Use This Instead	Compatibility Considerations
BPfrequencies	Still runs	Use <code>centerFrequencies</code>	Replace all instances of <code>BPfrequencies</code> with <code>centerFrequencies</code> .

See Also

`centerPeriods` | `cwtfilterbank` | `freqz` | `powerbw`

Introduced in R2018a

BPperiods

CWT filter bank bandpass periods

Note BPperiods is not recommended and may be removed in a future release. Use centerPeriods instead.

Syntax

```
p = BPperiods(fb)
```

Description

`p = BPperiods(fb)` returns the wavelet bandpass periods, `p`, for the continuous wavelet transform (CWT) filter bank, `fb`.

Examples

Wavelet Filter Bank Bandpass Periods

Create two CWT filter banks. Set the sampling period of the first filter bank to 0.5 seconds, and the sampling frequency of the second filter bank to 2 Hz.

```
fb = cwtfilterbank('SamplingPeriod',seconds(0.5));  
fb2 = cwtfilterbank('SamplingFrequency',2);
```

Obtain the bandpass center periods of both filter banks. Inspect the first five elements of each.

```
bp = centerPeriods(fb);  
bp2 = centerPeriods(fb2);  
bp(1:5)
```

```
ans = 5x1 duration array  
    1.1517 sec
```

```
1.2344 sec
1.323  sec
1.418  sec
1.5197 sec
```

```
bp2(1:5)
```

```
ans = 5×1
```

```
1.1517
1.2344
1.3230
1.4180
1.5197
```

Obtain the bandpass center frequencies of the second filter bank. Inspect the reciprocals of the first five elements.

```
f2 = centerFrequencies(fb2);
1./f2(1:5)
```

```
ans = 5×1
```

```
1.1517
1.2344
1.3230
1.4180
1.5197
```

Input Arguments

fb — Continuous wavelet transform filter bank

object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

p — Wavelet bandpass filter periods

real-valued vector | duration array

Wavelet bandpass filter periods, returned as a real-valued vector of length N_s where N_s is the number of scales in the filter bank.

If `SamplingPeriod` is specified, `p` is a duration array with the same units and format as `SamplingPeriod`. If `SamplingFrequency` is specified, `p` is in seconds.

Compatibility Considerations

BPperiods will be removed

Not recommended starting in R2018b

The `BPperiods` object function of `cwtfilterbank` has been renamed `centerPeriods`. The functionality remains unchanged. `BPperiods` will be removed in a future release.

Functionality	What Happens When You Use This Functionality?	Use This Instead	Compatibility Considerations
BPperiods	Still runs	Use centerPeriods	Replace all instances of BPperiods with centerPeriods.

See Also

`centerFrequencies` | `cwtfilterbank` | `freqz` | `powerbw`

Introduced in R2018a

bswfun

Biorthogonal scaling and wavelet functions

Syntax

```
[PHIS,PSIS,PHIA,PSIA,XVAL] = bswfun(LoD,HiD,LoR,HiR)
bswfun(LoD,HiD,LoR,HiR,ITER)
bswfun(LoD,HiD,LoR,HiR,'plot')
bswfun(LoD,HiD,LoR,HiR,ITER,'plot')
bswfun(LoD,HiD,LoR,HiR,'plot',ITER)
```

Description

`[PHIS,PSIS,PHIA,PSIA,XVAL] = bswfun(LoD,HiD,LoR,HiR)` returns approximations on the grid `XVAL` of the two pairs of biorthogonal scaling and wavelet functions. `PHIS` and `PSIS` are the scaling and wavelet functions constructed from the decomposition filters, `LoD` and `HiD`. `PHIA` and `PSIA` are the scaling and wavelet functions constructed from the reconstruction filters, `LoR` and `HiR`.

`bswfun(LoD,HiD,LoR,HiR,ITER)` computes the two pairs of scaling and wavelet functions using `ITER` iterations.

`bswfun(LoD,HiD,LoR,HiR,'plot')` or `bswfun(LoD,HiD,LoR,HiR,ITER,'plot')` or `bswfun(LoD,HiD,LoR,HiR,'plot',ITER)` computes and plots the functions.

Examples

Biorthogonal Scaling and Wavelet from Lifting Scheme

This example shows how to obtain the biorthogonal scaling and wavelet functions corresponding to a lifting scheme. Obtain the lifting scheme for the CDF 3/1 wavelet.

```
lscdf = liftwave('cdf3.1');
```

Display the lifting scheme, which consists of two primal and one dual step.

```
Sc = displs(lscdf);
```

```
Sc
```

```
Sc = 6x50 char array
```

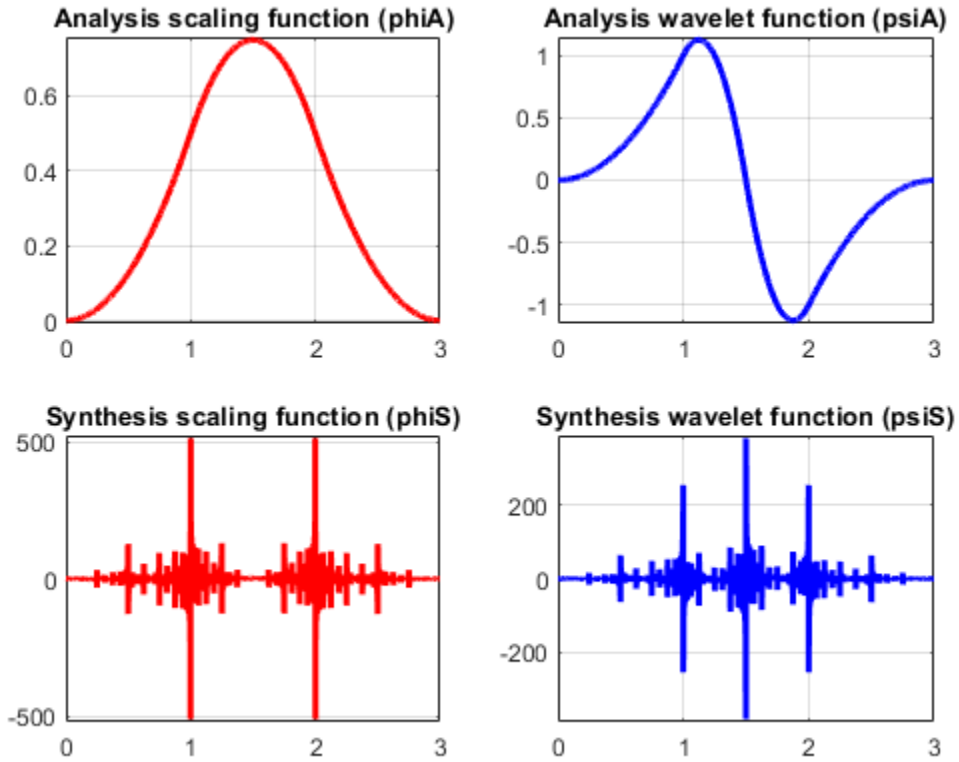
```
'lscdf = {...'
''p'          [ -0.33333333]      [-1]
''d'          [ -0.37500000 -1.12500000] [1]
''p'          [  0.44444444]      [0]
' [  2.12132034] [  0.47140452]      []
'};
```

Obtain the decomposition and reconstruction filters from the lifting scheme.

```
[LoD,HiD,LoR,HiR] = ls2filt(lscdf);
```

Visualize the scaling and wavelet function and their duals.

```
bswfun(LoD,HiD,LoR,HiR,'plot');
```



Algorithms

This function uses the cascade algorithm.

See Also

wavefun

Introduced before R2006a

centerFrequencies

CWT filter bank bandpass center frequencies

Syntax

```
bpcf = centerFrequencies(fb)
```

Description

`bpcf = centerFrequencies(fb)` returns the wavelet bandpass center frequencies `bpcf` for the CWT filter bank `fb`. Frequencies are ordered from high to low. Frequencies are in cycles/sample if a sampling frequency or sampling period is not specified. If a sampling frequency is specified, `bpcf` has units of hertz. If a sampling period is specified, `bpcf` has units cycles/unit time where the time unit is the same as the duration `SamplingPeriod`.

Examples

Wavelet Bandpass Center Frequencies

Create a CWT filter bank.

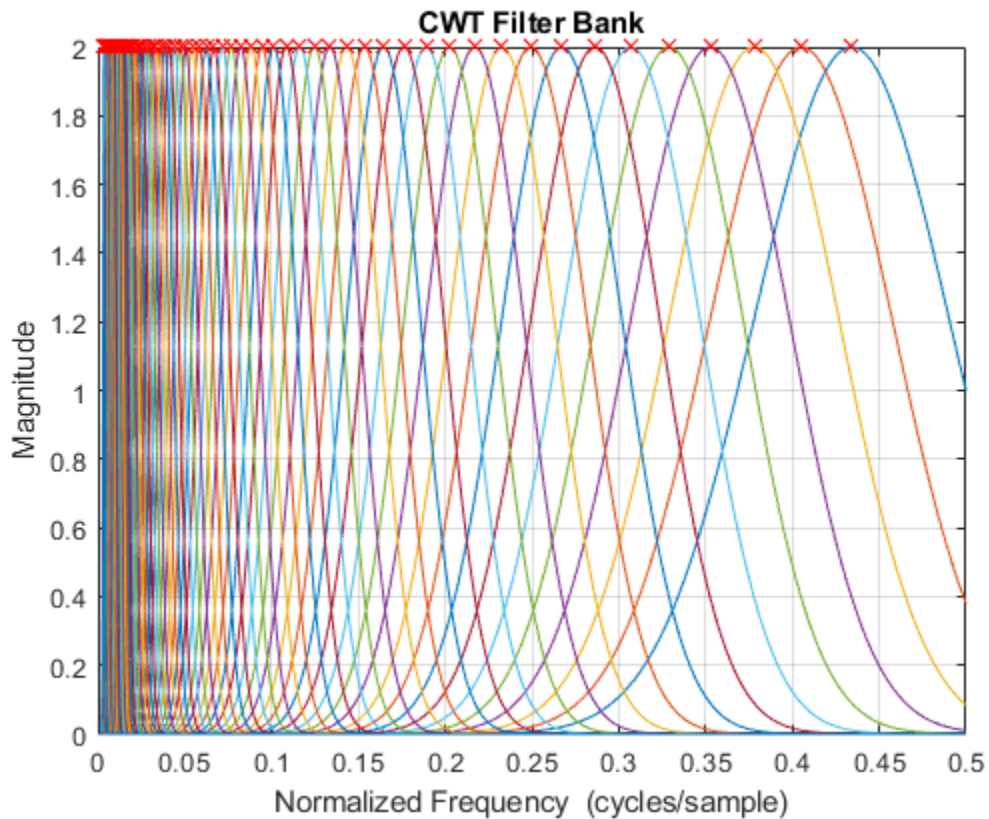
```
fb = cwtfilterbank;
```

Return the bandpass center frequencies.

```
bpcf = centerFrequencies(fb);
```

Plot the frequency responses of the filter bank and the bandpass center frequencies. Observe that the bandpass center frequencies correspond to the peaks of the frequency response of each wavelet in the filter bank.

```
freqz(fb)  
hold on  
plot(bpcf, 2*ones(size(bpcf)), 'rx')
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

bpcf — Wavelet bandpass center frequencies

real-valued vector

Wavelet bandpass center frequencies, returned as a real-valued vector of length `Ns` where `Ns` is the number of scales in the filter bank. Frequencies are ordered from high to low. Frequencies are in cycles/sample if a sampling frequency or sampling period is not specified. If a sampling frequency is specified, `bpcf` has units of hertz. If a sampling period is specified, `bpcf` has units cycles/unit time where the time unit is the same as the duration `SamplingPeriod`.

See Also

`centerPeriods` | `cwtfilterbank` | `freqz` | `powerbw`

Introduced in R2018b

centerFrequencies

Wavelet scattering bandpass center frequencies

Syntax

```
F = centerFrequencies(sf)
F = centerFrequencies(sf,filterbanks)
```

Description

`F = centerFrequencies(sf)` returns the wavelet bandpass center frequencies for all filter banks of the scattering decomposition framework, `sf`. The output `F` is a cell array with `Nfb` elements, where `Nfb` is the number of scattering filter banks. Each element of `F` is a real-valued vector. If you specify a sampling frequency in `sf`, `F` is in hertz. If you do not specify a sampling frequency, `F` is in cycles/sample.

If there is only one filter bank in the scattering framework, `F` is a real-valued vector containing the wavelet bandpass center frequencies.

`F = centerFrequencies(sf,filterbanks)` returns the wavelet bandpass center frequencies for the specified `filterbanks`. The argument `filterbanks` is a scalar or vector with all the elements between 1 and `Nfb` inclusive, where `Nfb` is the number of scattering filter banks.

Examples

Wavelet Bandpass Center Frequencies

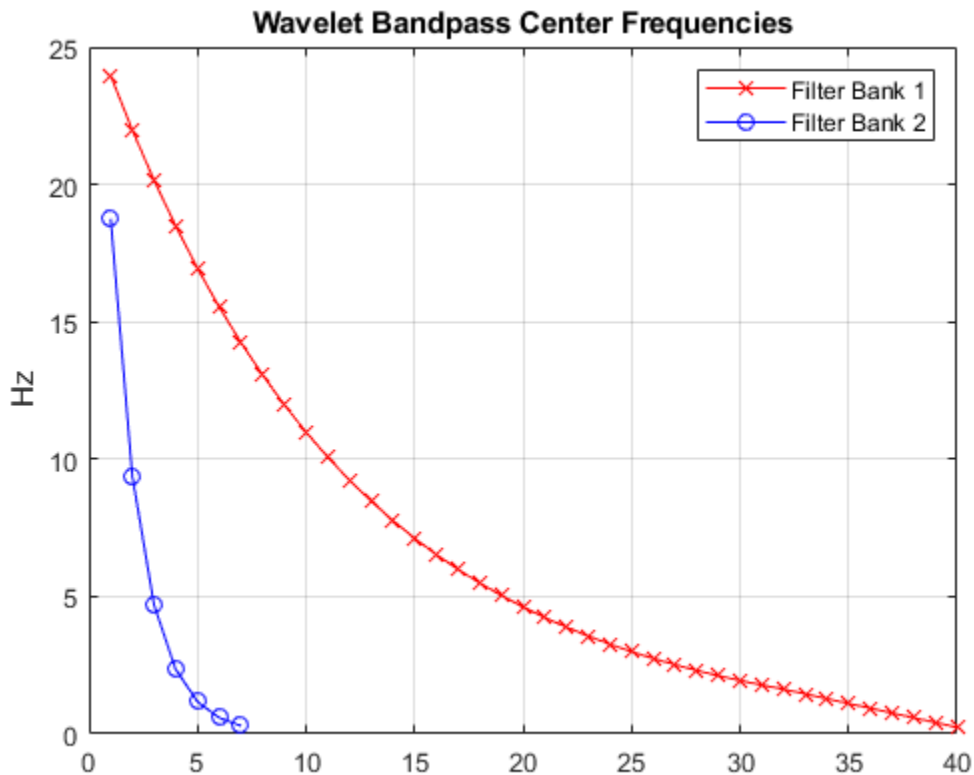
Create a scattering decomposition framework with a sampling frequency of 50 Hz.

```
sf = waveletScattering('SamplingFrequency',50)
sf =
    waveletScattering with properties:
```

```
SignalLength: 1024
InvarianceScale: 10.2400
QualityFactors: [8 1]
    Boundary: "periodic"
SamplingFrequency: 50
    Precision: "double"
OversamplingFactor: 0
```

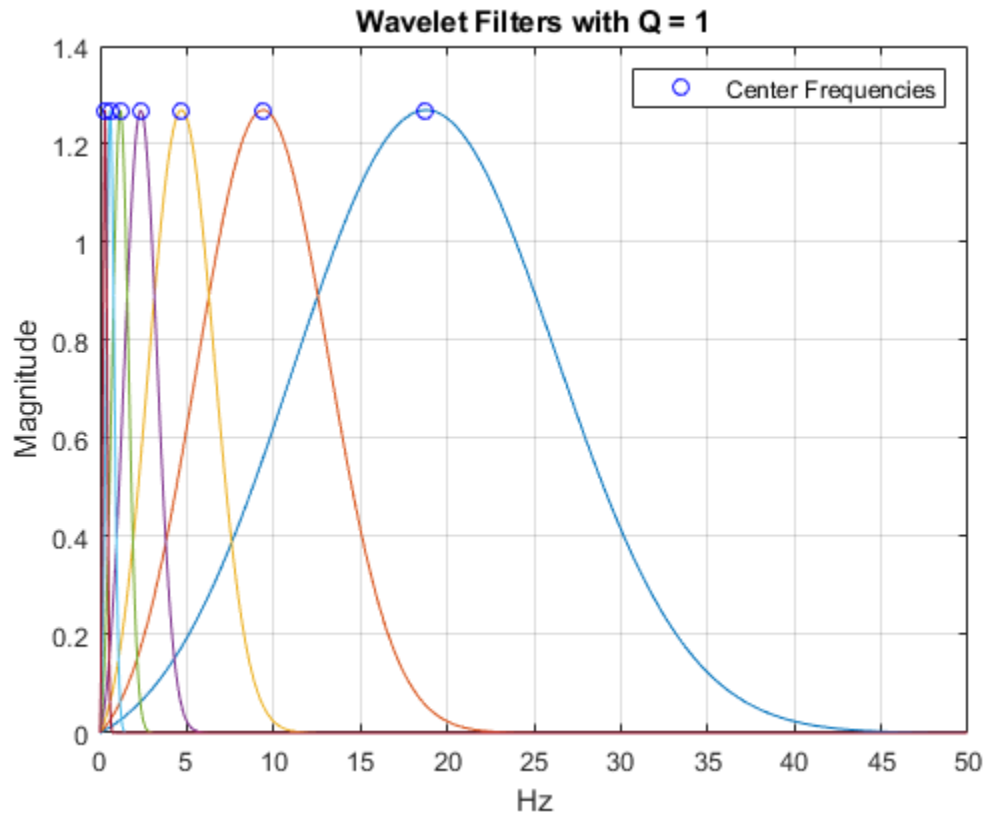
Plot the wavelet bandpass center frequencies for all the filter banks.

```
bpcf = centerFrequencies(sf);
plot(bpcf{1}, 'rx-')
hold on
plot(bpcf{2}, 'bo-')
grid on
title('Wavelet Bandpass Center Frequencies')
legend('Filter Bank 1', 'Filter Bank 2')
ylabel('Hz')
```



Plot the wavelets filters used in computing the second-order scattering coefficients.

```
orderCoef = 2;
[filters,f] = filterbank(sf);
figure
plot(f, filters{orderCoef+1}.psift)
grid on
title('Wavelet Filters with Q = 1')
xlabel('Hz')
ylabel('Magnitude')
hold on
pl = plot(bpcf{orderCoef},max(filters{orderCoef+1}.psift), 'bo');
legend(pl, 'Center Frequencies')
```



Input Arguments

sf — Scattering decomposition framework

waveletScattering object

Scattering decomposition framework, specified as a waveletScattering object.

filterbanks — Filter bank indices

positive integer | vector of integers

Filter bank indices, specified as a positive integer or vector of integers. Elements of `filterbanks` are integers between 1 and Nfb inclusive, where Nfb is the number of scattering filter banks.

Example: `F = centerFrequencies(sf, [1 2])` returns the wavelet bandpass center frequencies for the first two filter banks in `sf`.

Data Types: `double`

See Also

`waveletScattering`

Introduced in R2018b

centerPeriods

CWT filter bank bandpass center periods

Syntax

```
p = centerPeriods(fb)
```

Description

`p = centerPeriods(fb)` returns the wavelet bandpass center periods, `p`, for the continuous wavelet transform (CWT) filter bank, `fb`.

Examples

Wavelet Filter Bank Bandpass Periods

Create two CWT filter banks. Set the sampling period of the first filter bank to 0.5 seconds, and the sampling frequency of the second filter bank to 2 Hz.

```
fb = cwtfilterbank('SamplingPeriod',seconds(0.5));  
fb2 = cwtfilterbank('SamplingFrequency',2);
```

Obtain the bandpass center periods of both filter banks. Inspect the first five elements of each.

```
bp = centerPeriods(fb);  
bp2 = centerPeriods(fb2);  
bp(1:5)
```

```
ans = 5x1 duration array  
    1.1517 sec  
    1.2344 sec  
    1.323  sec  
    1.418  sec
```

```
1.5197 sec
```

```
bp2(1:5)
```

```
ans = 5×1
```

```
1.1517  
1.2344  
1.3230  
1.4180  
1.5197
```

Obtain the bandpass center frequencies of the second filter bank. Inspect the reciprocals of the first five elements.

```
f2 = centerFrequencies(fb2);  
1./f2(1:5)
```

```
ans = 5×1
```

```
1.1517  
1.2344  
1.3230  
1.4180  
1.5197
```

Input Arguments

fb — Continuous wavelet transform filter bank

object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

p — Wavelet bandpass center periods

real-valued vector | duration array

Wavelet bandpass center periods, returned as a real-valued vector of length `Ns` where `Ns` is the number of scales in the filter bank.

If `SamplingPeriod` is specified, `p` is a duration array with the same units and format as `SamplingPeriod`. If `SamplingFrequency` is specified, `p` is in seconds.

See Also

`centerFrequencies` | `cwtfilterbank` | `freqz` | `powerbw`

Introduced in R2018b

centfrq

Wavelet center frequency

Syntax

```
FREQ = centfrq('wname')  
FREQ = centfrq('wname',ITER)  
[FREQ,XVAL,RECFREQ] = centfrq('wname',ITER,'plot')
```

Description

`FREQ = centfrq('wname')` returns the center frequency in hertz of the wavelet function, `'wname'` (see `wavefun` for more information).

For `FREQ = centfrq('wname',ITER)`, `ITER` is the number of iterations performed by the function `wavefun`, which is used to compute the wavelet.

`[FREQ,XVAL,RECFREQ] = centfrq('wname',ITER,'plot')` returns, in addition, the associated center frequency based approximation `RECFREQ` on the 2^{ITER} points grid `XVAL` and plots the wavelet function and `RECFREQ`.

Examples

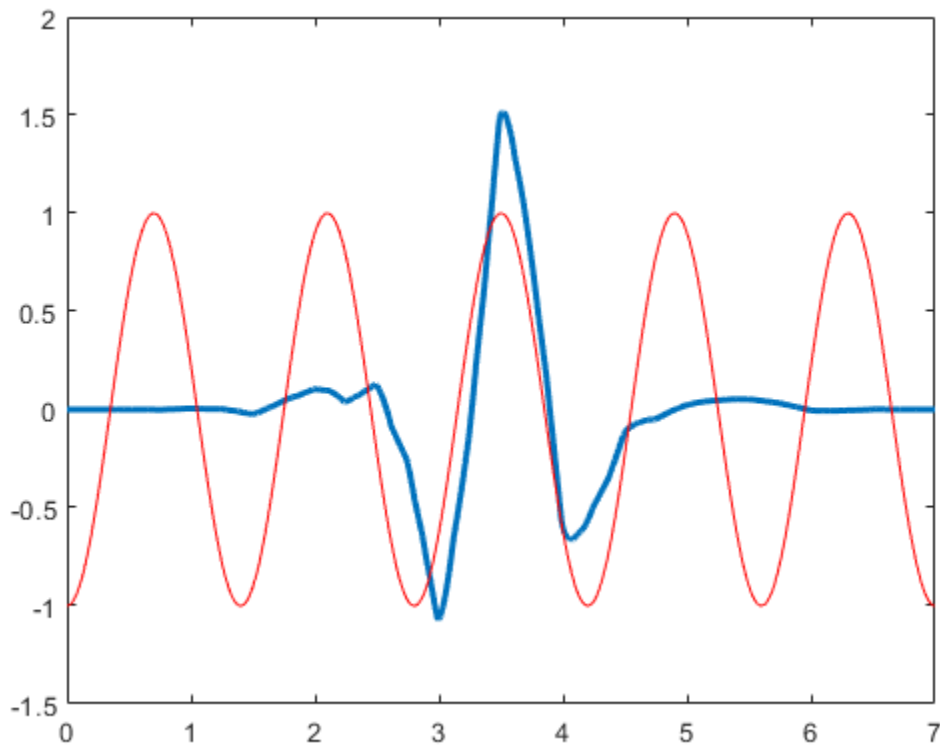
Determine Center Frequency

This example shows how to determine the center frequency in hertz for Daubechies' least-asymmetric wavelet with 4 vanishing moments.

```
cfreq = centfrq('sym4');
```

Obtain the wavelet and create a sine wave with a frequency equal to the center frequency, `cfreq`, of the wavelet. Use a starting phase of $-\pi$ for the sine wave to visualize how the oscillation in the sine wave matches the oscillation in the wavelet.

```
[~,psi,xval] = wavefun('sym4');  
y = cos(2*pi*cfreq*xval-pi);  
plot(xval,psi,'linewidth',2);  
hold on;  
plot(xval,y,'r');
```



Convert Scales to Frequencies

This example shows to convert scales to frequencies for the Morlet wavelet. There is an approximate inverse relationship between scale and frequency. Specifically, scale is

inversely proportional to frequency with the constant of proportionality being the center frequency of the wavelet.

Construct a vector of scales with 32 voices per octave over 5 octaves for data sampled at 1 kHz.

```
Fs = 1000;  
numvoices = 32;  
a0 = 2^(1/numvoices);  
numoctaves = 5;  
scales = a0.^(numvoices:numvoices*numoctaves).*1/Fs;
```

Convert the scales to approximate frequencies in hertz for the Morlet wavelet.

```
Frq = centfrq('morl')./scales;
```

You can also use `scal2frq` to convert scales to approximate frequencies in hertz.

See Also

`scal2frq` | `wavefun`

Introduced before R2006a

cfs2wpt

Wavelet packet tree construction from coefficients

Syntax

Description

CFS2WPT builds a wavelet packet tree (T) and the related analyzed signal or image (X) using the following input information:

WNAME: name of the wavelet used for the analysis

SIZE_OF_DATA: size of the analyzed signal or image

TN_OF_TREE: vector containing the terminal node indices of the tree

ORDER: 2 for a signal or 4 for an image

CFS: coefficients used to reconstruct the original signal or image. *CFS* is optional. When CFS2WPT is used without the *CFS* input parameter, the wavelet packet tree structure (T) is generated, but all the tree coefficients are null (including X).

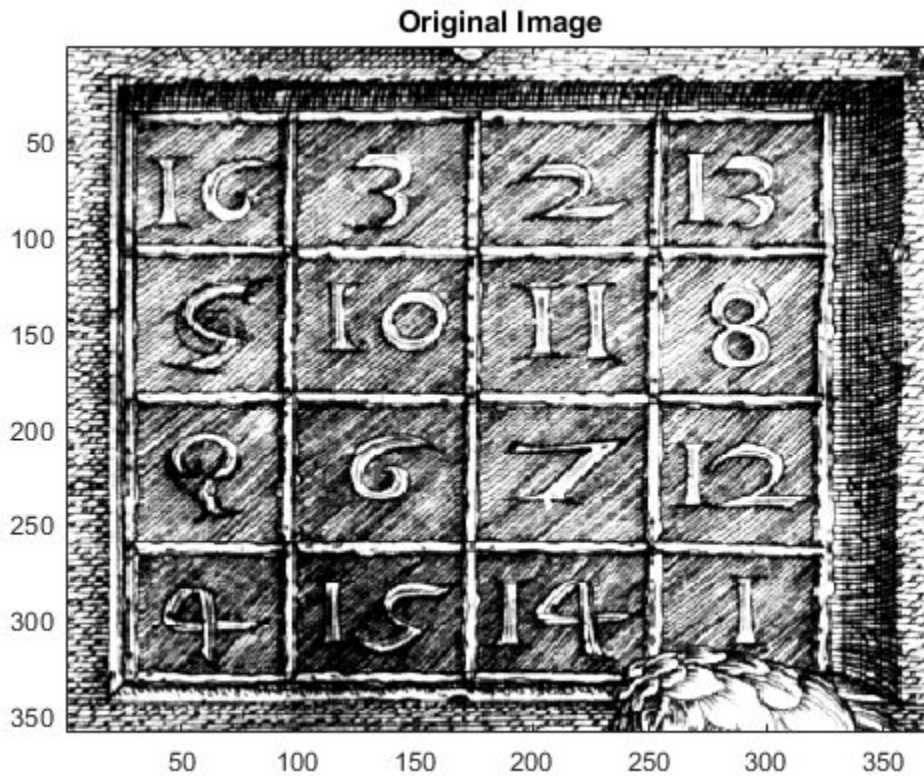
Examples

Build Wavelet Packet Tree

This example shows how to build a wavelet packet tree in two ways: 1.) By filling the wavelet packet tree with coefficients, and 2.) By creating the wavelet packet tree and using `write`

Load an image and obtain the wavelet packet decomposition down to level 2 with the 'sym4' wavelet.

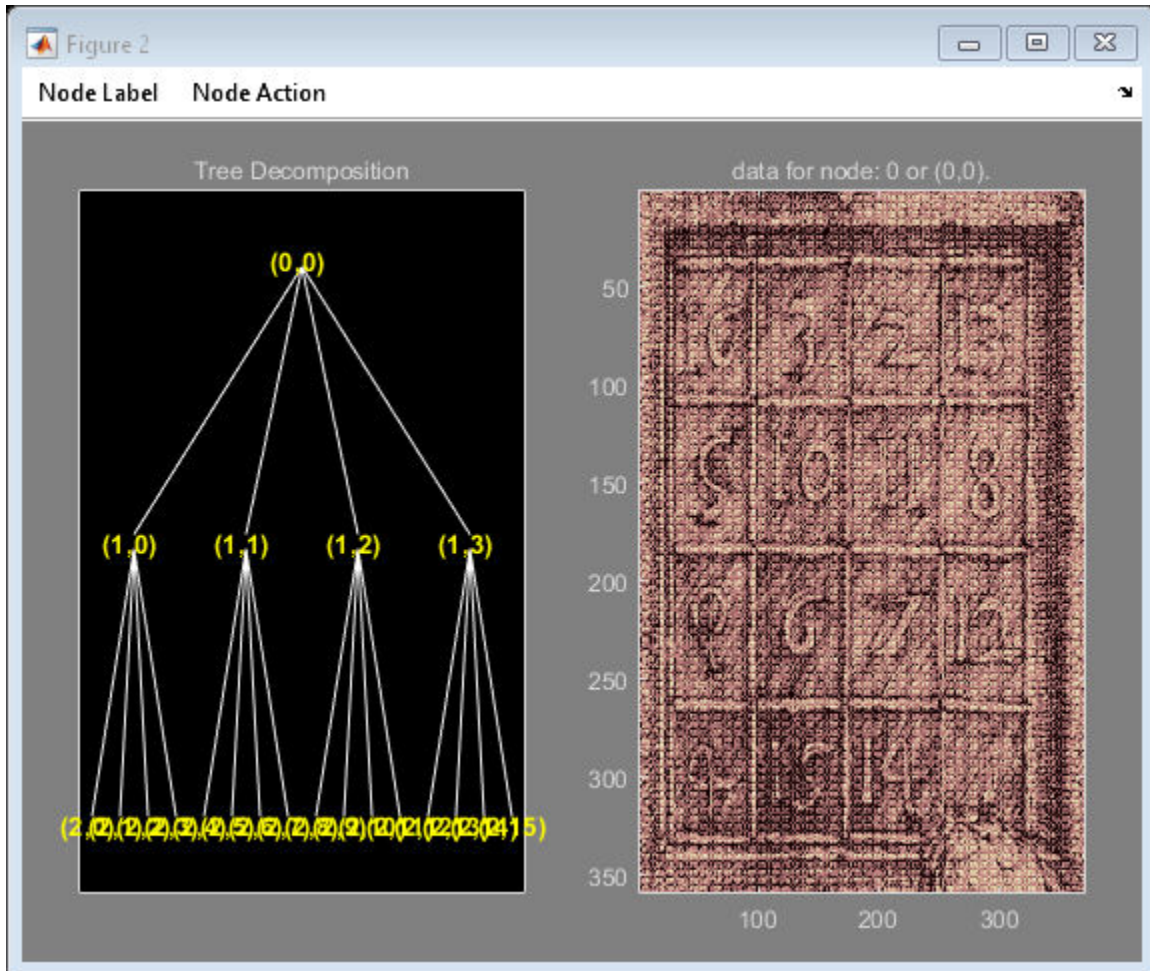
```
load detail;  
imagesc(X); colormap gray; title('Original Image');
```



```
Tr = wptdec2(X,2,'sym4');
```

Read the coefficients from the wavelet packet tree. Add $N(0, 40^2)$ noise to the coefficients and plot the new wavelet packet tree.

```
cfs = read(Tr,'allcfs');
noisyCfs = cfs + 40*rand(size(cfs));
noisyT = cfs2wpt('sym4',size(X),tnodes(Tr),4,noisyCfs);
plot(noisyT)
```



To illustrate building a wavelet packet tree using `write`, construct an admissible binary wavelet packet tree with terminal nodes [2 3 9 10]. The analyzing wavelet is 'sym4' and the signal length is 1024.

```
tr = cfs2wpt('sym4',[1 1024],[2 3 9 10'],2);
```

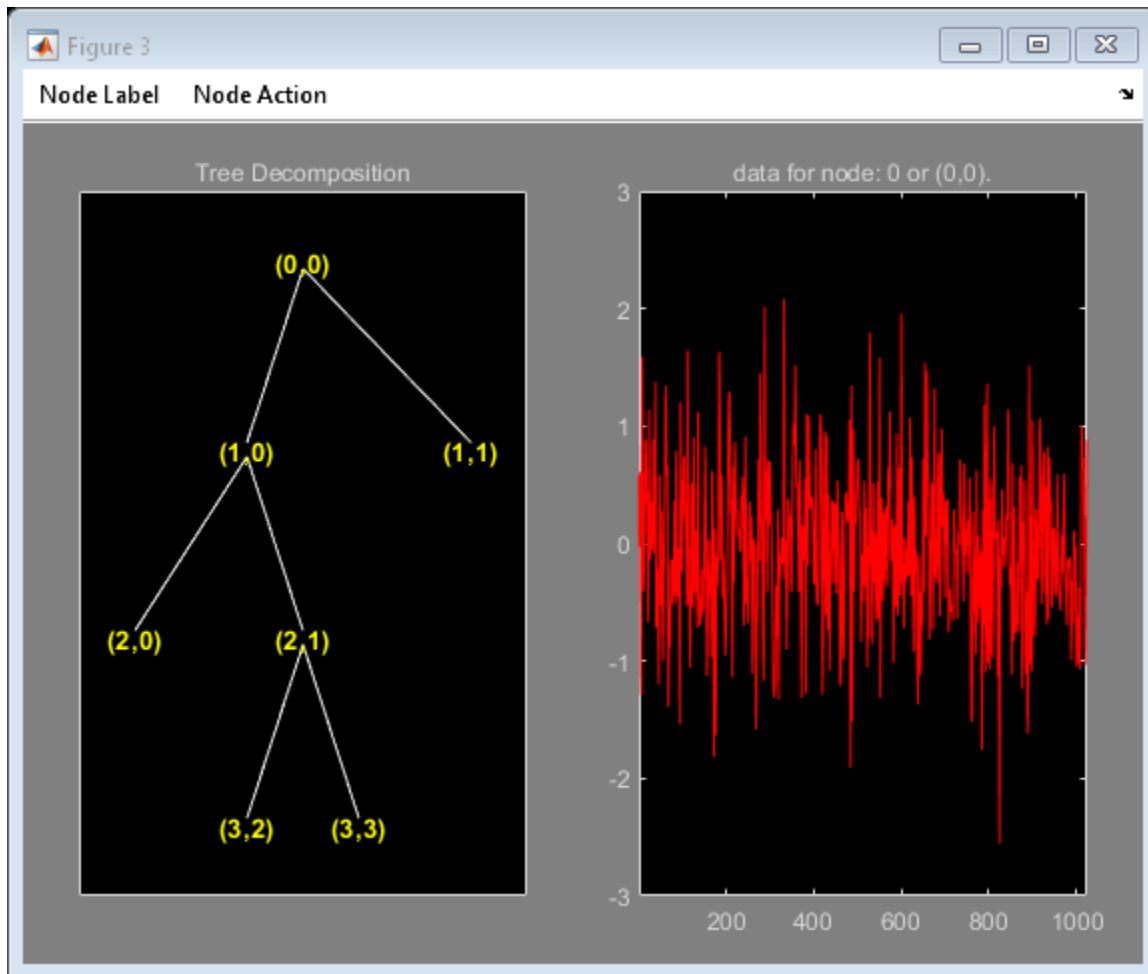
Fill terminal nodes [3 9] with $N(0, 1)$ coefficients.

```
sN = read(tr,'sizes',[3,9]);
sN3 = sN(1,:); sN9 = sN(2,:);
```

```
cfsN3 = randn(sN3);  
cfsN9 = randn(sN9);  
tr = write(tr, 'cfs', 3, cfsN3, 'cfs', 9, cfsN9);
```

Plot the resulting wavelet packet tree and synthesized signal.

```
plot(tr)
```



Introduced before R2006a

cgauwavf

Complex Gaussian wavelet

Syntax

```
[psi,x] = cgauwavf(lb,ub,n)
[psi,x] = cgauwavf(lb,ub,n,p)
[psi,x] = cgauwavf(lb,ub,n,wname)
```

Description

`[psi,x] = cgauwavf(lb,ub,n)` returns the 1st order derivative of the complex-valued Gaussian wavelet, `psi`, on an `n`-point regular grid, `x`, for the interval `[lb,ub]`. The effective support of the complex-valued Gaussian wavelets is `[-5, 5]`.

`[psi,x] = cgauwavf(lb,ub,n,p)` returns the `p`th derivative. `p` is an integer from 1 through 8.

The complex Gaussian function is defined as $C_p e^{-ix} e^{-x^2}$. C_p is such that the 2-norm of the `p`th derivative of `psi` is equal to 1.

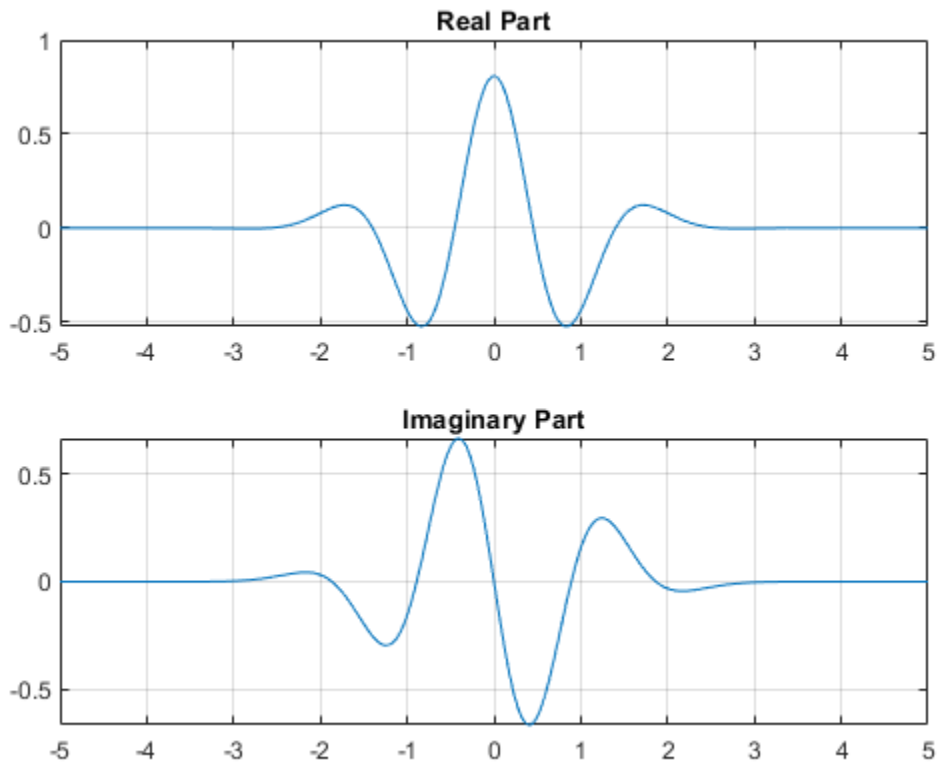
`[psi,x] = cgauwavf(lb,ub,n,wname)` used the valid wavelet family short name `wname` plus the order of the derivative in a character vector or string scalar, such as `'cgau4'`. To see valid character vectors for complex-valued Gaussian wavelets, use `waveinfo('cgau')` or use `wavemngr('read',1)` and refer to the Complex Gaussian section.

Examples

Create Complex Gaussian Wavelet

This example shows how to create a complex-valued Gaussian wavelet of order 4. The wavelet has an effective support of `[-5,5]` and is constructed using 1,000 samples.

```
lb = -5;  
ub = 5;  
n = 1000;  
order = 4;  
[psi,x] = cgauwavf(lb,ub,n,order);  
subplot(2,1,1)  
plot(x,real(psi))  
title('Real Part')  
grid on  
subplot(2,1,2)  
plot(x,imag(psi))  
title('Imaginary Part')  
grid on
```



Input Arguments

lb — Left endpoint

real number

Left endpoint of the closed interval, specified as a real number. `lb` is strictly less than `ub`.

Data Types: `double`

ub — Right endpoint

real number

Right endpoint of the closed interval, specified as a real number. `ub` is strictly greater than `lb`.

Data Types: `double`

n — Number of regularly spaced points

positive integer

Number of regularly spaced points in the interval `[lb,ub]`, specified as a positive integer. The derivative of the complex-valued Gaussian wavelet is evaluated at these points.

Data Types: `double`

p — Derivative

positive integer

Positive integer defining the order of the derivative of the complex-valued Gaussian, specified as a positive integer. `p` is an integer from 1 through 8.

wname — Gaussian wavelet

character vector | string scalar

Gaussian wavelet to evaluate, specified as a character vector or string scalar. `wname` is of the form `'cgauN'` where `N` is an integer that denotes the order of the derivative of the complex-valued Gaussian. `N` is an integer from 1 through 8.

Example: `'cgau4'` denotes the fourth derivative of the complex-valued Gaussian wavelet.

Output Arguments

psi — Derivative of complex-valued Gaussian wavelet

complex-valued vector

Derivative of the complex-valued Gaussian wavelet, returned as a complex-valued 1-by-N vector.

x — Sample points

real-valued vector

Sample points where the derivative of the complex-valued Gaussian wavelet is evaluated, returned as a real-valued 1-by-N vector. The sample points are evenly distributed between lb and ub.

See Also

`waveinfo` | `wavemngr`

Introduced before R2006a

chgwdeccfs

Change multisignal 1-D decomposition coefficients

Syntax

```
DEC = chgwdeccfs(DEC, 'ca', COEFS)
DEC = chgwdeccfs(DEC, 'cd', COEFS, LEV)
DEC = chgwdeccfs(DEC, 'all', CA, CD)
DEC = chgwdeccfs(DEC, 'all', V)
DEC = chgwdeccfs(..., IDXSIG)
```

Description

`DEC = chgwdeccfs(DEC, 'ca', COEFS)` replaces the approximation coefficients at level `DEC.level` with those contained in the matrix `COEFS`. If `COEFS` is a single value V , all coefficients are replaced by V .

`DEC = chgwdeccfs(DEC, 'cd', COEFS, LEV)` replaces the detail coefficients at level `LEV` with those contained in the matrix `COEFS`. If `COEFS` is a single value V , then `LEV` can be a vector of levels and all the coefficients that belong to these levels are replaced by V . `LEV` must be such that $1 \leq LEV \leq DEC.level$

`DEC = chgwdeccfs(DEC, 'all', CA, CD)` replaces all the approximation and detail coefficients. `CA` must be a matrix and `CD` must be a cell array of length `DEC.level`.

If `COEFS` (or `CA` or `CD`) is a single number, then it replaces all the related coefficients. Otherwise, `COEFS` (or `CA`, or `CD`) must be a matrix of appropriate size.

For a real value V , `DEC = chgwdeccfs(DEC, 'all', V)` replaces all the coefficients by V .

`DEC = chgwdeccfs(..., IDXSIG)` replaces the coefficients for the signals whose indices are given by the vector `IDXSIG`. If the initial data are stored row-wise or column-wise in a matrix `X`, then `IDXSIG` contains the row or column indices, respectively, of the data.

Examples

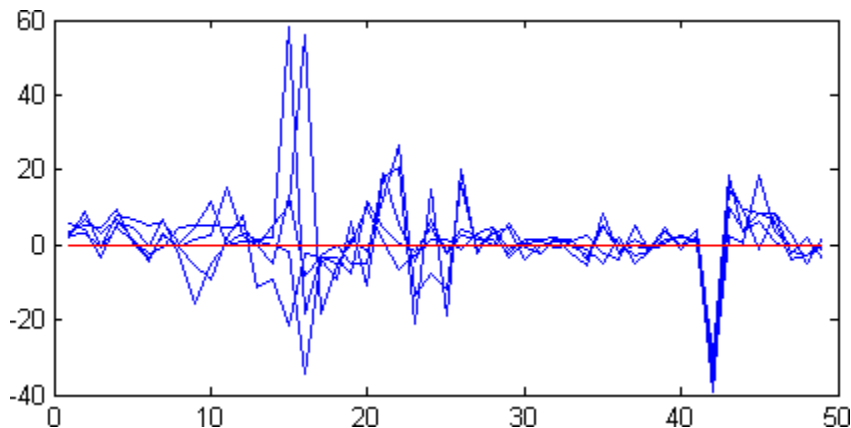
```
% Load original 1D-multisignal
load thinker

% Perform a decomposition at level 2 using wavelet db2
dec = mdwtdec('r',X,2,'db2');

% Change the coefficients of details at level 1.
% Replace all values by 0.
decBIS = chgwdeccfs(dec,'cd',0,1);

% Change the coefficients of details at level 1 and
% level 2 for signals 31 to 35. Replace all values by 0.
decTER = chgwdeccfs(dec,'cd',0,1:2,31:35);

% Compare original and new coefficients for details
% at level 1 for signals 31 to 35.
plot(dec.cd{1}(31:35,:),'b'); hold on;
plot(decTER.cd{1}(31:35,:),'r')
```



See Also

mdwtdec | mdwtrec

Introduced in R2007a

cmddenoise

Interval-dependent denoising

Syntax

```
sigden = cmddenoise(sig,wname,level)
sigden = cmddenoise(sig,wname,level,sorh)
sigden = cmddenoise(sig,wname,level,sorh,nb_inter)
sigden = cmddenoise(sig,wname,level,sorh,nb_inter,thrParamsIn)

[sigden,coefs] = cmddenoise(____)
[sigden,coefs,thrParamsOut] = cmddenoise(____)

[sigden,coefs,thrParamsOut,int_DepThr_Cell] = cmddenoise(sig,wname,
level,sorh,nb_inter)
[sigden,coefs,thrParamsOut,int_DepThr_Cell,BestNbofInt] =
cmddenoise(sig,wname,level,sorh,nb_inter)
```

Description

`sigden = cmddenoise(sig,wname,level)` returns the denoised signal, `sigden`, obtained from an interval-dependent denoising of the signal, `sig`, using the orthogonal or biorthogonal wavelet and scaling filters, `wname`. `cmddenoise` thresholds the wavelet (detail) coefficients down to `level`, `level`, and reconstructs a signal approximation using the modified detail coefficients. `cmddenoise` partitions the signal into intervals based on variance change points in the first level detail coefficients and thresholds each interval separately. The location and number of variance change points are automatically selected using a penalized contrast function [2]. The minimum delay between change points is 10 samples. Thresholds are obtained using a minimax threshold rule and soft thresholding is used to modify the wavelet coefficients [1].

`sigden = cmddenoise(sig,wname,level,sorh)` returns the denoised signal, `sigden`, using the thresholding method, `sorh`, to modify the wavelet coefficients. Valid choices for `sorh` are 's' for soft thresholding or 'h' for hard thresholding.

`sigden = cmddenoise(sig,wname,level,sorh,nb_inter)` returns the denoised signal, `sigden`, with the number of denoising intervals as a positive integer between 1

and 6: $1 \leq \text{nb_inter} \leq 6$. For $\text{nb_inter} \geq 2$, `cmddenoise` estimates the location of the change points with a contrast function [2].

`sigden = cmddenoise(sig,wname,level,sorh,nb_inter,thrParamsIn)` returns the denoised signal, `sigden`, with the denoising intervals and corresponding thresholds specified as a cell array of matrices with length equal to `level`. Each element of the cell array contains the interval and threshold information for the corresponding level of the wavelet transform. The elements of `thrParamsIn` are N-by-3 matrices with N equal to the number of intervals. The 1st and 2nd columns contain the beginning and ending indices of the intervals and the 3rd column contains the corresponding threshold value. If you specify `thrParamsIn`, `cmddenoise` ignores the value of `nb_inter`.

`[sigden,coefs] = cmddenoise(___)` returns the approximation (scaling) and detail (wavelet) coefficients, `coefs`. The organization of `coefs` is identical to the structure returned by `wavedec`. This syntax can include any of the input arguments used in previous syntaxes.

`[sigden,coefs,thrParamsOut] = cmddenoise(___)` returns a cell array, `thrParamsOut`, with length equal to `level`. Each element of `thrParamsOut` is an N-by-3 matrix. The row dimension of the matrix elements is the number of intervals and is determined by the value of the input arguments. Each row of the matrix contains the beginning and end points (indices) of the thresholded interval and the corresponding threshold value.

`[sigden,coefs,thrParamsOut,int_DepThr_Cell] = cmddenoise(sig,wname,level,sorh,nb_inter)` returns a cell array, `int_DepThr_Cell`, with length equal to 6. `int_DepThr_Cell` contains interval and threshold information assuming the number of change points ranges from 0 to 5. The N-th element of `int_DepThr_Cell` is a N-by-3 matrix containing the interval information assuming N-1 change points. Each row of the matrix contains the beginning and end points (indices) of the thresholded interval and the corresponding threshold value. Attempting to output `int_DepThr_Cell` if you use the input argument, `thrParamsIn`, results in an error.

`[sigden,coefs,thrParamsOut,int_DepThr_Cell,BestNbofInt] = cmddenoise(sig,wname,level,sorh,nb_inter)` returns the optimal number of signal intervals based on the estimated variance change points in the level-1 detail coefficients. To estimate the number of change points, `cmddenoise` assumes the total number is less than or equal to 6 and uses a penalized contrast [2]. Attempting to output `BestNbofInt` if you use the input argument, `thrParamsIn`, results in an error.

Examples

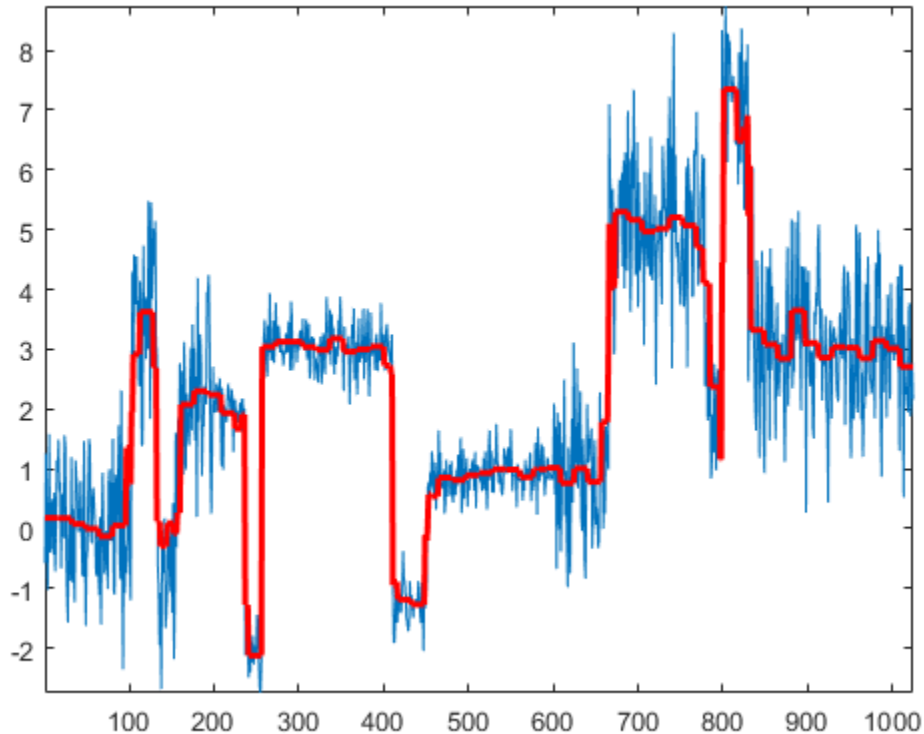
Denoising Blocks Signal with Haar Wavelet

Load the noisy blocks signal, `nblocr1.mat`. The signal consists of a piecewise constant signal in additive white Gaussian noise. The variance of the additive noise differs in three disjoint intervals.

```
load nblocr1;
```

Apply interval-dependent denoising down to level 4 using the Haar wavelet. `cmdddenoise` automatically determines the optimal number and locations of the variance change points. Plot the denoised and original signal for comparison.

```
sigden = cmdddenoise(nblocr1, 'db1', 4);  
plot(nblocr1);  
hold on;  
plot(sigden, 'r', 'linewidth', 2);  
axis tight;
```



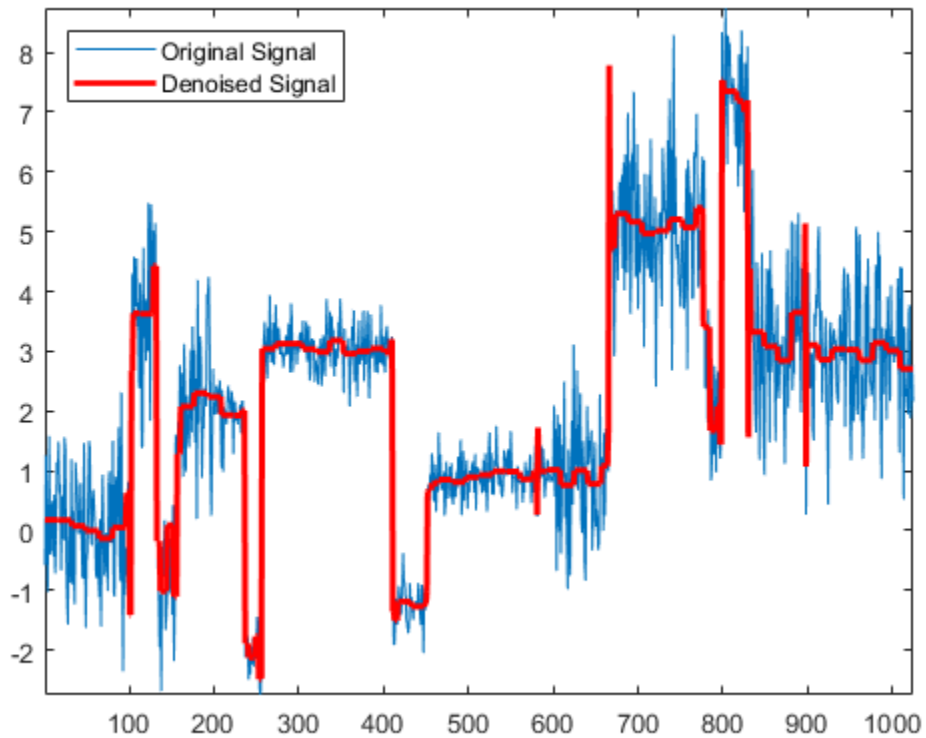
Denoising Blocks Signal with Hard Thresholding

Load the noisy blocks signal, `nblocr1.mat`. The signal consists of a piecewise constant signal in additive white Gaussian noise. The variance of the additive noise differs in three disjoint intervals.

```
load nblocr1;
```

Apply interval-dependent denoising down to level 4 using the Haar wavelet and a hard thresholding rule. `cmddenoise` automatically determines the optimal number and locations of the intervals. Plot the original and denoised signals.

```
sorh = 'h';  
sigden = cmdenoise(nblocr1,'dbl',4,sorh);  
plot(nblocr1);  
hold on;  
plot(sigden,'r','linewidth',2);  
axis tight;  
legend('Original Signal','Denoised Signal','Location','NorthWest');
```



Specify the Number of Intervals

Create a signal sampled at 1 kHz. The signal consists of a series of bumps of various widths.

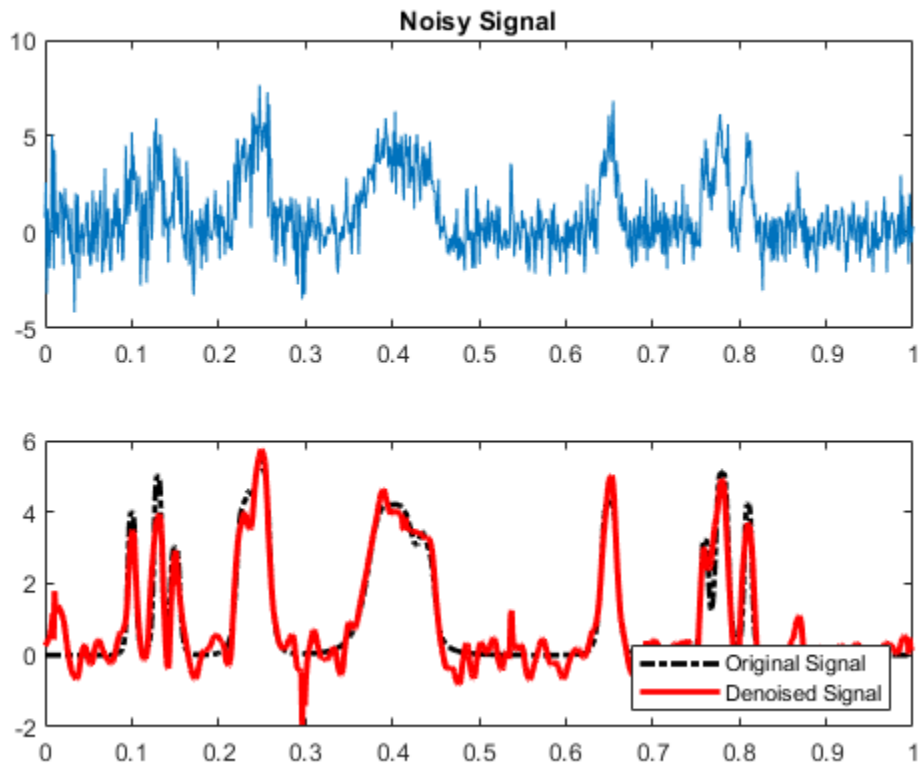
```
t = [0.1 0.13 0.15 0.23 0.25 0.40 0.44 0.65 0.76 0.78 0.81];
h = [4 -5 3 -4 5 -4.2 2.1 4.3 -3.1 5.1 -4.2];
h = abs(h);
len = 1000;
w = 0.01*[0.5 0.5 0.6 1 1 3 1 1 0.5 0.8 0.5];
tt = linspace(0,1,len);
x = zeros(1,len);
for j=1:11
    x = x + ( h(j) ./ (1+ ((tt-t(j))/w(j)).^4));
end
```

Add white Gaussian noise with different variances to two disjoint segments of the signal. Add zero-mean white Gaussian noise with variance equal to 2 to the signal segment from 0 to 0.3 seconds. Add zero-mean white Gaussian noise with unit variance to the signal segment from 0.3 seconds to 1 second. Set the random number generator to the default settings for reproducible results.

```
rng default;
nv1 = sqrt(2).*randn(size(tt)).*(tt<=0.3);
nv2 = randn(size(tt)).*(tt>0.3);
xx = x+nv1+nv2;
sigden = cmdddenoise(xx,'sym5',5,'s',2);
```

Apply interval-dependent denoising using the Daubechies' least-asymmetric wavelet with 5 vanishing moments down to level 3. Set the number of intervals to 2. Plot the noisy signal, original signal, and denoised signal for comparison.

```
sigden = cmdddenoise(xx,'sym5',3,'s',2);
subplot(211)
plot(tt,xx); title('Noisy Signal');
subplot(212)
plot(tt,x,'k-.','linewidth',2);
hold on;
plot(tt,sigden,'r','linewidth',2);
legend('Original Signal','Denoised Signal','Location','SouthEast');
```



Specify Intervals and Thresholds

Load the example signal `nbump1.mat`. The variance of the additive noise differs in three disjoint intervals.

```
load nbump1.mat;
```

Use a level-5 multiresolution analysis. Create a cell array of length 5 consisting of 3-by-3 matrices. The first two elements of each row contain the beginning and ending indices of the interval and the last element of each row is the corresponding threshold.

```

wname = 'sym4';
level = 5;
sorh = 's';
thrParamsIn = {...
    [...
    1     207     1.0482; ...
    207   613     2.5110; ...
    613  1024     1.0031; ...
    ]; ...
    [...
    1     207     1.04824; ...
    207   613     3.8718; ...
    613  1024     1.04824; ...
    ]; ...
    [...
    1     207     1.04824; ...
    207   613     1.99710; ...
    613  1024     1.65613; ...
    ]; ...
    [...
    1     207     1.04824; ...
    207   613     2.09117; ...
    613  1024     1.04824; ...
    ]; ...
    [...
    1     207     1.04824; ...
    207   613     1.78620; ...
    613  102     1.04824; ...
    ]; ...
    };

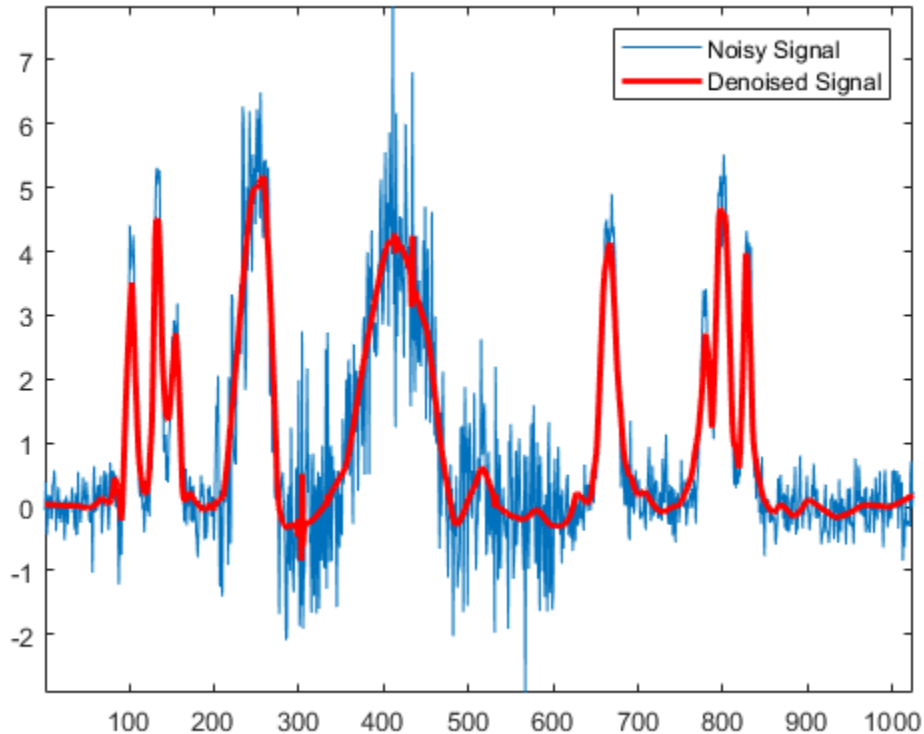
```

Denoise the signal using the threshold settings and the Daubechies' least-asymmetric wavelet with 4 vanishing moments. Use a soft thresholding rule. Plot the noisy and denoised signals for comparison.

```

wname = 'sym4';
level = 5;
sorh = 's'; sigden = cmdddenoise(nbumpr1,wname,level,sorh,...
    NaN,thrParamsIn);
plot(nbumpr1); hold on;
plot(sigden,'r','linewidth',2); axis tight;
legend('Noisy Signal','Denoised Signal','Location','NorthEast');

```



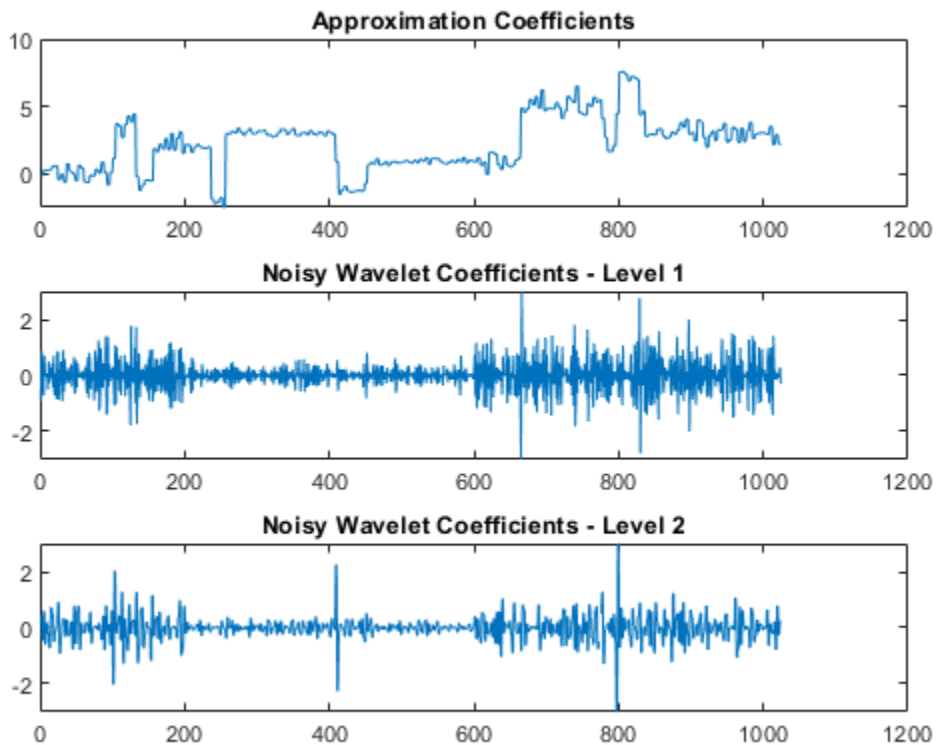
Return Denoised Wavelet Coefficients

Load the example signal `nblocr1.mat`. Use the Haar wavelet and decompose the signal down to level 2. Obtain the discrete wavelet transform and denoise the signal. Return the wavelet coefficients of the noisy and denoised signals.

```
load nblocr1.mat;  
[sigden,coefs] = cmdenoise(nblocr1,'db1',2);  
[C,L] = wavedec(nblocr1,2,'db1');
```

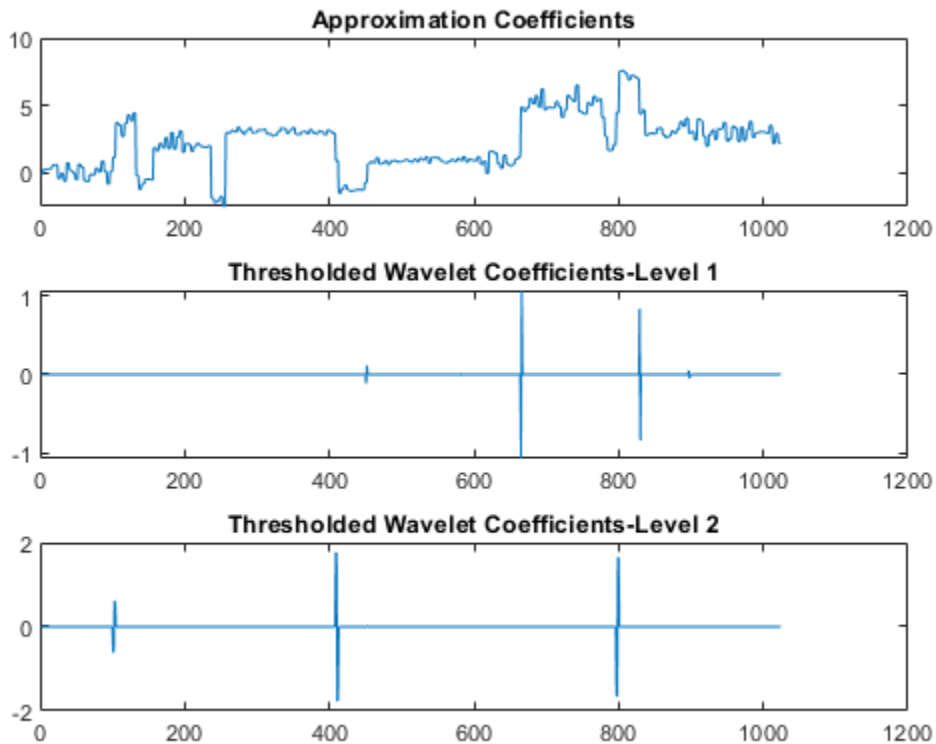

Plot reconstructions based on the level-2 approximation and level-2 and level-1 detail coefficients for the noisy signal.

```
app = wrcoef('a',C,L,'db1',2);  
subplot(3,1,1);  
plot(app); title('Approximation Coefficients');  
for nn = 1:2  
    det = wrcoef('d',C,L,'db1',nn);  
    subplot(3,1,nn+1)  
    plot(det); title(['Noisy Wavelet Coefficients - Level '...  
        num2str(nn)]);  
end
```



Plot reconstructions based on the approximation and detail coefficients for the denoised signal at the same levels.

```
figure;  
app = wrcoef('a',coefs,L,'db1',2);  
subplot(3,1,1);  
plot(app); title('Approximation Coefficients');  
for nn = 1:2  
    det = wrcoef('d',coefs,L,'db1',nn);  
    subplot(3,1,nn+1)  
    plot(det);  
    title(['Thresholded Wavelet Coefficients-Level '...  
          num2str(nn)]);  
end
```

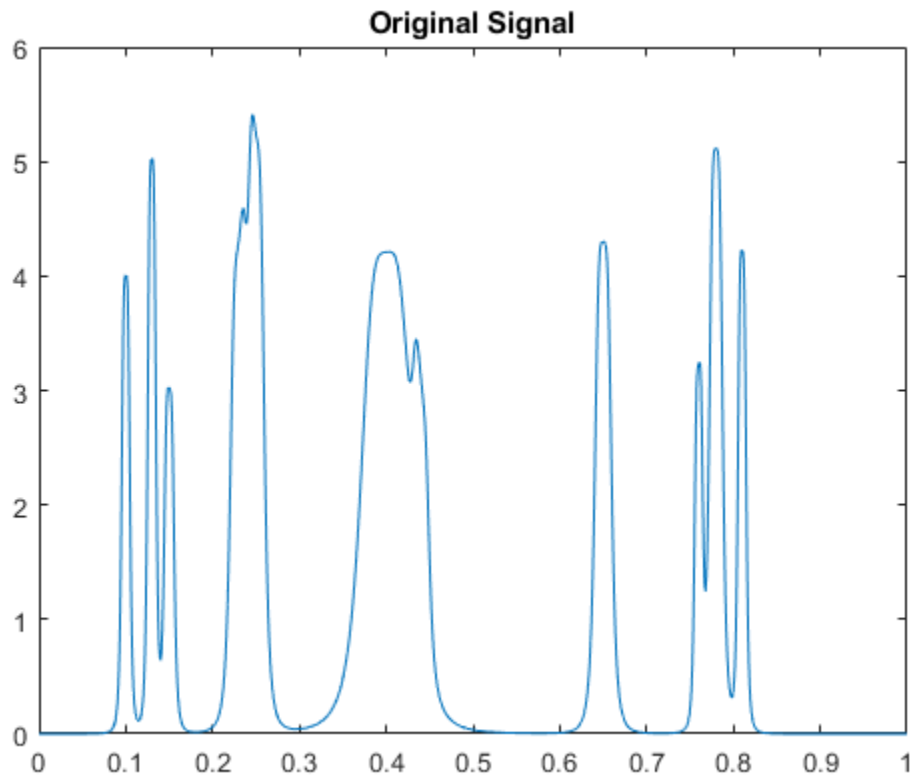


The approximation coefficients are identical in the noisy and denoised signal, but most of the detail coefficients in the denoised signal are close to zero.

Output Intervals and Thresholds

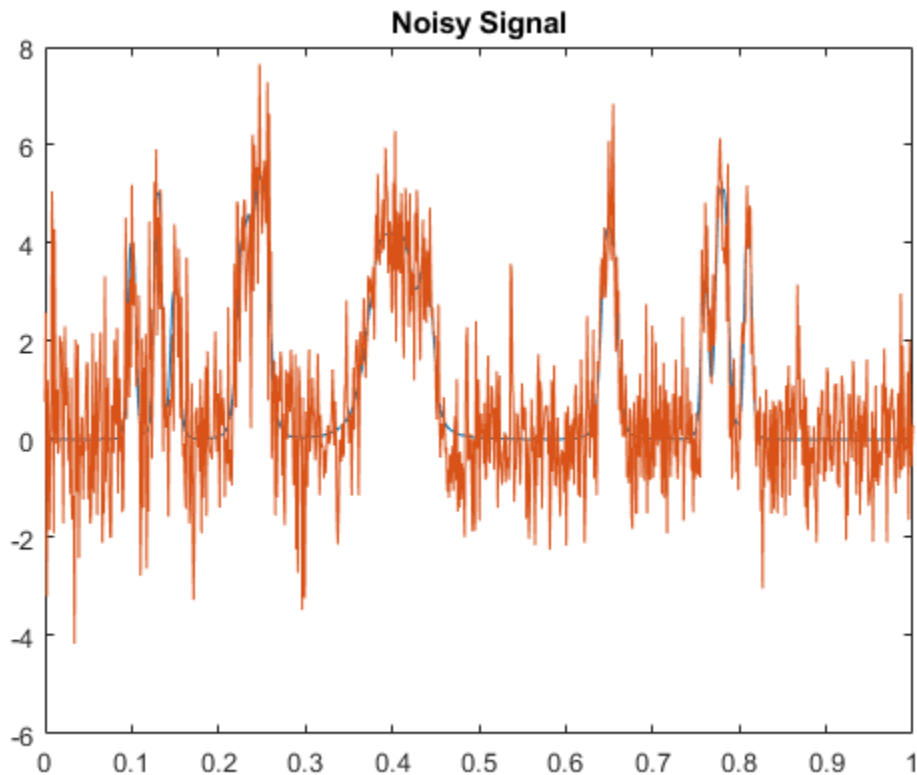
Create a signal sampled at 1 kHz. The signal consists of a series of bumps of various widths.

```
t = [0.1 0.13 0.15 0.23 0.25 0.40 0.44 0.65 0.76 0.78 0.81];
h = [4 -5 3 -4 5 -4.2 2.1 4.3 -3.1 5.1 -4.2];
h = abs(h);
len = 1000;
w = 0.01*[0.5 0.5 0.6 1 1 3 1 1 0.5 0.8 0.5];
tt = linspace(0,1,len); x = zeros(1,len);
for j=1:11
    x = x + ( h(j) ./ (1+ ((tt-t(j))/w(j)).^4));
end
plot(tt,x);
title('Original Signal');
hold on;
```



Add white Gaussian noise with different variances to two disjoint segments of the signal. Add zero-mean white Gaussian noise with variance equal to 2 to the signal segment from 0 to 0.3 seconds. Add zero-mean white Gaussian noise with unit variance to the signal segment from 0.3 seconds to 1 second. Set the random number generator to the default settings for reproducible results.

```
rng default;  
nv1 = sqrt(2).*randn(size(tt)).*(tt<=0.3);  
nv2 = randn(size(tt)).*(tt>0.3);  
xx = x+nv1+nv2;  
plot(tt,xx);  
title('Noisy Signal');
```



Apply interval-dependent denoising using the Daubechies' least- asymmetric wavelet with 4 vanishing moments down to level 5. Automatically choose the number of intervals and output the result.

```
[sigden,coefs,thrParamsOut] = cmddenoise(xx,'sym4',5);
thrParamsOut{1}
```

```
ans = 2x3
```

```
103 ×
```

```
0.0010    0.2930    0.0036
0.2930    1.0000    0.0028
```

`cmdnoise` identifies one variance change point in the 1st level detail coefficients defining two intervals. The first interval contains samples 1 to 293. The second interval contains samples 293 to 1000. This is close to the true variance change point, which occurs at sample 299.

Partition Signal into Increasing Numbers of Intervals with Thresholds

Load the example signal, `nbump1.mat`. Partition the signal into 1 to 6 intervals assuming 0 to 5 change points. Compute the thresholds for each interval. Using the Daubechies' least-asymmetric wavelet with 4 vanishing moments return the intervals and corresponding thresholds. Display the results.

```
load nbump1.mat;
[sigden,~,~,int_DepThr_Cell] = cmdnoise(nbump1,'sym4',1);
format bank;
disp('          Begin          End          Threshold ');
          Begin          End          Threshold
cellfun(@disp,int_DepThr_Cell,'UniformOutput',false);
```

1.00	1024.00	1.36
1.00	613.00	1.73
613.00	1024.00	1.00
1.00	207.00	1.05
207.00	613.00	2.51
613.00	1024.00	1.00
1.00	207.00	1.05
207.00	597.00	2.52
597.00	627.00	1.69
627.00	1024.00	0.97
1.00	207.00	1.05
207.00	613.00	2.51
613.00	695.00	1.20
695.00	725.00	0.59
725.00	1024.00	1.05
1.00	207.00	1.05

207.00	597.00	2.52
597.00	627.00	1.69
627.00	695.00	1.19
695.00	725.00	0.59
725.00	1024.00	1.05

Detect Number of Change Points

Load the example signal, `nbump1.mat`. The signal has two variance change points, which results in three intervals. Use `cmdddenoise` to detect the number of change points.

```
load nbump1.mat;
[sigden,~,thrParamsOut,~,bestNbofInt] = ...
    cmdddenoise(nbump1,'sym4',1);
fprintf('Found %d change points.\n',bestNbofInt-1);
```

Found 2 change points.

Input Arguments

sig — Signal for interval-dependent denoising

1-D row or column vector

Input signal, specified as a 1-D row or column vector. `sig` is the real-valued input signal for interval-dependent denoising. The elements of `sig` are assumed to be equally spaced in time or space. If `sig` contains unequally-sampled data, `cmdddenoise` is not appropriate. Use a lifting transform instead. See `lwt` for details.

Data Types: double

wname — Wavelet name

character vector | string scalar

Wavelet name, specified as a character vector or string scalar. `wname` is any valid orthogonal or biorthogonal wavelet. You can use the command: `wtype = wavemngr('fields',wname,'type','file');` to determine if the wavelet name is valid to use with `cmdddenoise`. Valid wavelet names return a 1 or 2 for `wtype`.

Example: `'bior2.2', 'db4', 'sym4'`

Data Types: `char`

level — Level of the decimated wavelet transform (multiresolution analysis)

positive integer

Wavelet transform (multiresolution analysis) level, specified as a positive integer. `level` gives the level of the multiresolution decomposition of the input signal using the decimated 1-D discrete wavelet transform, `wavedec`.

Data Types: `double`

sorh — Threshold rule

's' (default) | 'h'

Thresholding rule, specified as a character array. `sorh` is the threshold rule used in the modification of the detail coefficients. Valid choices for `sorh` are 's' (default) and 'h' for soft and hard thresholding.

nb_inter — Number of intervals

positive integer in the set {1,2,3,4,5,6} | NaN

Number of intervals, specified as a positive integer less than 7. `cmddenoise` divides the input signal into `nb_inter` intervals. `cmddenoise` determines the location of the `nb_inter` change points using a contrast function [2]. If you enter NaN for `nb_inter`, `cmddenoise` ignores the input. If you use the input argument `thrParamsIn`, `cmddenoise` disregards any value you enter for `nb_inter`.

Data Types: `double`

thrParamsIn — Intervals and thresholds by level

cell array of matrices

Intervals and thresholds by level, specified as a cell array of matrices equal in length to `level`. Each element of `thrParamsIn` contains the interval and threshold information for the corresponding level of the multiresolution analysis. The elements of `thrParamsIn` are N-by-3 matrices with N equal to the number of intervals. The 1st and 2nd columns contain the beginning and ending indices of the intervals and the 3rd column contains the corresponding threshold value. If you specify `thrParamsIn`, you cannot specify the output arguments `int_DepThr_Cell` or `BestNbofInt`.

Data Types: `cell`

Output Arguments

sigden — Denoised signal

1-D row or column vector

`sigden` is the denoised version of the input `sig`. `sigden` is a 1-D row vector equal in length to `sig`.

coefs — Approximation coefficients and thresholded wavelet coefficients

1-D row vector of approximation coefficients and thresholded wavelet coefficients

`coefs` is a row vector of approximation (scaling) and thresholded detail (wavelet) coefficients. The ordering of the approximation and detail coefficients by level in `coefs` is the same as the output of `wavedec`. `cmddenoise` does not apply thresholding to the approximation coefficients.

Data Types: `double`

thrParamsOut — Intervals and thresholds by level

cell array of matrices

`thrParamsOut` is a cell array of matrices equal in length to `level`. Each element of the cell array contains the interval and threshold information for the corresponding level of the multiresolution analysis. The elements of `thrParamsOut` are N-by-3 matrices with N equal to the number of intervals. N is determined by the value of the input arguments. The 1st and 2nd columns contain the beginning and ending indices of the intervals and the 3rd column contains the corresponding threshold value.

Data Types: `cell`

int_DepThr_Cell — Intervals and thresholds assuming 0 to 5 change points

cell array of matrices

`int_DepThr_Cell` contains interval and threshold information assuming the number of change points ranges from 0 to 5. The N-th element of `int_DepThr_Cell` is a N-by-3 matrix containing the interval information assuming N-1 change points. Each row of the matrix contains the beginning and ending indices of the thresholded interval and the corresponding threshold value. Attempting to output `int_DepThr_Cell` if you input the number of intervals and thresholds, `thrParamsIn`, results in an error.

`int_DepThr_Cell{BestNbofInt}` or `int_DepThr_Cell{nb_inter}` is equal to the matrix elements of `thrParamsOut`.

Data Types: `cell`

BestNbofInt — Optimal number of intervals

positive integer ≤ 6

`BestNbofInt` is the optimal number of intervals based on estimated change points in the variance of the level-1 detail coefficients. The number and location of the change points are estimated using a penalized contrast method [2]. Attempting to output `BestNbofInt` if you input the number of intervals and thresholds, `thrParamsIn`, results in an error.

References

- [1] Donoho, D. and Johnstone, I. “Ideal spatial adaptation by wavelet shrinkage”, *Biometrika*, 1994, 81,3, 425-455.
- [2] Lavielle, M. “Detection of multiple changes in a sequence of dependent variables”, *Stochastic Processes and their Applications*, 1999, 83, 79-102.

See Also

Functions

`thselect` | `wavedec` | `wdenoise` | `wthresh` | `wvarchg`

Apps

Wavelet Signal Denoiser

Introduced in R2010a

cmorwavf

Complex Morlet wavelet

Syntax

```
[PSI,X] = cmorwavf(LB,UB,N)
[PSI,X] = cmorwavf(LB,UB,N,FB,FC)
```

Description

[PSI,X] = cmorwavf(LB,UB,N) returns the complex Morlet wavelet, PSI, with time-decay parameter, FB, and center frequency, FC, both equal to 1. The general expression for the complex Morlet wavelet is

$$\text{PSI}(X) = ((\pi \cdot \text{FB})^{-0.5}) \cdot \exp(2 \cdot \pi \cdot i \cdot \text{FC} \cdot X) \cdot \exp(-(X^2)/\text{FB})$$

X is evaluated on an N -point regular grid in the interval [LB,UB].

[PSI,X] = cmorwavf(LB,UB,N,FB,FC) returns values of the complex Morlet wavelet defined by a positive time-decay parameter, FB, and positive center frequency, FC.

FB controls the decay in the time domain and the corresponding energy spread (bandwidth) in the frequency domain. FB is the inverse of the variance in the frequency domain. Increasing FB makes the wavelet energy more concentrated around the center frequency and results in slower decay of the wavelet in the time domain. Decreasing FB results in faster decay of the wavelet in the time domain and less energy spread in the frequency domain. The value of FB does not affect the center frequency. When converting from scale to frequency, only the center frequency affects the frequency values. The energy spread or bandwidth parameter affects how localized the wavelet is in the frequency domain.

Examples

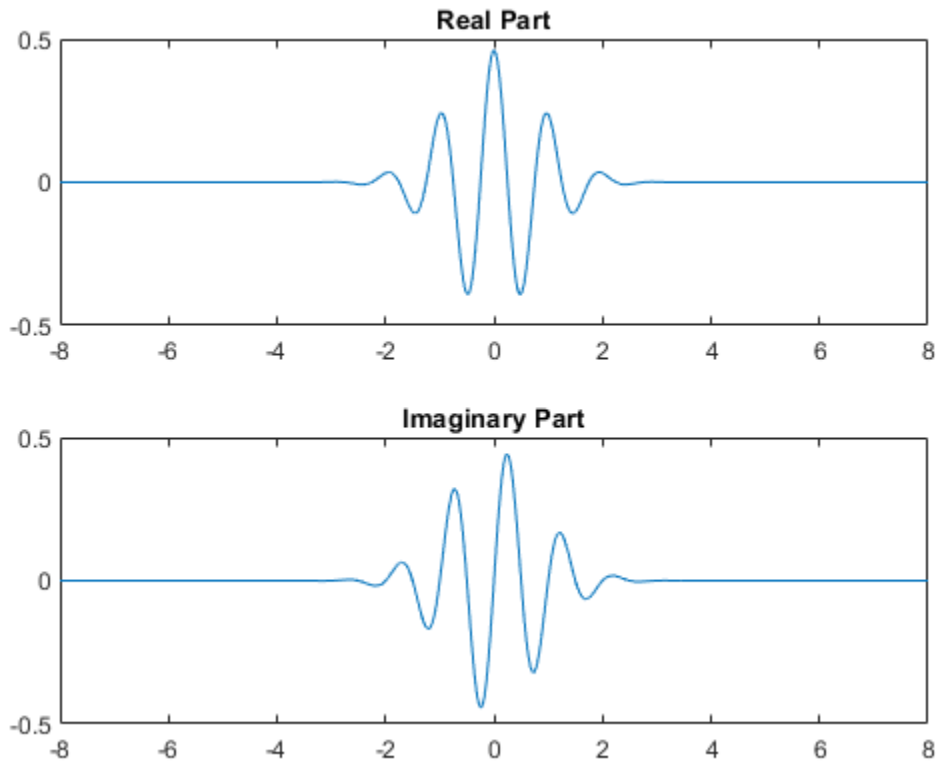
Complex Morlet Wavelet

Construct a complex-valued Morlet wavelet with a bandwidth parameter of 1.5 and a center frequency of 1. Set the effective support to $[-8, 8]$ and the length of the wavelet to 1000.

```
N = 1000;  
Lb = -8;  
Ub = 8;  
fb = 1.5;  
fc = 1;  
[psi,x] = cmorwavf(Lb,Ub,N,fb,fc);
```

Plot the real and imaginary parts of the wavelet.

```
subplot(2,1,1)  
plot(x,real(psi)); title('Real Part');  
subplot(2,1,2)  
plot(x,imag(psi)); title('Imaginary Part');
```

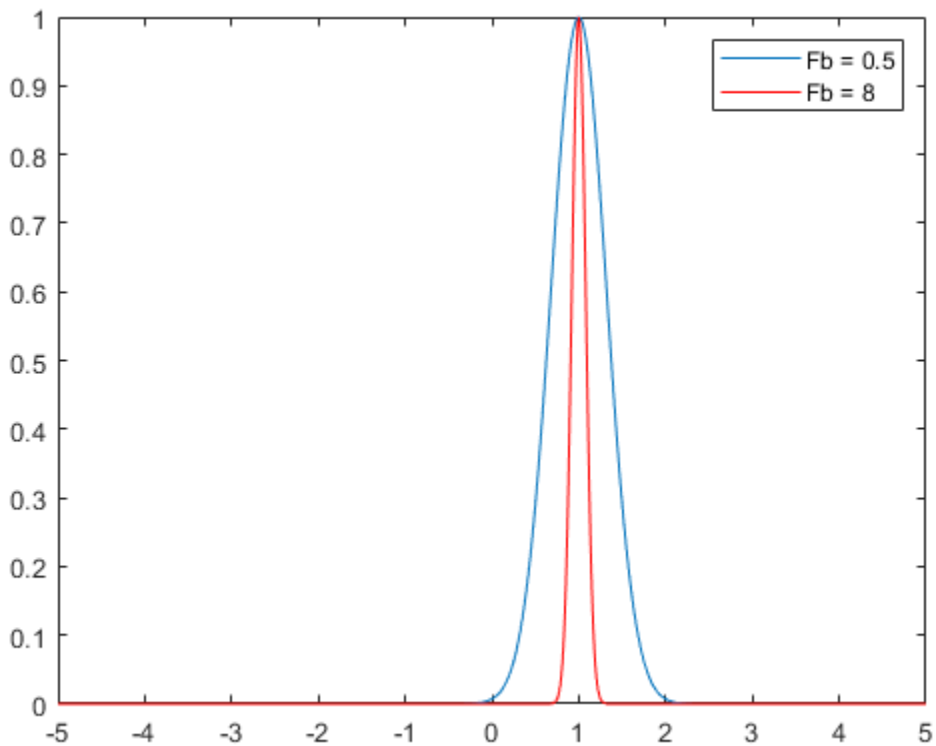


Effect of Bandwidth Parameter on Morlet Wavelet Shape

This example shows how the complex Morlet wavelet shape in the frequency domain is affected by the value of the bandwidth parameter (Fb). Both wavelets have a center frequency of 1. One wavelet has an Fb value of 0.5 and the other wavelet has a value of 8.

```
f = -5:.01:5;  
Fc = 1;  
Fb1 = 0.5;  
Fb2 = 8;  
psihat1 = exp(-pi^2*Fb1*(f-Fc).^2);
```

```
psihat2 = exp(-pi^2*Fb2*(f-Fc).^2);  
plot(f,psihat1)  
hold on;  
plot(f,psihat2,'r')  
legend('Fb = 0.5','Fb = 8')
```



The Fb bandwidth parameter for the complex Morlet wavelet is the inverse of the variance in frequency. Therefore, increasing Fb results in a narrower concentration of energy around the center frequency.

References

Teolis, A. (1998), *Computational signal processing with wavelets*, Birkhauser, p. 65.

See Also

waveinfo

Introduced before R2006a

coefficientSize

Size of image scattering coefficients

Syntax

```
sz = coefficientSize(sf)
```

Description

`sz = coefficientSize(sf)` returns the scattering coefficient sizes for the wavelet image scattering framework, `sf`. The output `sz` is a two-element row vector that gives the scattering coefficient output size in the row and column dimensions. For an RGB image, the actual output size is `[sz(1) sz(2) 3]`.

Examples

Scattering Coefficient Sizes for Image Scattering Framework

This example shows how to determine the scattering coefficient sizes of an image scattering framework.

Create a wavelet image scattering framework with an image size of 128-by-64. Obtain the coefficient sizes of the framework.

```
sf = waveletScattering2('ImageSize',[128 64]);  
sz = coefficientSize(sf)
```

```
sz = 1×2
```

```
    16     8
```

Create a second wavelet image scattering framework with an image size of 128-by-64 and an oversampling factor equal to 1. Obtain the coefficient sizes of the framework. Since the

oversampling factor is equal to 1, the scattering transform of the second framework returns 2-by-2-by- P as many coefficients for each scattering path with respect to the critically sampled number.

```
sf2 = waveletScattering2('ImageSize',[128 64],'OversamplingFactor',1);  
sz = coefficientSize(sf2)
```

```
sz = 1×2
```

```
    32    16
```

Input Arguments

sf — Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

See Also

paths | waveletScattering2

Introduced in R2019a

coifwavf

Coiflet wavelet filter

Syntax

```
f = coifwavf(wname)
```

Description

`f = coifwavf(wname)` returns the scaling filter `f` associated with the Coiflet wavelet specified by `wname`. `f` is a real-valued vector.

Examples

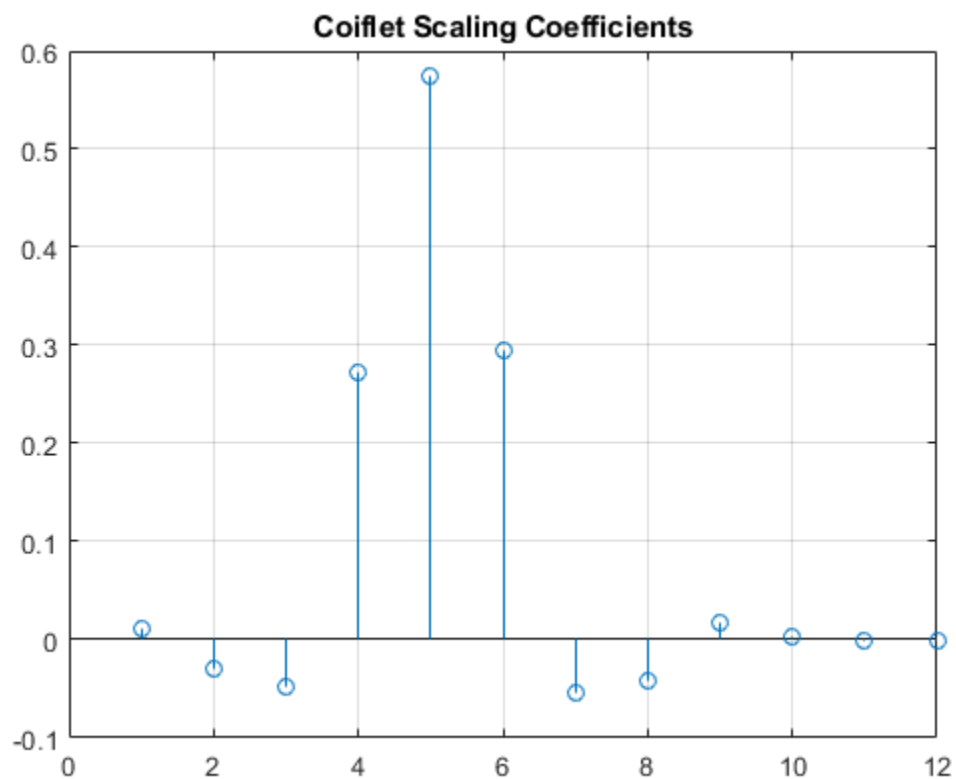
Coiflet Wavelet Filter

Set the Coiflet wavelet name.

```
wname = 'coif2';
```

Compute and plot the scaling filter coefficients associated with the Coiflet.

```
f = coifwavf(wname);  
stem(f)  
grid on  
title('Coiflet Scaling Coefficients')
```



Input Arguments

wname — Name of Coiflet

character vector | string scalar

Name of Coiflet, specified as 'coifN' where N is an integer between 1 and 5.

Example: 'coif3'

See Also

waveinfo

Introduced before R2006a

concatenate

Concatenate two or more labeled signal sets

Syntax

```
lssnew = concatenate(lss1,...,lssN)
```

Description

`lssnew = concatenate(lss1,...,lssN)` concatenates N labeled signal set objects, `lss1,...,lssN`, and returns a labeled signal set `lssnew` containing all the members and label values of the input sets.

Examples

Concatenate Labeled Signal Sets

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
      NumMembers: 2  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [2x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Create a new signal set with the same data source, time information, and labels as `lss`.

```
newlss = lss
newlss =
  labeledSignalSet with properties:
        Source: {2x1 cell}
      NumMembers: 2
    TimeInformation: "sampleRate"
      SampleRate: 4000
        Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Concatenate the two signal sets.

```
lssconcat = concatenate(lss,newlss)
lssconcat =
  labeledSignalSet with properties:
        Source: {4x1 cell}
      NumMembers: 4
    TimeInformation: "sampleRate"
      SampleRate: 4000
        Labels: [4x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Input Arguments

`lss1, ..., lssN` — Input labeled signal sets

`labeledSignalSet` objects

Input labeled signal sets, specified as `labeledSignalSet` objects. All input sets must have the same time information settings, label definitions, and data source type.

Output Arguments

lssnew — Concatenated labeled signal set

labeledSignalSet object

Concatenated labeled signal set, returned as a labeledSignalSet object. The set lssnew contains a signal source, label definitions, and label values that are independent of the input labeled signal sets.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

conofinf

Cone of influence

Syntax

```
cone = conofinf(wname,scales,LenSig,SigVal)
[cone,PL,PR] = conofinf(wname,scales,LenSig,SigVal)
[cone,PL,PR,PLmin,PRmax] = conofinf(wname,scales,LenSig,SigVal)
[PLmin,PRmax] = conofinf(wname,scales,LenSig)
[...] = conofinf(...,'plot')
```

Description

`cone = conofinf(wname,scales,LenSig,SigVal)` returns the cone of influence (COI) for the wavelet `wname` at the scales in `scales` and positions in `SigVal`. `LenSig` is the length of the input signal. If `SigVal` is a scalar, `cone` is a matrix with row dimension `length(scales)` and column dimension `LenSig`. If `SigVal` is a vector, `cone` is cell array of matrices.

`[cone,PL,PR] = conofinf(wname,scales,LenSig,SigVal)` returns the left and right boundaries of the cone of influence at scale 1 for the points in `SigVal`. `PL` and `PR` are `length(SigVal)`-by-2 matrices. The left boundaries are $(1-PL(:,2))./PL(:,1)$ and the right boundaries are $(1-PR(:,2))./PR(:,1)$.

`[cone,PL,PR,PLmin,PRmax] = conofinf(wname,scales,LenSig,SigVal)` returns the equations of the lines that define the minimal left and maximal right boundaries of the cone of influence. `PLmin` and `PRmax` are 1-by-2 row vectors where `PLmin(1)` and `PRmax(1)` are the slopes of the lines. `PLmin(2)` and `PRmax(2)` are the points where the lines intercept the scale axis.

`[PLmin,PRmax] = conofinf(wname,scales,LenSig)` returns the slope and intercept terms for the first-degree polynomials defining the minimal left and maximal right vertices of the cone of influence.

`[...] = conofinf(...,'plot')` plots the cone of influence.

Input Arguments

wname

wname is a character vector or string scalar corresponding to a valid wavelet. To verify that wname is a valid wavelet, `wavemngr('fields',wname)` must return a struct array with a `type` field of 1 or 2, or a nonempty `bound` field.

scales

scales is a vector of scales over which to compute the cone of influence. Larger scales correspond to stretched versions of the wavelet and larger boundary values for the cone of influence.

LenSig

LenSig is the signal length and must exceed the maximum of SigVal.

SigVal

SigVal is a vector of signal values at which to compute the cone of influence. The largest value of SigVal must be less than the signal length, LenSig. If SigVal is empty, conofinf returns the slope and intercept terms for the minimal left and maximal right vertices of the cone of influence.

Output Arguments

cone

cone is the cone of influence. If SigVal is a scalar, cone is a matrix. The row dimension is equal to the number of scales and column dimension equal to the signal length, LenSig. If SigVal is a vector, cone is a cell array of matrices. The elements of each row of the matrix are equal to 1 in the interval around SigVal corresponding to the cone of influence.

PL

PL is the minimum value of the cone of influence on the position (time) axis.

PR

PR is the maximum value of the cone of influence on the position (time) axis.

PLmin

PLmin is a 1-by-2 row vector containing the slope and scale axis intercept of the line defining the minimal left vertex of the cone of influence. PLmin(1) is the slope and PLmin(2) is the point where the line intercepts the scale axis.

PRmax

PRmax is a 1-by-2 row vector containing the slope and scale axis intercept of the line defining the maximal right vertex of the cone of influence. PRmax(1) is the slope and PRmax(2) is the point where the line intercepts the scale axis.

Examples

Cone of Influence for Mexican Hat or Ricker Wavelet

Load the data.

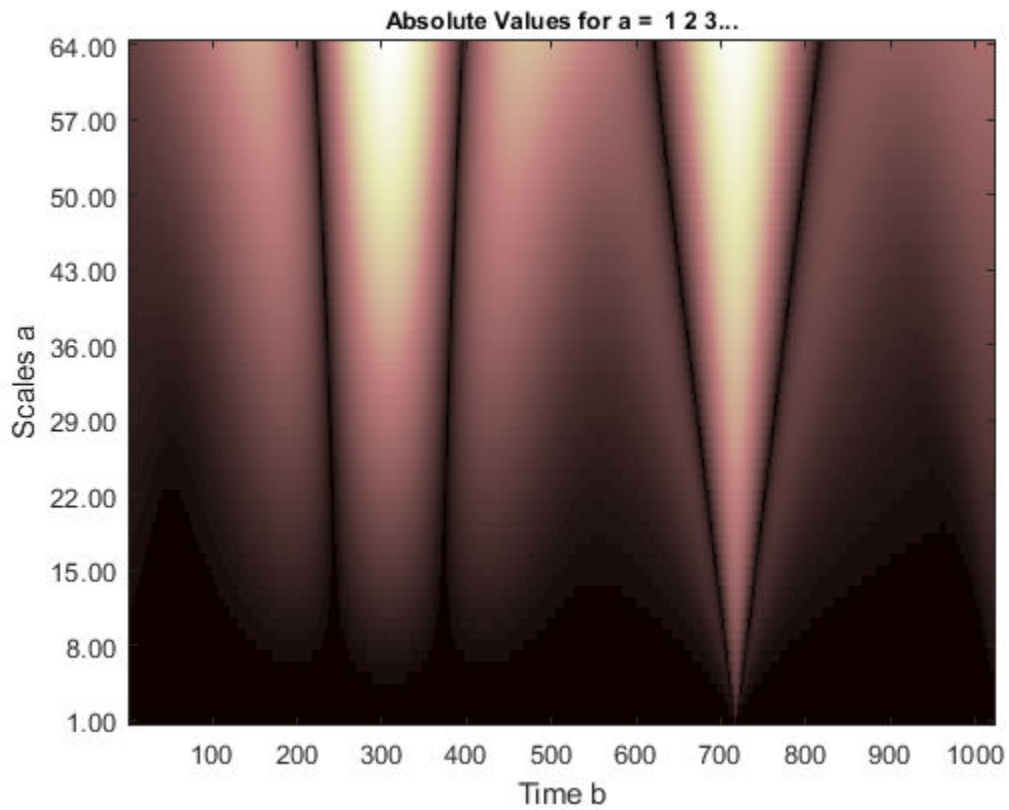
```
load cuspamax  
signal = cuspamax;
```

Set up the wavelet.

```
wname = 'mexh';  
scales = 1:64;  
lenSIG = length(signal);  
x = 500;
```

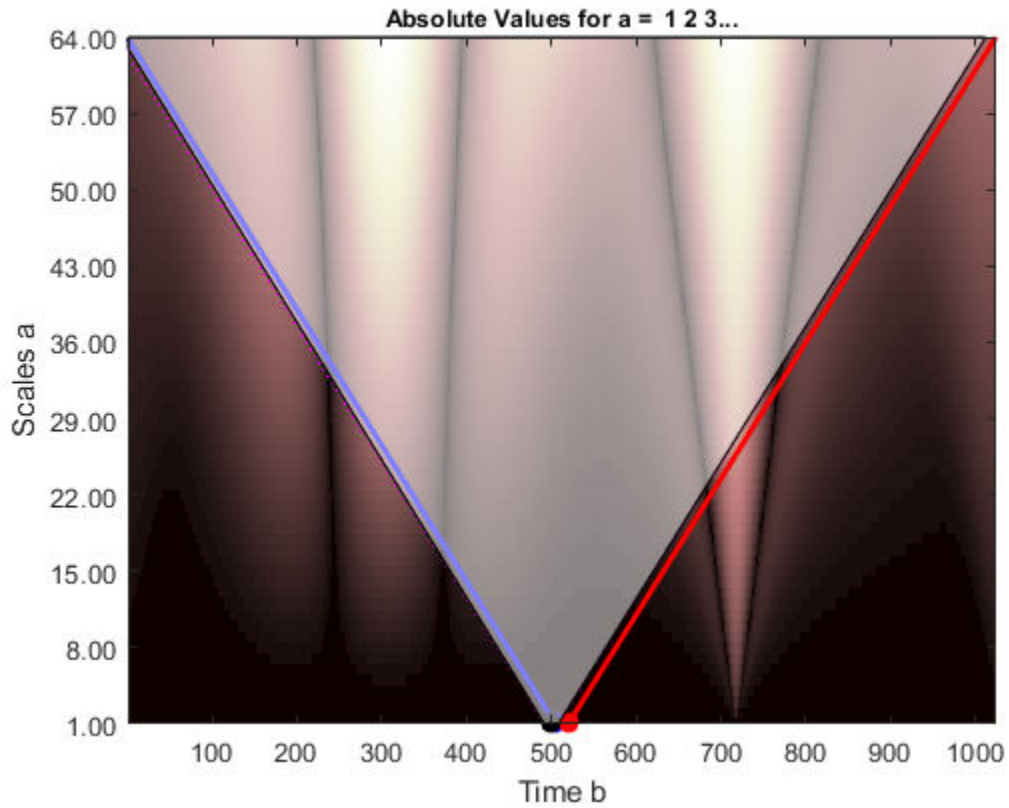
Plot the wavelet.

```
figure;  
cwt(signal,scales,wname,'plot');
```



Plot the cone of influence.

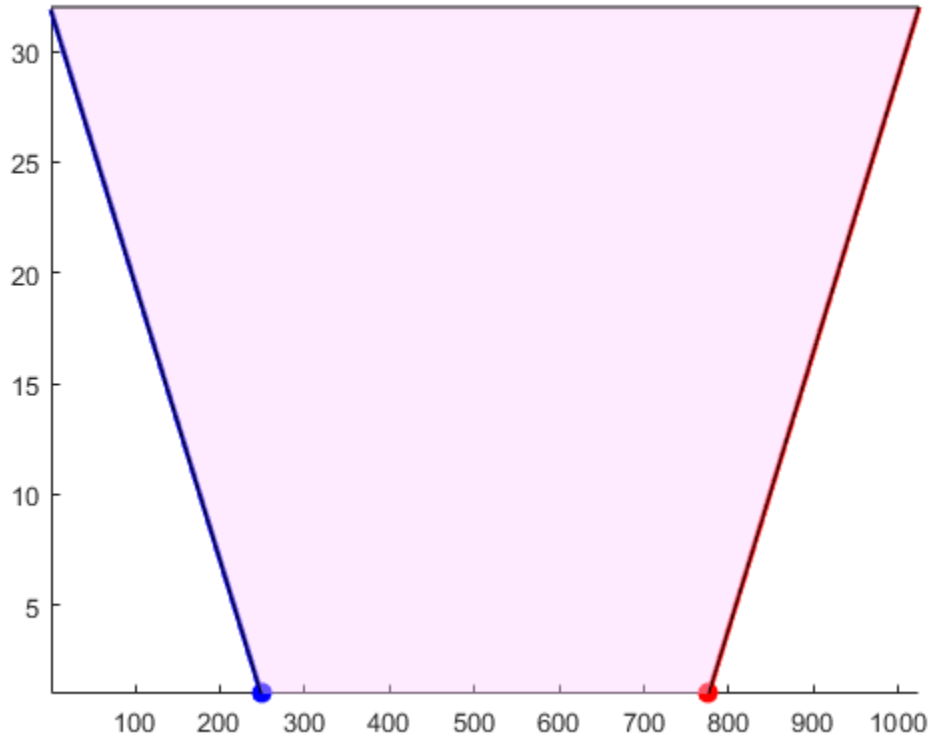
```
hold on  
[cone,PL,PR,Pmin,Pmax] = conofinf(wname,scales,lenSIG,x,'plot');  
set(gca,'Xlim',[1 lenSIG])
```



Cone of Influence Minimal and Maximal Vertices

Return the left minimal and right maximal vertices for the cone of influence (Morlet wavelet).

```
[PLmin,PRmax] = conofinf('morl',1:32,1024,[],'plot');
```



PLmin

PLmin = 1×2

-0.1245 32.0000

PRmax

PRmax = 1×2

0.1250 -96.0000

Definitions

Cone of Influence

Let $\psi(t)$ be an admissible wavelet. Assume that the effective support of $\psi(t)$ is $[-B, B]$. Letting u denote the translation parameter and s denote the scale parameter, the dilated and translated wavelet is:

$$\psi_{u,s}(t) = \frac{1}{\sqrt{s}} \psi\left(\frac{t-u}{s}\right)$$

and has effective support $[u-sB, u+sB]$. The cone of influence (COI) is the set of all t included in the effective support of the wavelet at a given position and scale. This set is equivalent to:

$$|t - u| \leq sB$$

At each scale, the COI determines the set of wavelet coefficients influenced by the value of the signal at a specified position.

References

Mallat, S. *A Wavelet Tour of Signal Processing*, London:Academic Press, 1999, p. 174.

See Also

cwt | wavsupport

Topics

“Continuous and Discrete Wavelet Transforms”

Introduced in R2010b

cqt

Constant-Q nonstationary Gabor transform

Syntax

```
cfs = cqt(x)
[cfs,f] = cqt(x)
[cfs,f,g,fshifts] = cqt(x)
[cfs,f,g,fshifts,fintervals] = cqt(x)
[cfs,f,g,fshifts,fintervals,bw] = cqt(x)
[___] = cqt( ___,Name,Value)
cqt( ___ )
```

Description

`cfs = cqt(x)` returns the constant-Q transform (CQT), `cfs`, of input signal `x`.

- If `x` is a vector, then `cqt` returns a matrix corresponding to the CQT.
- If `x` is a matrix, then `cqt` obtains the CQT for each column (independent channel) of `x`. The function returns a multidimensional array corresponding to the maximally redundant version of the CQT.

`[cfs,f] = cqt(x)` returns the approximate bandpass center frequencies, `f`, corresponding to the rows of `cfs`. The frequencies are ordered from 0 to 1 and are in cycles/sample.

`[cfs,f,g,fshifts] = cqt(x)` returns the Gabor frames, `g`, used in the analysis of `x` and the frequency shifts, `fshifts`, in discrete Fourier transform (DFT) bins between the passbands in the rows of `cfs`.

`cfs`, `g`, and `fshifts` are required inputs for the inversion of the CQT with `icqt`.

`[cfs,f,g,fshifts,fintervals] = cqt(x)` returns the frequency intervals, `fintervals`, corresponding the rows of `cfs`. The k th element of `fshifts` is the frequency shift in DFT bins between the $((k-1) \bmod N)$ and $(k \bmod N)$ element of `fintervals` with $k = 0, 1, 2, \dots, N-1$ where N is the number of frequency shifts.

Because MATLAB® indexes from 1, `fshifts(1)` contains the frequency shift between `fintervals{end}` and `fintervals{1}`, `fshifts(2)` contains the frequency shift between `fintervals{1}` and `fintervals{2}`, and so on.

`[cfs,f,g,fshifts,fintervals,bw] = cqt(x)` returns the bandwidth, `bw`, in DFT bins of the frequency intervals, `fintervals`.

`[___] = cqt(___ ,Name,Value)` returns the CQT with additional options specified by one or more `Name,Value` pair arguments, using any of the preceding syntaxes.

`cqt(___)` with no output arguments plots the CQT in the current figure. Plotting is supported for vector inputs only. If the input signal is real and `Fs` is the sampling frequency, the CQT is plotted over the range `[0, Fs/2]`. If the signal is complex, the CQT is plotted over the range `[0, Fs)`.

Examples

Constant-Q Transform Using Default Values

Load a signal and obtain the constant-Q transform.

```
load noisdopp
cfs = cqt(noisdopp);
```

Center Frequencies of the Constant-Q Transform

Load a real-valued signal and obtain the constant-Q transform. Return the approximate bandpass center frequencies.

```
load handel
[cfs,f] = cqt(y);
```

Plot on a logarithmic scale the bandpass center frequencies through the Nyquist frequency.

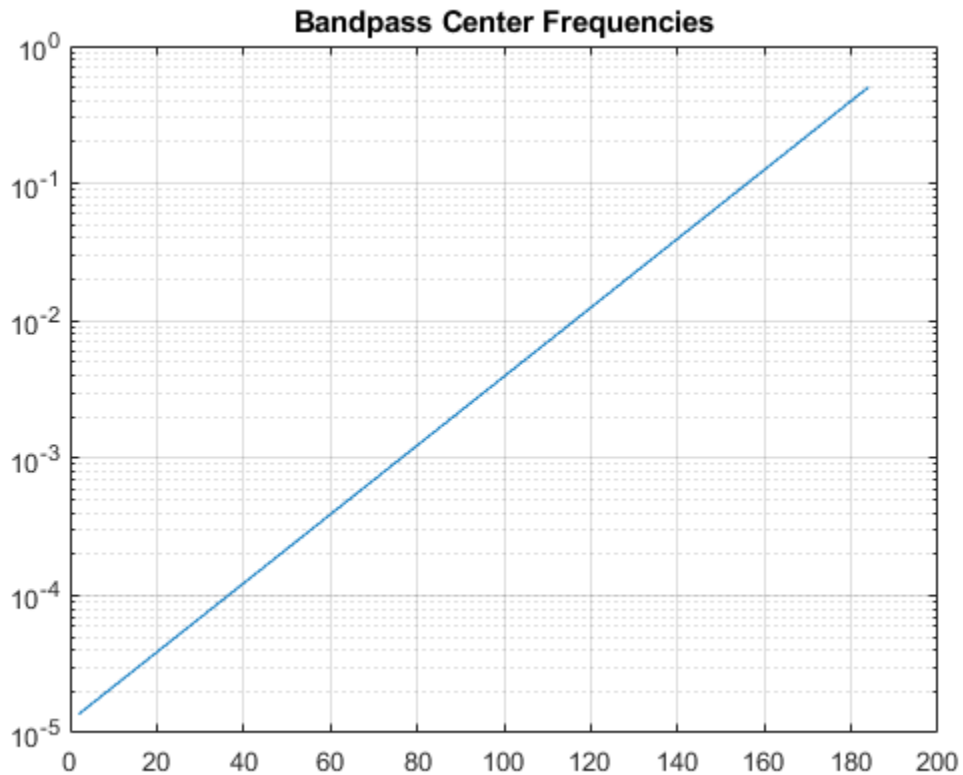
```
lfreq = length(f);
nyquistBin = floor(lfreq/2)+1;
plot(f(1:nyquistBin))
```



```

title('Bandpass Center Frequencies')
grid on
set(gca,'yscale','log')

```



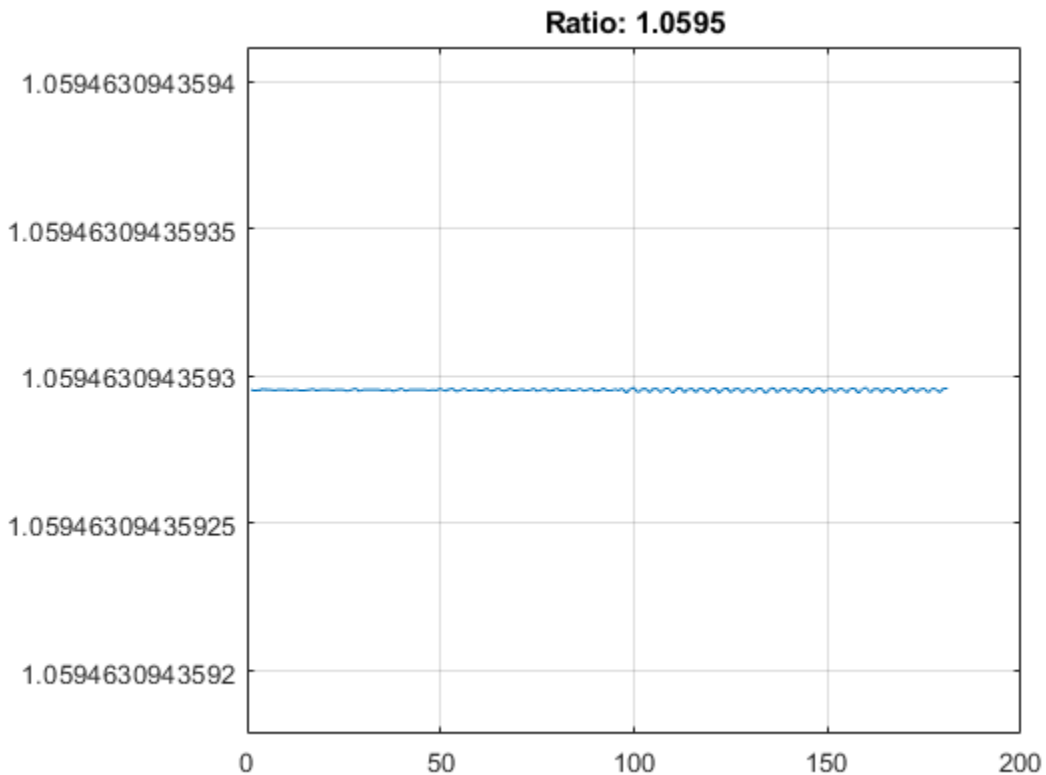
To confirm the ratios of consecutive pairs of frequencies are constant, plot the ratios.

Since `cqt` uses 12 bins per octave by default, the ratio should equal $2^{1/12}$. Since the DC and Nyquist frequencies are not members of the geometric sequence of center frequencies but are included in the frequency vector, exclude them from the plot.

```

figure
plot(f(3:nyquistBin-1)./f(2:nyquistBin-2))
grid on
title(['Ratio: ',num2str(2^(1/12))])

```



Visualize and Apply Constant-Q Transform Gabor Frames

Obtain the minimally redundant constant-Q transform of an audio signal. Use the Blackman-Harris window as the prototype function for the Gabor frames.

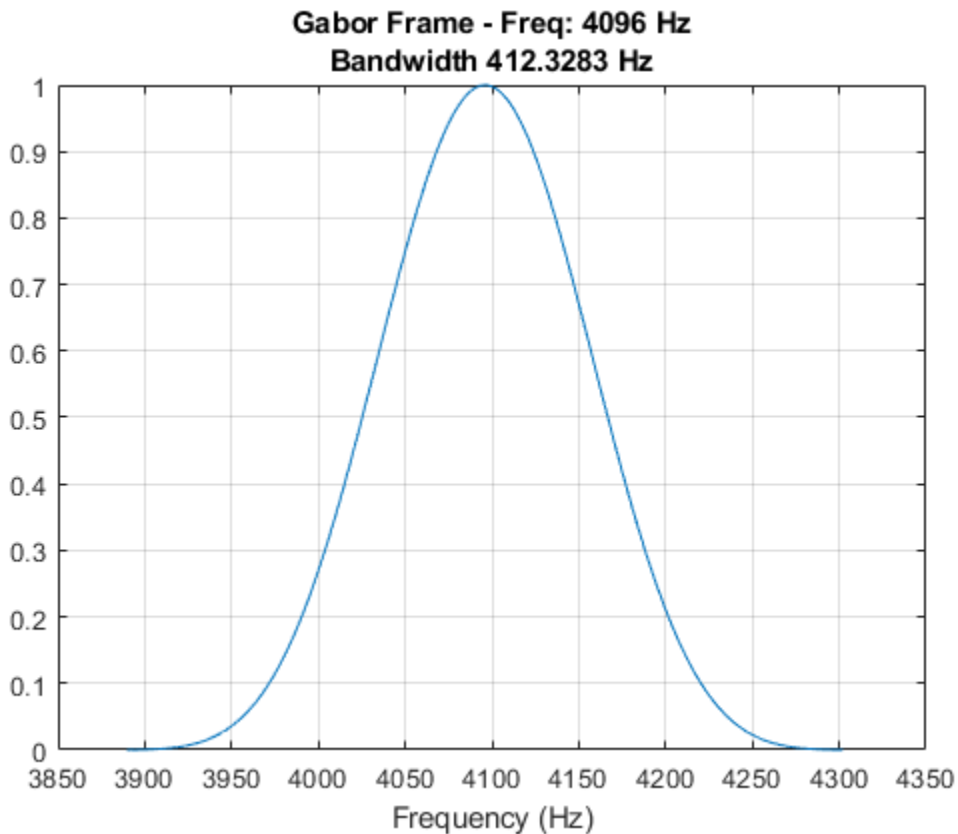
```
load handel
df = Fs/numel(y);
[cfs,f,g,fshifts,fintervals,bw] = cqt(y, 'SamplingFrequency',Fs, 'TransformType', 'sparse');
```

`cfs` is a cell array, where each element in the array corresponds to a bandpass center frequency and Gabor frame. Plot the Gabor frame associated with the Nyquist frequency.

```

lf = length(f);
ind = floor(lf/2)+1;
gFrame = fftshift(g{ind});
fvec = f(ind-1):df:f(ind+1)-df;
plot(fvec,gFrame)
xlabel('Frequency (Hz)')
grid on
title(['Gabor Frame - Freq: ',num2str(f(ind)), ' Hz'];['Bandwidth ',num2str(bw(ind)*Fs), ' Hz'])

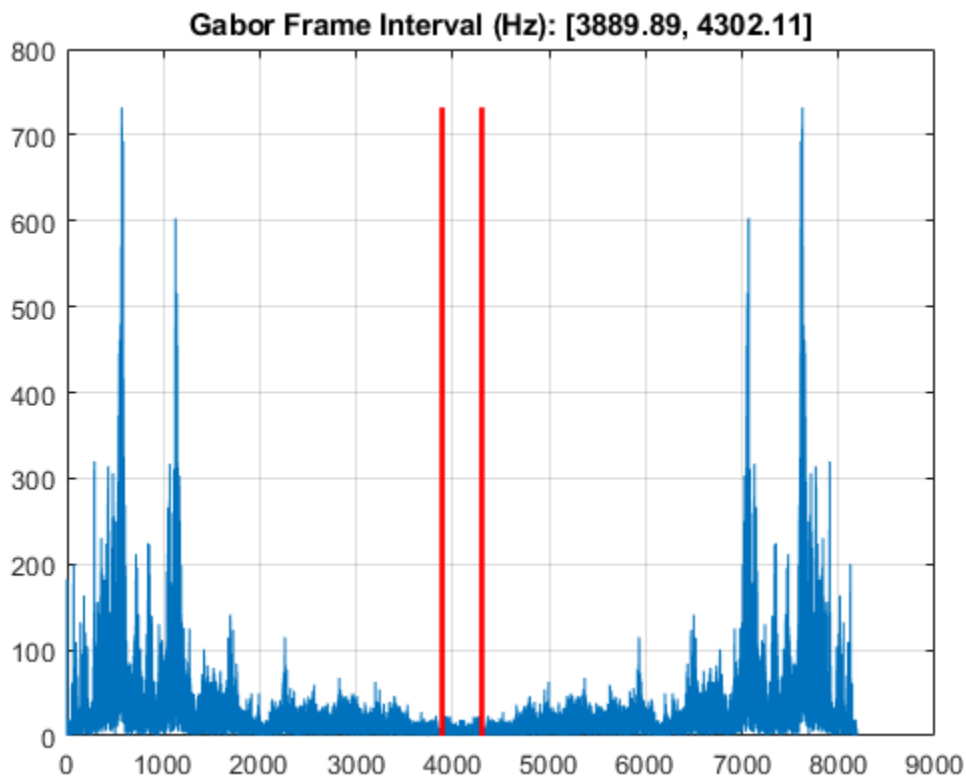
```



In the constant-Q transform, the Gabor frames are applied to the discrete Fourier transform of the input signal, and the inverse discrete Fourier transform is performed. The k -th Gabor frame is applied to the k -th frequency interval specified in `fintervals`. Take the discrete Fourier transform of the signal and plot its magnitude spectrum. Use

`fintervals` to indicate over which Fourier coefficients are the Gabor frame associated with the Nyquist frequency are applied.

```
yDFT = fft(y);  
lyDFT = length(yDFT);  
plot(Fs*(0:lyDFT-1)/lyDFT,abs(yDFT))  
grid on  
fIntervalGabor = fintervals{ind};  
mx = max(abs(yDFT));  
hold on  
plot([df*fIntervalGabor(1) df*fIntervalGabor(1)],[0 mx],'r-','LineWidth',2)  
plot([df*fIntervalGabor(end) df*fIntervalGabor(end)],[0 mx],'r-','LineWidth',2)  
str = sprintf('Gabor Frame Interval (Hz): [%3.2f, %3.2f]',df*fIntervalGabor(1),df*fInt  
title(str)
```



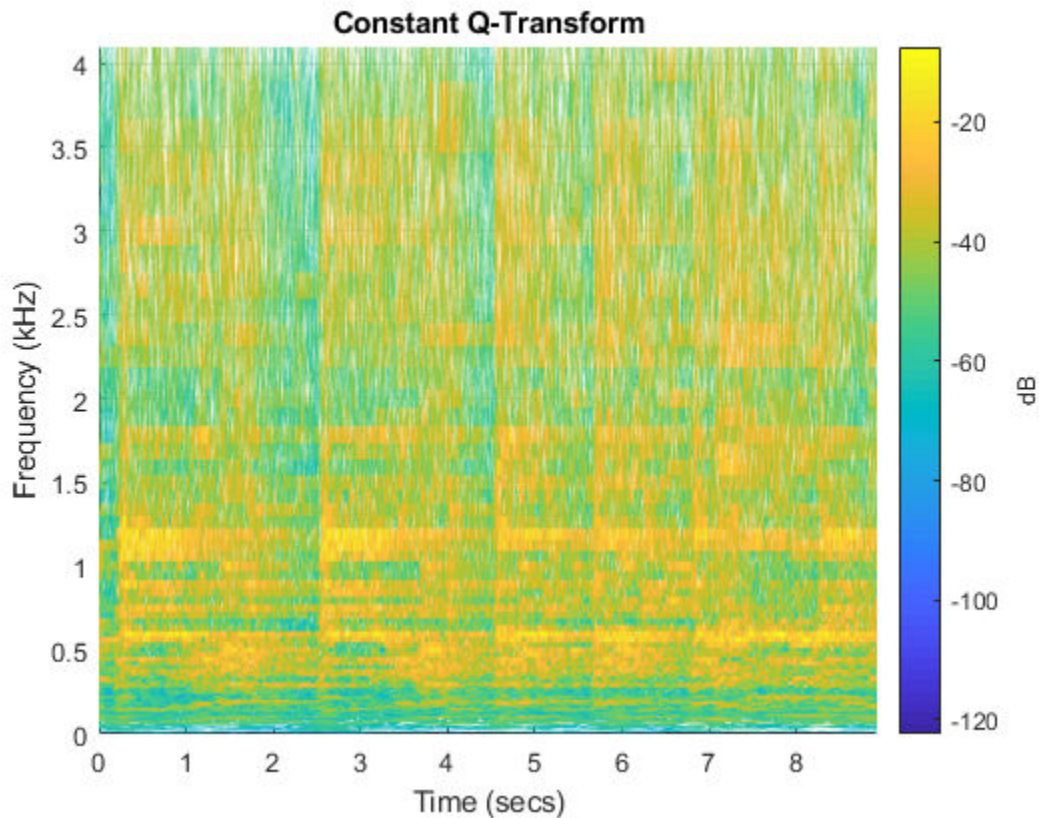
Window the Fourier coefficients in the interval with the Gabor frame, and take the inverse discrete Fourier transform. Normalize the result, and compare with computed constant-Q coefficients and confirm they are equal.

```
lGframe = length(gFrame);  
indx = 1:lGframe;  
indx = fftshift(indx);  
winDFT(indx) = yDFT(fIntervalGabor).*fftshift(gFrame(indx));  
cqCoefs = ifft(winDFT);  
cqCoefs = (2*lGframe/length(y))*cqCoefs;  
max(abs(cqCoefs(:)-cfs{ind}{:}))  
  
ans = 0
```

Constant-Q Transform of an Audio Signal

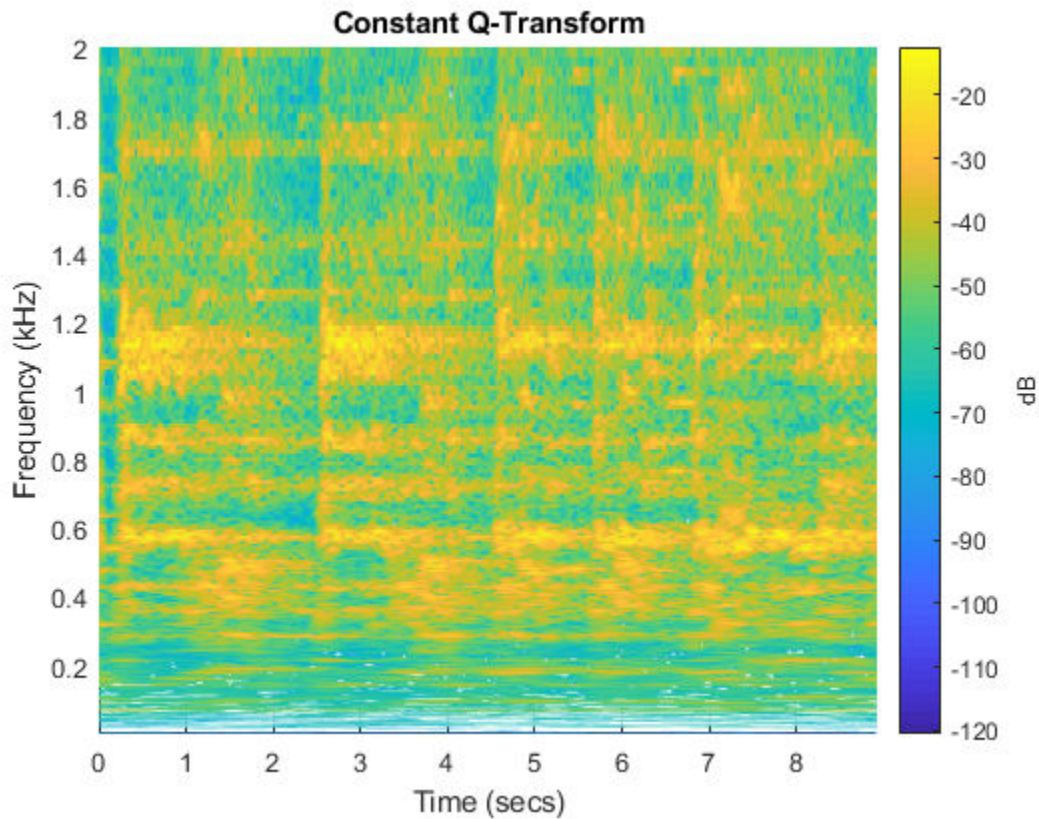
Load an audio signal. Plot the constant-Q transform using the maximally redundant version of the transform and using 12 bins per octave.

```
load handel  
cqt(y, 'SamplingFrequency', Fs)
```



Perform the CQT of the same signal using 48 bins per octave. Set the frequency range over which the CQT has a logarithmic frequency response to be the minimum allowable frequency to 2 kHz.

```
minFreq = Fs/length(y);  
maxFreq = 2000;  
figure  
cqt(y, 'SamplingFrequency',Fs, 'BinsPerOctave',48, 'FrequencyLimits',[minFreq maxFreq])
```



Input Arguments

x — Input signal

vector | matrix

Input signal, specified as a real or complex vector or matrix. **x** must have at least four samples.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'SamplingFrequency', 20, 'BinsPerOctave', 15`

SamplingFrequency — Sampling frequency

positive scalar

Sampling frequency, in Hz, specified as the comma-separated pair consisting of `'SamplingFrequency'` and a positive scalar.

BinsPerOctave — Number of bins per octave

12 (default) | positive integer from 1 to 96

Number of bins per octave to use in the CQT, specified as a positive integer from 1 to 96.

TransformType — Type of constant-Q transform

'full' (default) | 'sparse'

Type of constant-Q transform to perform, specified as the comma-separated pair consisting of `'TransformType'` and `'full'` or `'sparse'`. The sparse transform is the minimally redundant version of the constant-Q transform.

FrequencyLimits — Frequency limits

two-element real vector

Frequency limits over which the CQT has a logarithmic frequency response with the specified number of frequency bins per octave, specified as the comma-separated pair `'FrequencyLimits'` and a two-element real vector.

- The first element must be greater than or equal to F_s/N , where F_s is the sampling frequency and N is the length of the signal.
- The second element must be strictly less than the Nyquist frequency.

Window — Window to use as prototype function

'hann' (default) | 'hamming' | 'blackmanharris' | 'itersine' | 'bartlett'

Window to use as the prototype function for the nonstationary Gabor frames, specified as `'hann'`, `'hamming'`, `'blackmanharris'`, `'itersine'`, or `'bartlett'`. These

compactly support functions are defined in frequency. For normalized frequencies, they are defined on the interval $(-1/2, 1/2)$. If you specify a sampling frequency, F_s , they are defined on the interval $(-F_s/2, F_s/2)$.

Output Arguments

cfs — Constant-Q transform

matrix | multidimensional array | cell array | structure array

Constant-Q transform, returned as a matrix, multidimensional array, cell array, or structure array.

- If 'TransformType' is specified as 'full' without 'FrequencyLimits', **cfs** is a matrix or multidimensional array.
 - If **x** is a vector, then **cqt** returns a matrix corresponding to the CQT.
 - If **x** is a matrix, then **cqt** obtains the CQT for each column (independent channel) of **x**. The function returns a multidimensional array corresponding to the maximally redundant version of the CQT.

The array, **cfs**, corresponds to the maximally redundant version of the CQT. Each row of the pages of **cfs** corresponds to passbands with normalized center frequencies (cycles/sample) logarithmically spaced between 0 and 1. A normalized frequency of 1/2 corresponds to the Nyquist frequency. The number of columns, or hops, corresponds to the largest bandwidth center frequency, which usually occurs one frequency bin below or above the Nyquist bin.

- If 'TransformType' is specified as 'full' and you specify frequency limits, **cfs** is returned as a structure array with the following four fields.
 - **c** — Coefficient matrix of multidimensional array for the frequencies within the specified frequency limits. This includes both the positive and "negative" frequencies.
 - **DCcfs** — Coefficient vector or matrix for the passband from 0 to the lower frequency limit.
 - **Nyquistcfs** — Coefficient vector or matrix for the passband from the upper frequency limit to the Nyquist.
 - **NyquistBin** — DFT bin corresponding to the Nyquist frequency. This field is used when inverting the CQT.

- If 'TransformType' is specified as 'sparse', `cfs` is a cell array with the number of elements equal to the number of bandpass frequencies. Each element of the cell array, `cfs`, is a vector or matrix with the number of rows equal to the value of the bandwidth in DFT bins, `bw`.

`cfs`, `g`, and `fshifts` are required inputs for the inversion of the CQT with `icqt`.

f — Approximate bandpass center frequencies

real-valued vector

Approximate bandpass center frequencies corresponding to the rows of `cfs`, returned as a real-valued vector. The frequencies are ordered from 0 to 1 and are in cycles/sample. If you specified 'SamplingFrequency', then `f` is in Hertz.

g — Gabor frames

cell array of real-valued vectors

Gabor frames used in the analysis of `x`, returned as a cell array of real-valued vectors. Each vector in `g` corresponds to a row of `cfs`.

`cfs`, `g`, and `fshifts` are required inputs for the inversion of the CQT with `icqt`.

fshifts — Frequency shifts

real-valued vector

Frequency shifts in discrete Fourier transform bins, returned as a real-valued vector. The shifts are between the passbands in the rows of `cfs`.

`cfs`, `g`, and `fshifts` are required inputs for the inversion of the CQT with `icqt`.

fintervals — Frequency intervals

cell array of real-valued vectors

Frequency intervals corresponding to the rows of `cfs`, returned as a cell array. Each element in `fintervals` is a real-valued vector. The k th element of `fshifts` is the frequency shift in DFT bins between the $((k-1) \bmod N)$ and $(k \bmod N)$ element of `fintervals` with $k = 0, 1, 2, \dots, N-1$ where N is the number of frequency shifts. Because MATLAB indexes from 1, `fshifts(1)` contains the frequency shift between `fintervals{end}` and `fintervals{1}`, `fshifts(2)` contains the frequency shift between `fintervals{1}` and `fintervals{2}`, and so on.

bw — Bandwidths

real-valued vector

Bandwidths in DFT bins of the frequency intervals, `fintervals`, returned as a real-valued vector.

Algorithms

Nonstationary Gabor Frames

The theory of nonstationary Gabor (NSG) frames for frequency-adaptive analysis and efficient algorithms for analysis and synthesis using NSG frames are due to Dörfler, Holighaus, Grill, and Velasco [1],[2]. The algorithms used in CQT and ICQT were developed by Dörfler, Holighaus, Grill, and Velasco and are described in [1],[2]. In [3], Schörkhuber, Klapuri, Holighaus, and Dörfler develop and provide algorithms for a phase-corrected CQT transform which matches the CQT coefficients that would be obtained by naïve convolution. The Large Time-Frequency Analysis Toolbox (<https://github.com/ltfat>) provides an extensive suite of algorithms for nonstationary Gabor frames [4].

Perfect Reconstruction

To achieve the perfect reconstruction property of the constant-Q analysis with nonstationary Gabor frames, `cqt` internally prepends the zero frequency (DC) and appends the Nyquist frequency to the frequency interval. The negative frequencies are mirrored versions of the positive center frequencies and bandwidths

References

- [1] Holighaus, N., M. Dörfler, G. A. Velasco, and T. Grill. "A framework for invertible real-time constant-Q transforms." *IEEE Transactions on Audio, Speech, and Language Processing*. Vol. 21, No. 4, 2013, pp. 775-785.
- [2] Velasco, G. A., N. Holighaus, M. Dörfler, and T. Grill. "Constructing an invertible constant-Q transform with nonstationary Gabor frames." In *Proceedings of the 14th International Conference on Digital Audio Effects (DAFx-11)*. Paris, France: 2011.
- [3] Schörkhuber, C., A. Klapuri, N. Holighaus, and M. Dörfler. "A Matlab Toolbox for Efficient Perfect Reconstruction Time-Frequency Transforms with Log-Frequency Resolution." Submitted to the *AES 53rd International Conference on Semantic Audio*. London, UK: 2014.

- [4] Průša, Z., P. L. Søndergaard, N. Holighaus, C. Wiesmeyr, and P. Balazs. *The Large Time-Frequency Analysis Toolbox 2.0*. Sound, Music, and Motion, Lecture Notes in Computer Science 2014, pp 419-442.

See Also

`icqt`

Topics

“Nonstationary Gabor Frames and the Constant-Q Transform”

Introduced in R2018a

cwt

Continuous 1-D wavelet transform

Note See `cwt` for information on the older version of the `cwt`. The older version is no longer recommended.

Syntax

```
wt = cwt(x)
wt = cwt(x,wname)

[wt,f] = cwt( ____,fs)
[wt,period] = cwt( ____,ts)
[wt,f,coi] = cwt( ____,fs)
[wt,period,coi] = cwt( ____,ts)

[ ____ ] = cwt( ____,Name,Value)
[ ____,coi,fb] = cwt( ____ )
[ ____,fb,scalingcfs] = cwt( ____ )

cwt( ____ )
```

Description

`wt = cwt(x)` returns the continuous wavelet transform (CWT) of `x`. The input, `x`, is a double-precision real- or complex-valued vector, or a single-variable regularly sampled timetable and must have at least four samples. The CWT is obtained using the analytic Morse wavelet with the symmetry parameter (γ) equal to 3 and the time-bandwidth product equal to 60. `cwt` uses 10 voices per octave. The minimum and maximum scales are determined automatically based on the energy spread of the wavelet in frequency and time. If `x` is real-valued, `wt` is a 2-D matrix where each row corresponds to one scale. The column size of `wt` is equal to the length of `x`. If `x` is complex-valued, `wt` is a 3-D matrix, where the first page is the CWT for the positive scales (analytic part or counterclockwise component) and the second page is the CWT for the negative scales (anti-analytic part or clockwise component).

The `cwt` function uses L1 normalization. With L1 normalization, if you have equal amplitude oscillatory components in your data at different scales, they will have equal magnitude in the CWT. Using L1 normalization shows a more accurate representation of the signal. See “L1 Norm for CWT” on page 1-170 and “Sinusoid and Wavelet Coefficient Amplitudes” on page 1-145.

The `cwt` function supports GPU code generation. See “GPU Code Generation” on page 1-168 for more information.

`wt = cwt(x, wname)` uses the analytic wavelet specified by `wname` to compute the CWT. Valid options for `wname` are 'morse', 'amor', and 'bump', which specify the Morse, Morlet (Gabor), and bump wavelet, respectively. If you do not specify `wname`, `wname` defaults to 'morse'.

`[wt, f] = cwt(____, fs)` specifies the sampling frequency, `fs`, in Hz as a positive scalar. `cwt` uses `fs` to determine the scale-to-frequency conversions and returns the frequencies `f` in Hz. If you do not specify a sampling frequency, `cwt` returns `f` in cycles per sample. If the input `x` is complex, the scale-to-frequency conversions apply to both pages of `wt`. If `x` is a timetable, you cannot specify `fs`. `fs` is determined from the `RowTimes` of the timetable.

`[wt, period] = cwt(____, ts)` specifies the sampling period, `ts`, as a positive duration scalar. The duration can be in years, days, hours, minutes, or seconds. `cwt` uses `ts` to compute the scale-to-period conversion and returns the time periods in `period`. The array of durations in `period` has the same format property as `ts`. If the input `x` is complex, the scale-to-period conversions apply to both pages of `wt`. If `x` is a timetable, you cannot specify `ts`. `ts` is determined from the `RowTimes` of the timetable when you set the 'PeriodLimits' name-value pair.

`[wt, f, coi] = cwt(____, fs)` returns the cone of influence, `coi`, which shows where edge effects of the CWT become significant. The cone of influence for the CWT is in Hz. If the input `x` is complex, the cone of influence applies to both pages of `wt`.

`[wt, period, coi] = cwt(____, ts)` returns the cone of influence, `coi`, which shows where edge effects of the CWT become significant. The cone of influence for the CWT is in periods. If the input `x` is complex, the cone of influence applies to both pages of `wt`.

`[____, Name, Value] = cwt(____, Name, Value)` returns the CWT with additional options specified by one or more `Name, Value` pair arguments.

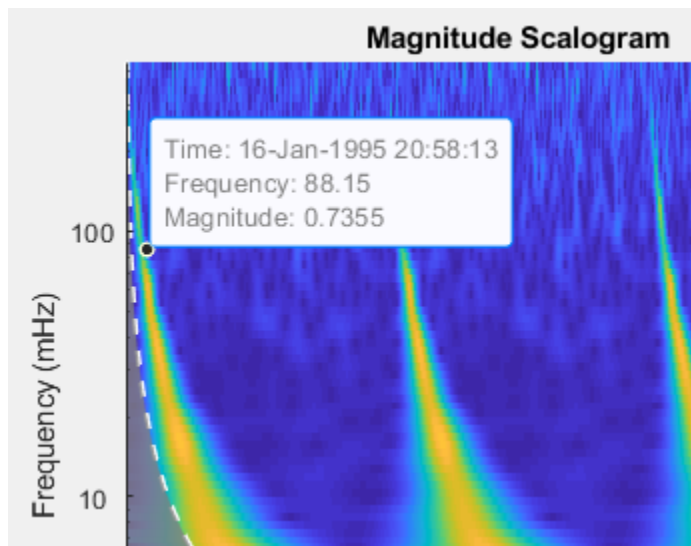
`[____, coi, fb] = cwt(____, coi, fb)` returns the filter bank used in the CWT. See `cwtfilterbank`.

`[____,fb,scalingcfs] = cwt(____)` returns the scaling coefficients if the analyzing wavelet is 'morse' or 'amor'. Scaling coefficients are not supported for the bump wavelet.

`cwt(____)` with no output arguments plots the CWT scalogram. The scalogram is the absolute value of the CWT as a function of time and frequency. Frequency is plotted on a logarithmic scale. The cone of influence showing where edge effects become significant is also plotted. Gray regions outside the dashed white line delineate regions where edge effects are significant. If the input signal is complex-valued, the positive (counterclockwise) and negative (clockwise) components are plotted in separate scalograms.

If you do not specify a sampling frequency or sampling period, the frequencies are plotted in cycles per sample. If you specify a sampling frequency, the frequencies are in Hz. If you specify a sampling period, the scalogram is plotted as a function of time and periods. If the input signal is a timetable, the scalogram is plotted as a function of time and frequency in hertz and uses the RowTimes as the basis for the time axis.

To see the time, frequency, and magnitude of a scalogram point, enable data tips in the figure axes toolbar and click the desired point in the scalogram.



Note Before plotting, `cwt` clears (`clf`) the current figure. To plot the scalogram in a subplot, use a plotting function.

Examples

Continuous Wavelet Transform Using Default Values

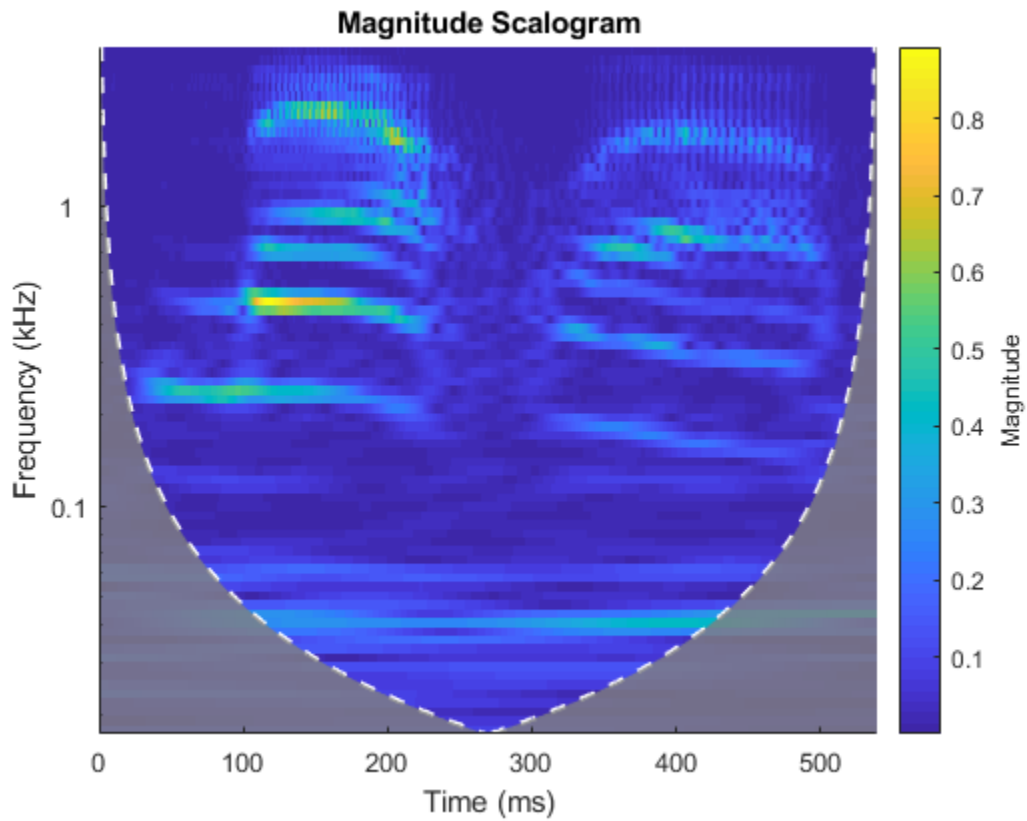
Obtain the continuous wavelet transform of a speech sample using default values.

```
load mtlb;  
w = cwt(mtlb);
```

Continuous Wavelet Transform Using Specified Wavelet

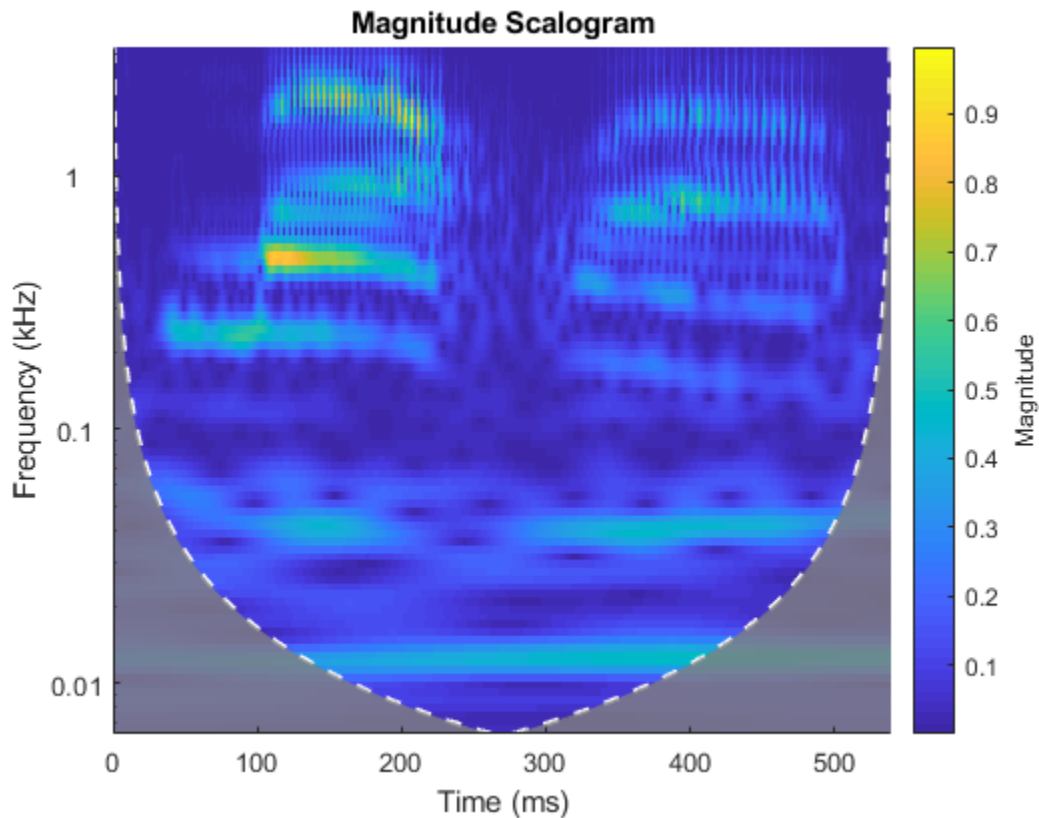
Obtain the continuous wavelet transform of a speech sample using the bump wavelet instead of the default Morse wavelet.

```
load mtlb;  
cwt(mtlb, 'bump', Fs);
```

Compare the result obtained from the CWT using the default Morse wavelet.

```
cwt(mtlb,Fs);
```



Continuous Wavelet Transform of Two Complex Exponentials

Create two complex exponentials, of different amplitudes, with frequencies of 32 and 64 Hz. The data is sampled at 1000 Hz. The two complex exponentials have disjoint support in time.

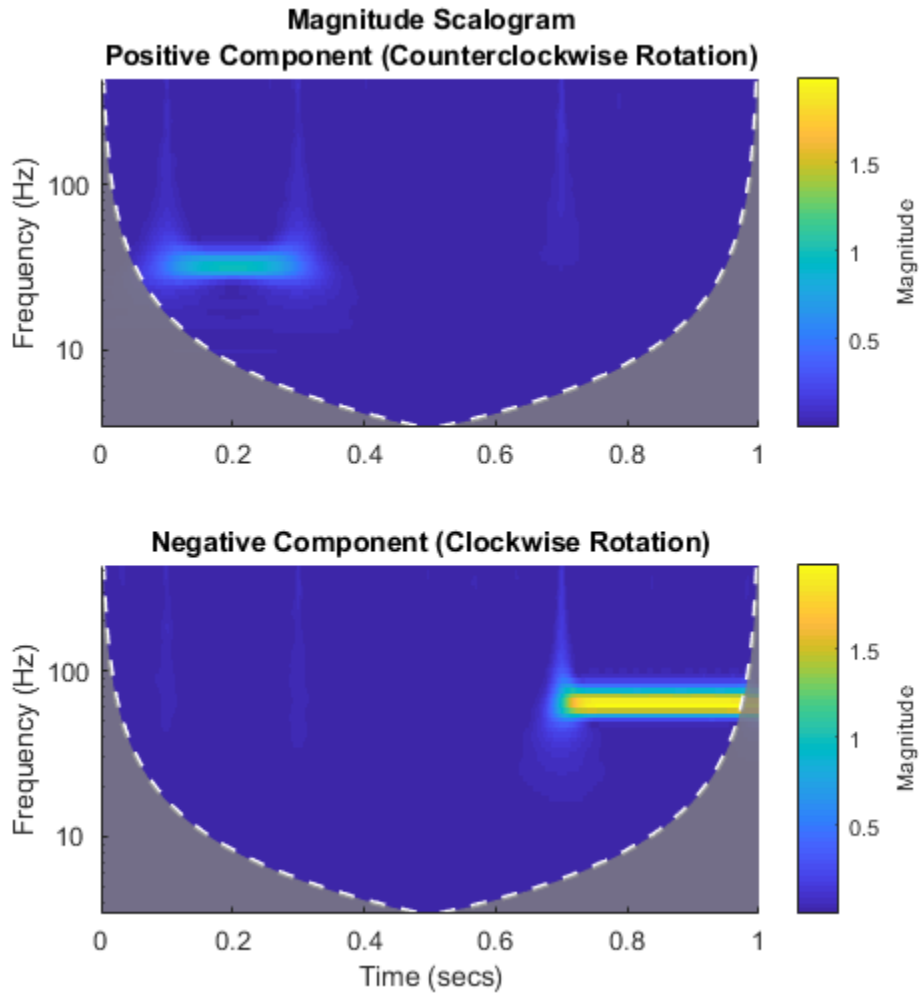
```
Fs = 1e3;
t = 0:1/Fs:1;
z = exp(1i*2*pi*32*t).*(t>=0.1 & t<0.3)+2*exp(-1i*2*pi*64*t).*(t>0.7);
```

Add complex white Gaussian noise with a standard deviation of 0.05.

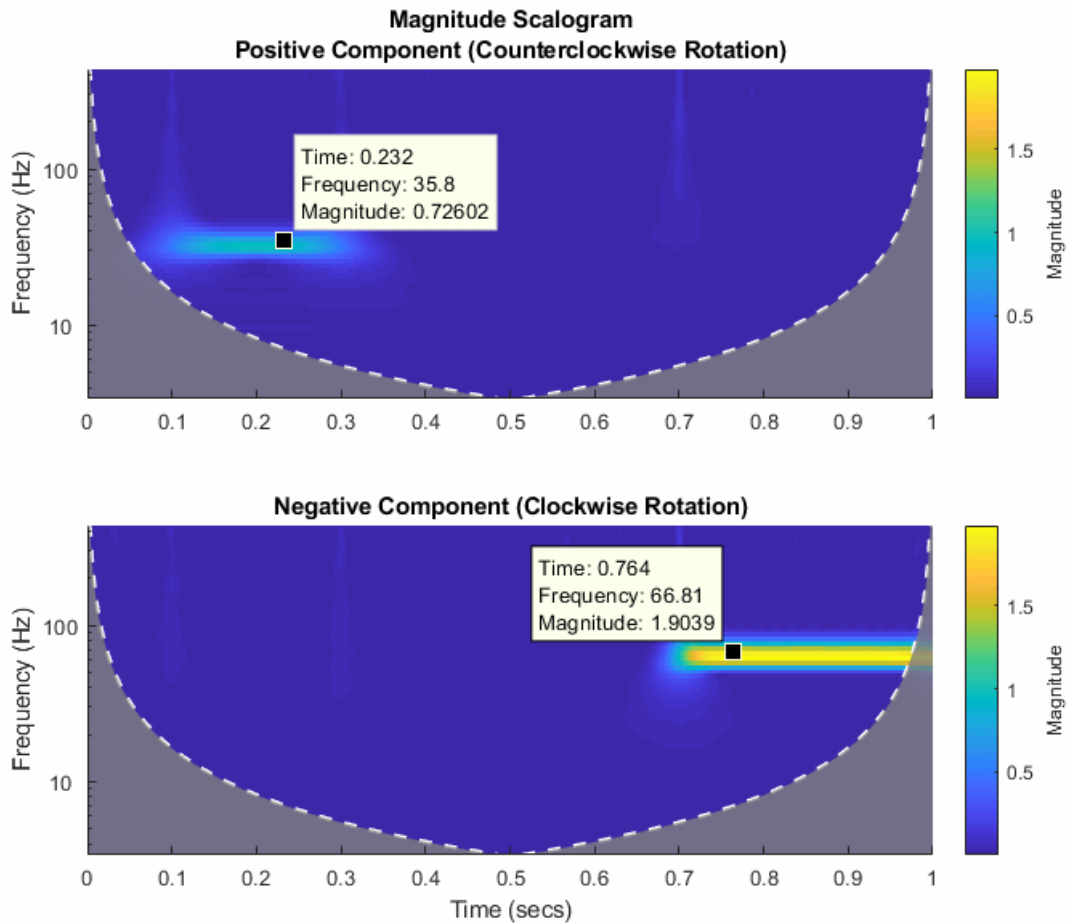
```
wgnNoise = 0.05/sqrt(2)*randn(size(t))+1i*0.05/sqrt(2)*randn(size(t));  
z = z+wgnNoise;
```

Obtain and plot the cwt using a Morse wavelet.

```
cwt(z, Fs)
```



Note the magnitudes of the complex exponential components in the colorbar are essentially their amplitudes even though they are at different scales. This is a direct result of the L1 normalization. You can verify this by executing this script and exploring each subplot with a data cursor.



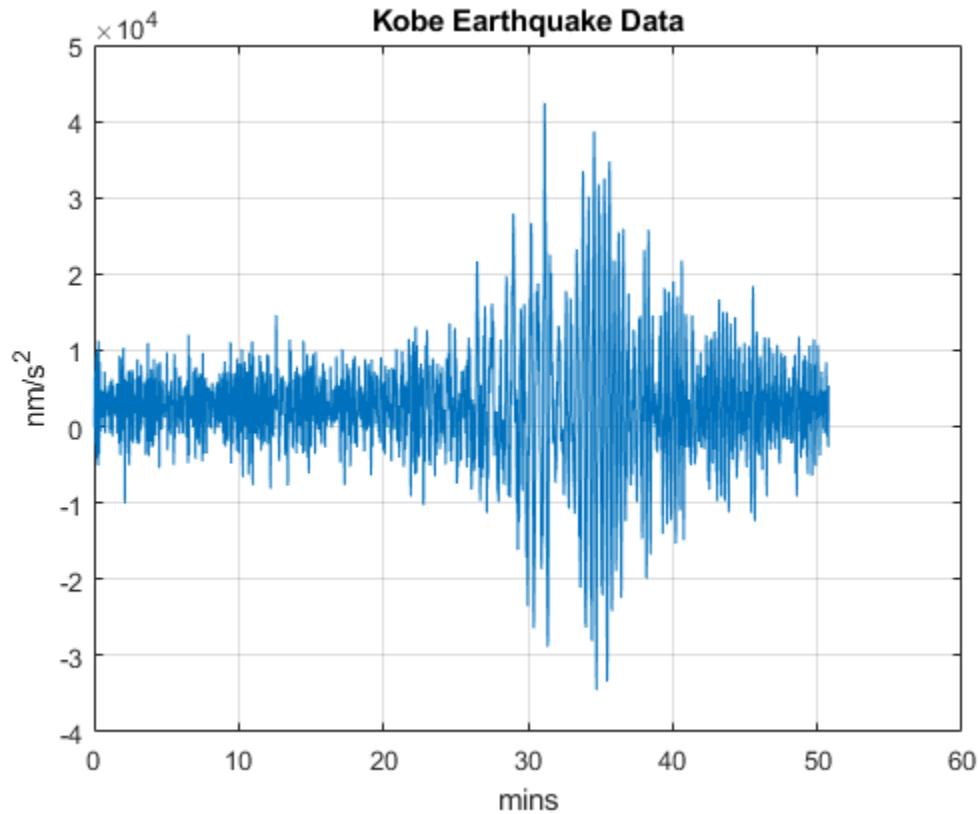
Continuous Wavelet Transform of Earthquake Data

Obtain the CWT of Kobe earthquake data. The sampling frequency is 1 Hz.

```
load kobe
```

Plot the earthquake data.

```
plot((1:numel(kobe))./60,kobe)  
xlabel('mins')  
ylabel('nm/s^2')  
grid on  
title('Kobe Earthquake Data')
```

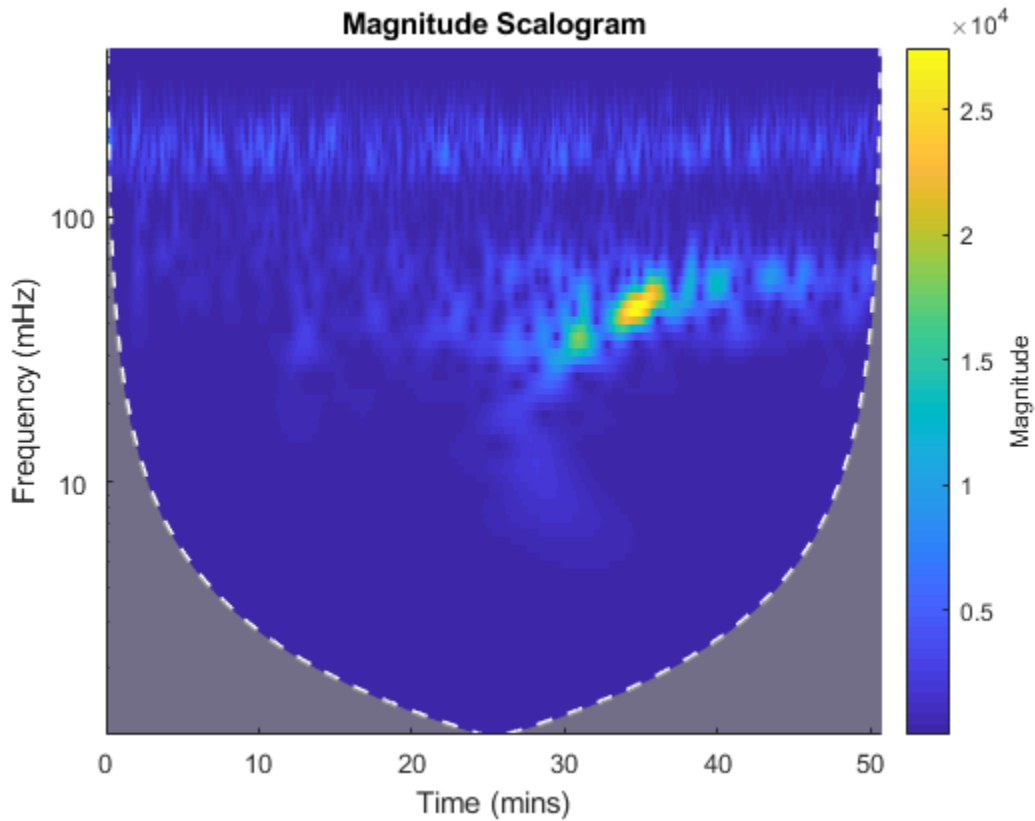


Obtain the CWT, frequencies, and cone of influence.

```
[wt,f,coi] = cwt(kobe,1);
```

Plot the data, including the cone of influence.

```
cwt(kobe,1)
```

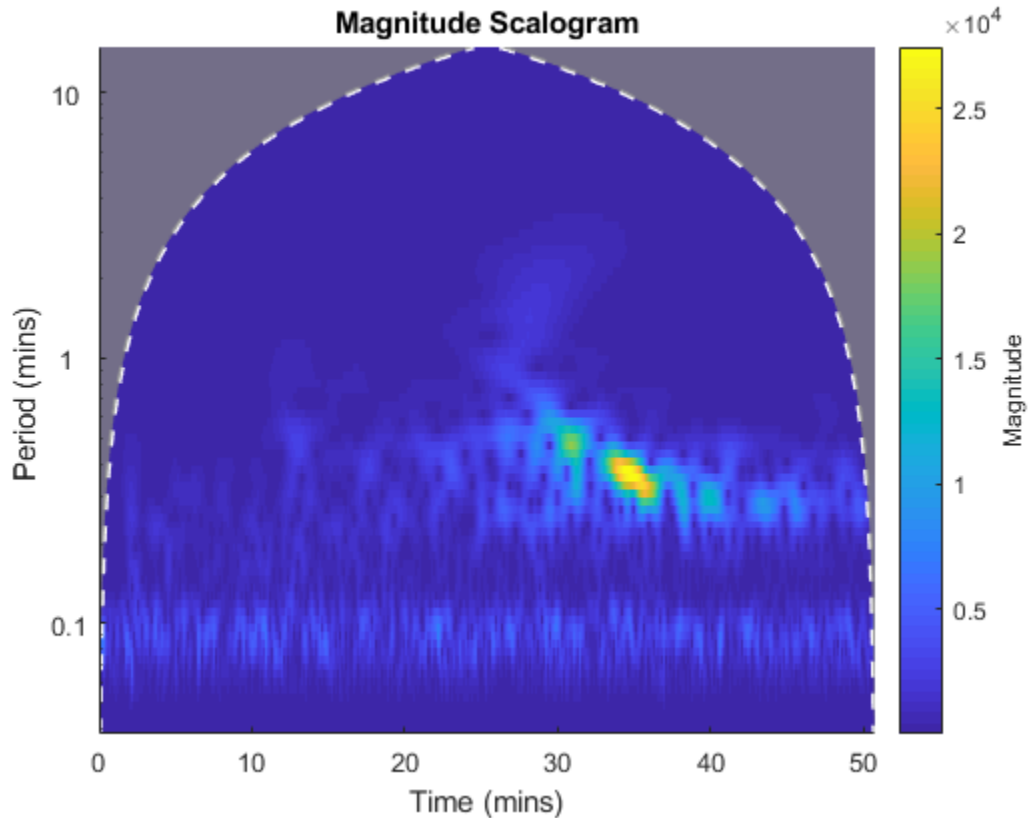


Obtain the CWT, time periods, and cone of influence by specifying a sampling period instead of a sampling frequency.

```
[wt,periods,coi] = cwt(kobe,minutes(1/60));
```

View the same data by specifying a sampling period input instead of a frequency.

```
cwt(kobe,minutes(1/60))
```

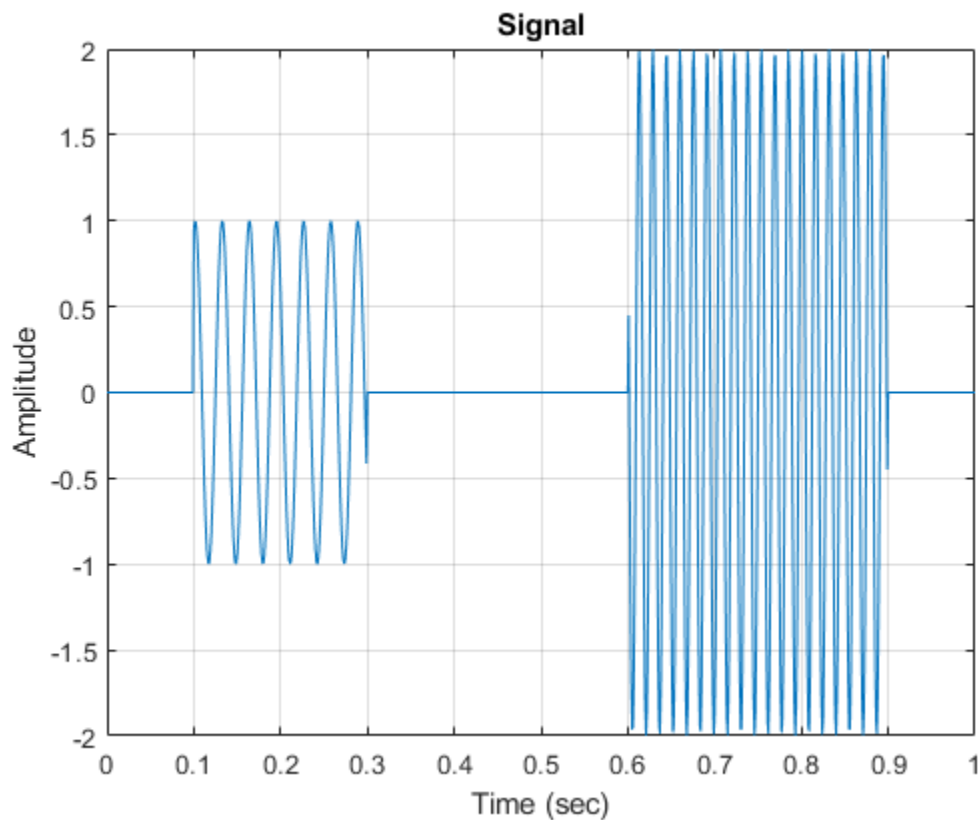


Sinusoid and Wavelet Coefficient Amplitudes

This example shows that the amplitudes of oscillatory components in a signal agree with the amplitudes of the corresponding wavelet coefficients.

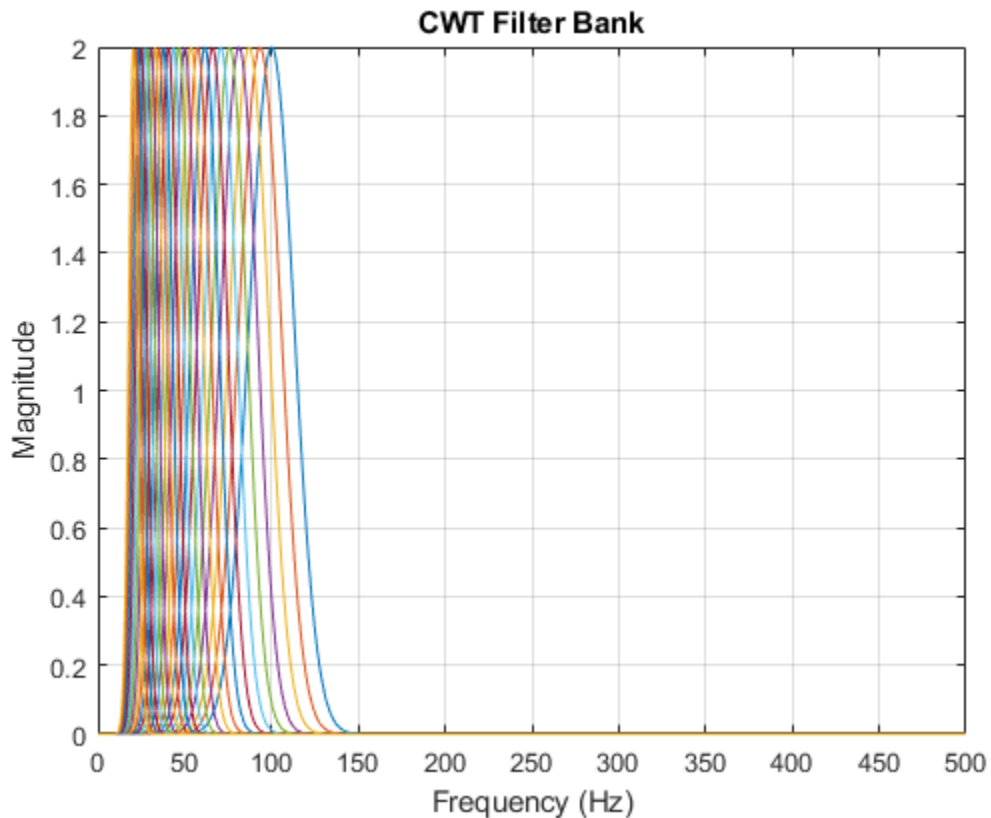
Create a signal composed of two sinusoids with disjoint support in time. One sinusoid has a frequency of 32 Hz and amplitude equal to 1. The other sinusoid has a frequency of 64 Hz and amplitude equal to 2. The signal is sampled for one second at 1000 Hz. Plot the signal.

```
frq1 = 32;  
amp1 = 1;  
frq2 = 64;  
amp2 = 2;  
  
Fs = 1e3;  
t = 0:1/Fs:1;  
x = amp1*sin(2*pi*frq1*t).*(t>=0.1 & t<0.3)+amp2*sin(2*pi*frq2*t).*(t>0.6 & t<0.9);  
  
plot(t,x)  
grid on  
xlabel('Time (sec)')  
ylabel('Amplitude')  
title('Signal')
```



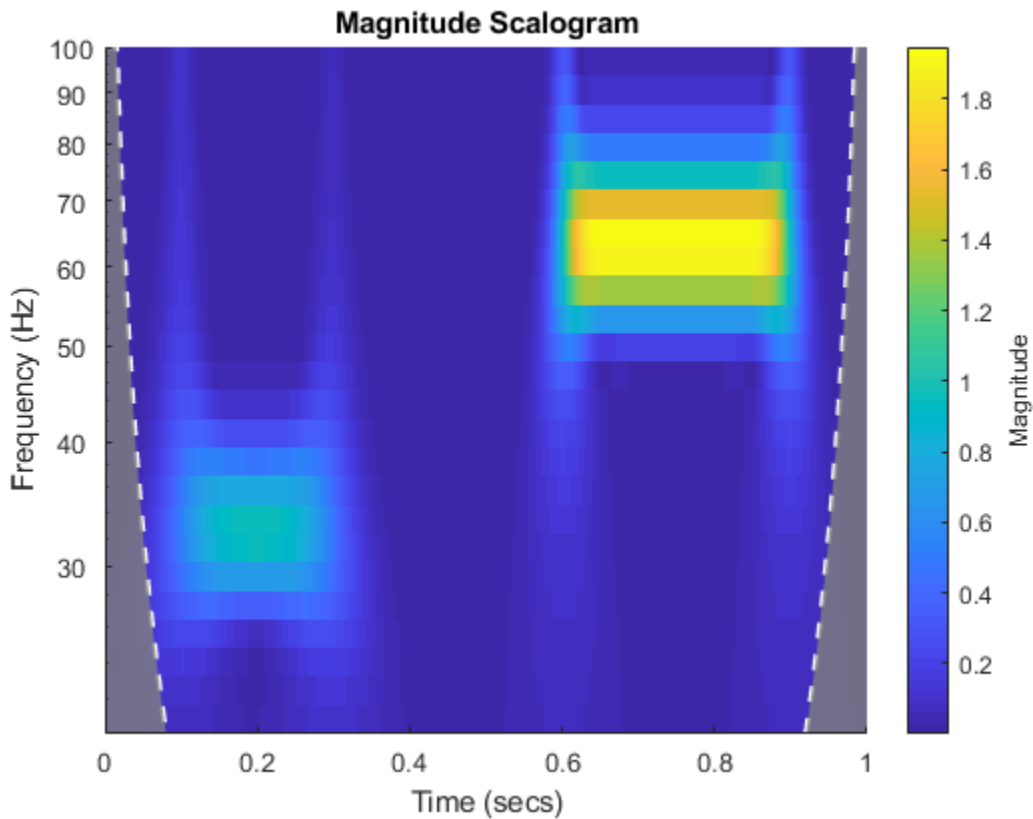
Create a CWT filter bank that can be applied to the signal. Since the signal component frequencies are known, set the frequency limits of the filter bank to a narrow range that includes the known frequencies. To confirm the range, plot the magnitude frequency responses for the filter bank.

```
fb = cwtfilterbank('SignalLength',numel(x),'SamplingFrequency',Fs,...  
    'FrequencyLimits',[20 100]);  
figure  
freqz(fb)
```

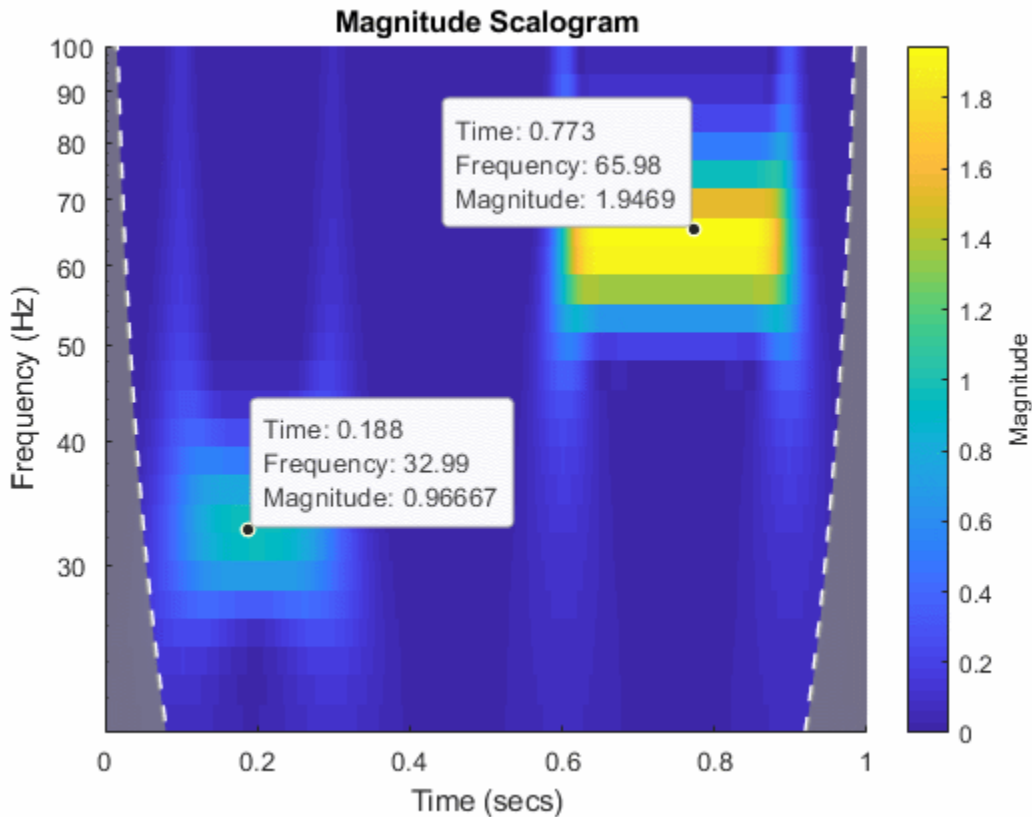


Use `cwt` and the filter bank to plot the scalogram of the signal.

```
figure  
cwt(x,'FilterBank',fb)
```



Execute this script and use a data cursor to confirm that the amplitudes of the wavelet coefficients are essentially equal to the amplitudes of the sinusoidal components.



Using CWT Filter Bank on Multiple Time Series

This example shows how using a CWT filter bank improves computational efficiency when taking the CWT of multiple time series.

Load the seismograph data recorded during the 1995 Kobe earthquake. The data are seismograph (vertical acceleration, nm/sq.sec) measurements recorded at Tasmania University, Hobart, Australia on 16 January 1995 beginning at 20:56:51 (GMT) and continuing for 51 minutes at 1 second intervals. Create a CWT filter bank that can be applied to the data.

```
load kobe
fb = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1);
```

Use the `cwt` function and take the CWT of the data 250 times. Display the elapsed time used.

```
num = 250;
tic;
for k=1:num
    cfs = cwt(kobe);
end
toc
```

Elapsed time is 6.551628 seconds.

Now use the `wt` object function of the filter bank to take the CWT of the data. Confirm using the filter bank is faster.

```
tic;
for k=1:num
    cfs = wt(fb, kobe);
end
toc
```

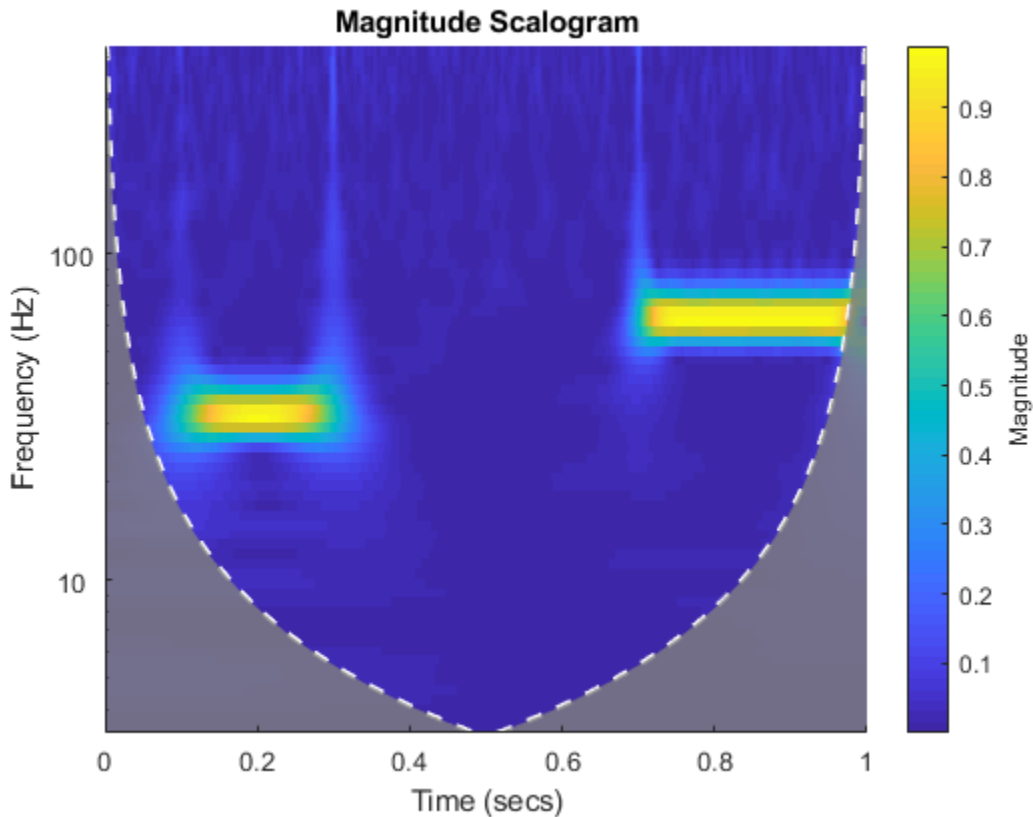
Elapsed time is 3.782376 seconds.

Change Default Frequency Axis Labels

This example shows how to change the default frequency axis labels for the CWT when you obtain a plot with no output arguments.

Create two sine waves with frequencies of 32 and 64 Hz. The data is sampled at 1000 Hz. The two sine waves have disjoint support in time. Add white Gaussian noise with a standard deviation of 0.05. Obtain and plot the CWT using the default Morse wavelet.

```
Fs = 1e3;
t = 0:1/Fs:1;
x = cos(2*pi*32*t).*(t>=0.1 & t<0.3)+sin(2*pi*64*t).*(t>0.7);
wgnNoise = 0.05*randn(size(t));
x = x+wgnNoise;
cwt(x, 1000)
```



The plot uses a logarithmic frequency axis because frequencies in the CWT are logarithmic. In MATLAB, logarithmic axes are in powers of 10 (decades). You can use `cwtfreqbounds` to determine what the minimum and maximum wavelet bandpass frequencies are for a given signal length, sampling frequency, and wavelet.

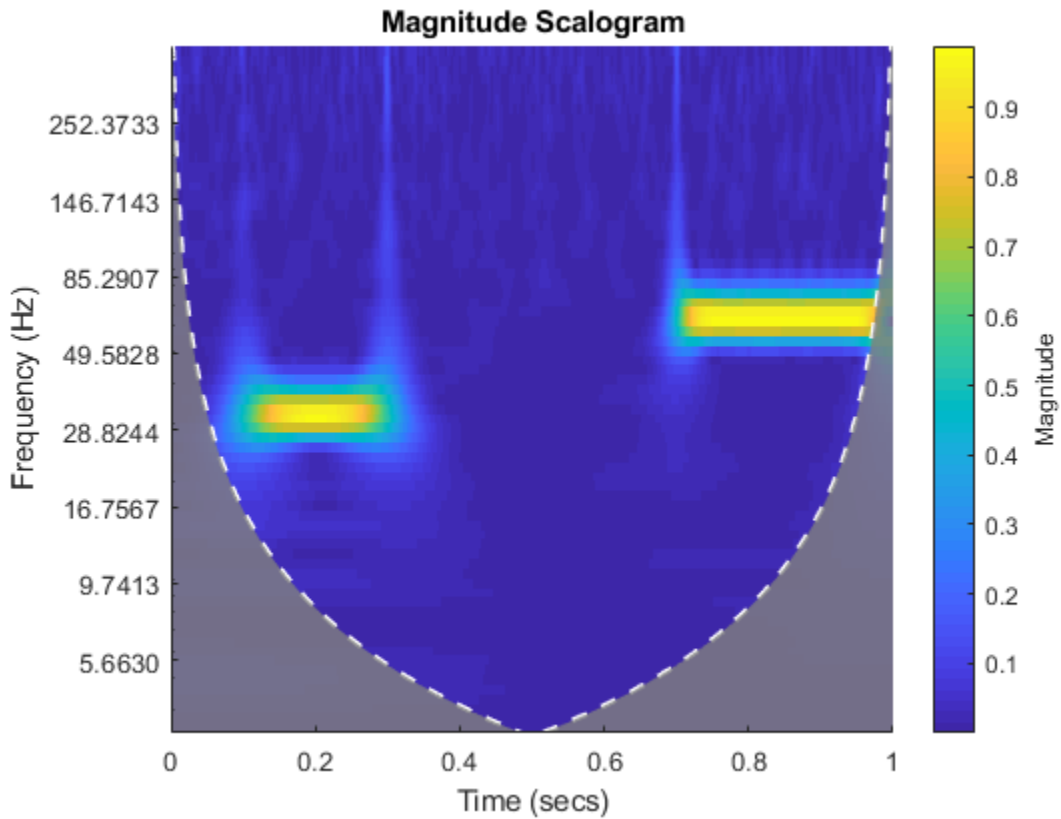
```
[minf,maxf] = cwtfreqbounds(numel(x),1000);
```

You see that by default MATLAB has placed frequency ticks at 10 and 100 because those are the powers of 10 between the minimum and maximum frequencies. If you wish to add more frequency axis ticks, you can obtain a logarithmically spaced set of frequencies between the minimum and maximum frequencies using the following.

```
numfreq = 10;
freq = logspace(log10(minf),log10(maxf),numfreq);
```

Next, get the handle to the current axes and replace the frequency axis ticks and labels with the following.

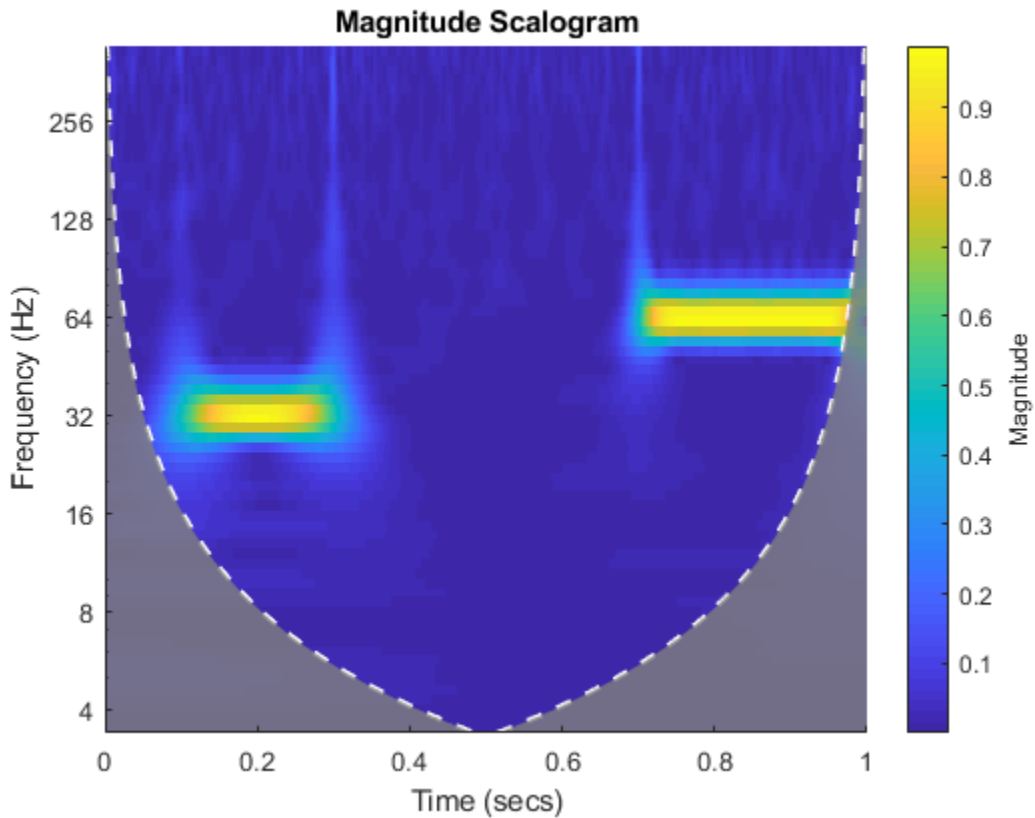
```
AX = gca;  
AX.YTickLabelMode = 'auto';  
AX.YTick = freq;
```



In the CWT, frequencies are computed in powers of two. To create the frequency ticks and tick labels in powers of two, you can do the following.

```
newplot;  
cwt(x,1000);  
AX = gca;  
freq = 2.^(round(log2(minf)):round(log2(maxf)));
```

```
AX.YTickLabelMode = 'auto';
AX.YTick = freq;
```

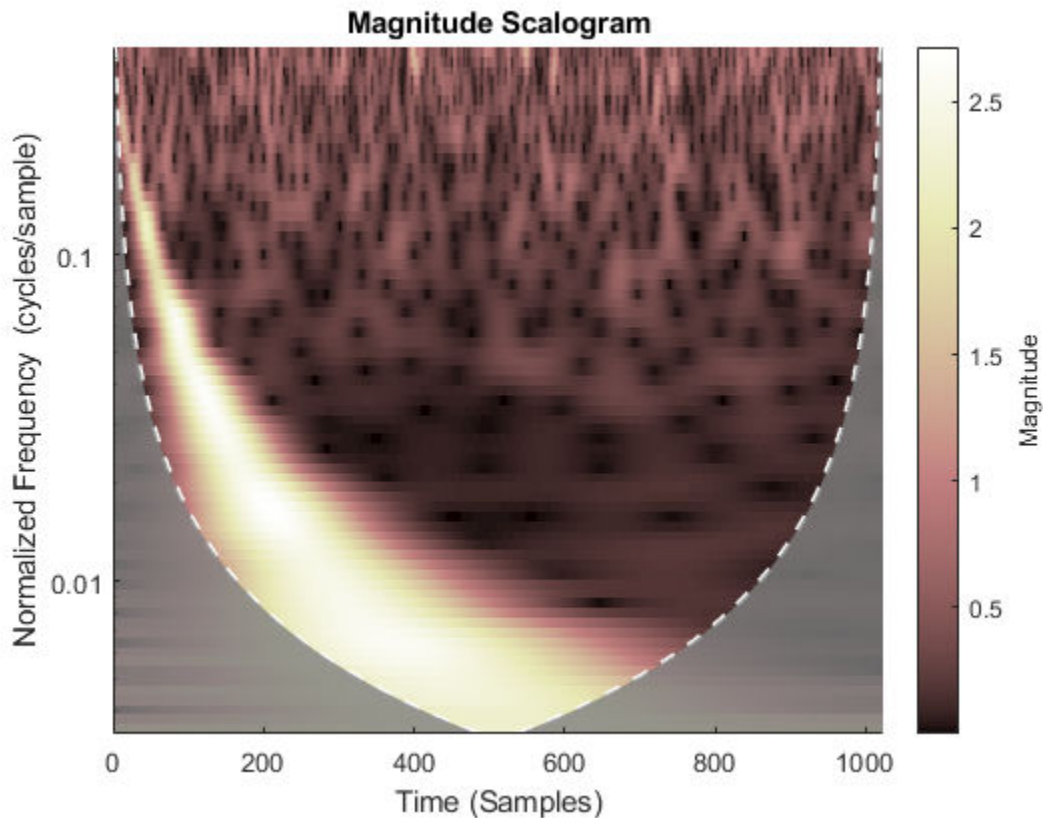


Change Scalogram Coloration

This example shows how to scale scalogram values by maximum absolute value at each level for plotting.

Load in a signal and display the default scalogram. Change the colormap to pink(240).

```
load noisdopp
cwt(noisdopp)
colormap(pink(240))
```



Take the CWT of the signal and obtain the wavelet coefficients and frequencies.

```
[cfs, frq] = cwt(noisdopp);
```

To efficiently find the maximum value of the coefficients at each frequency (level), first transpose the absolute value of the coefficients. Find the minimum value at every level. At each level, subtract the level's minimum value.

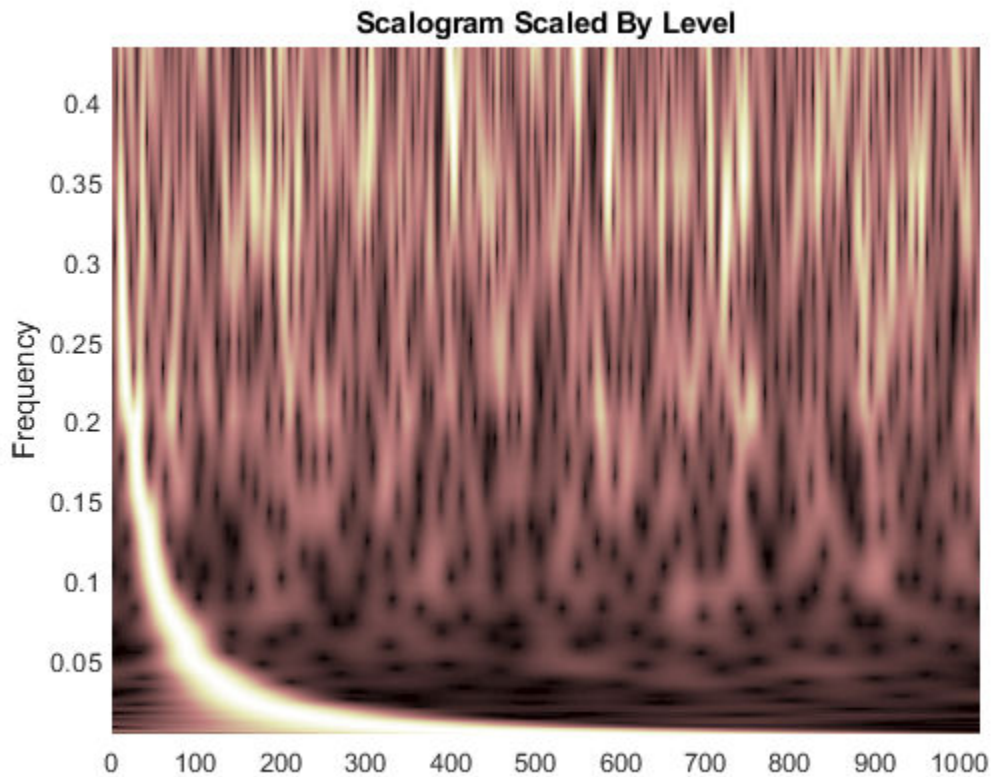
```
tmp1 = abs(cfs);
t1 = size(tmp1,2);
tmp1 = tmp1';
minv = min(tmp1);
tmp1 = (tmp1-minv(ones(1,t1),:));
```


Find the maximum value at every level of `tmp1`. For each level, divide every value by the maximum value at that level. Multiply the result by the number of colors in the colormap. Set equal to 1 all zero entries. Transpose the result.

```
maxv = max(tmp1);
maxvArray = maxv(ones(1,t1),:);
indx = maxvArray<eps;
tmp1 = 240*(tmp1./maxvArray);
tmp2 = 1+fix(tmp1);
tmp2(indx) = 1;
tmp2 = tmp2';
```

Display the result. The scalogram values are now scaled by the maximum absolute value at each level. Frequencies are displayed on a linear scale.

```
t = 0:length(noisdopp)-1;
pcolor(t,frq,tmp2);
shading interp
ylabel('Frequency')
title('Scalogram Scaled By Level')
colormap(pink(240))
```



Changing the Time-bandwidth Product

This example shows that increasing the time-bandwidth product P^2 of the Morse wavelet creates a wavelet with more oscillations under its envelope. All other things being equal, increasing P^2 makes the wavelet more frequency-localized as a result.

Create two filter banks. One filter bank has the default `TimeBandwidth` value of 60. The second filter bank has a `TimeBandwidth` value of 10. The `SignalLength` for both filter banks is 4096 samples.

```
sigLen = 4096;
fb60 = cwtfilterbank('SignalLength',sigLen);
fb10 = cwtfilterbank('SignalLength',sigLen,'TimeBandwidth',10);
```

Obtain the time-domain wavelets for the filter banks.

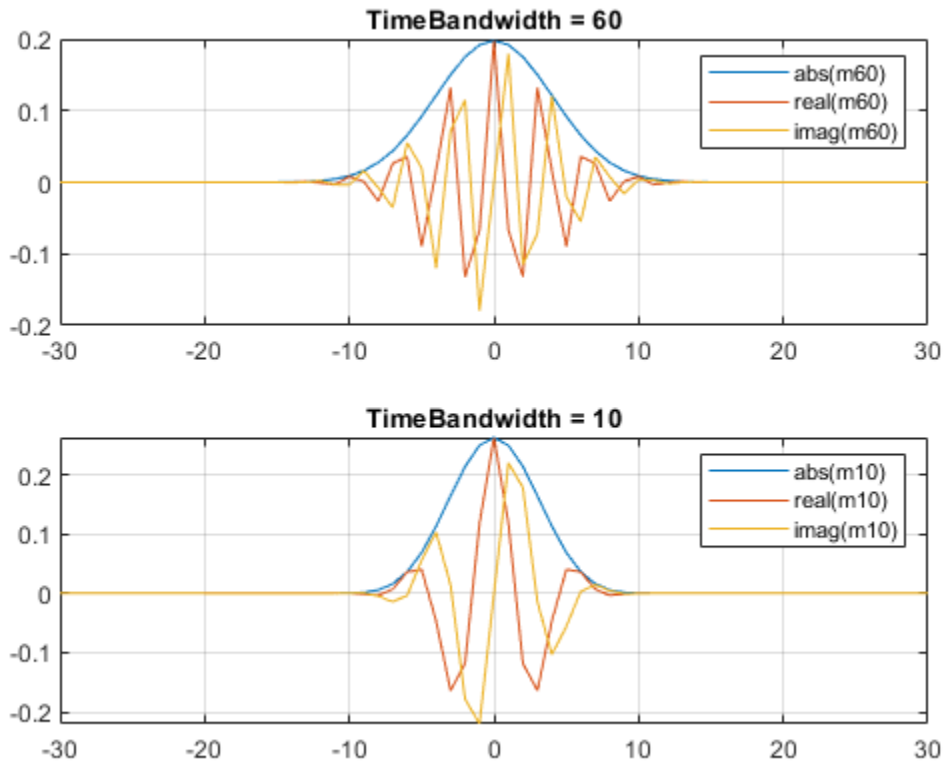
```
[psi60,t] = wavelets(fb60);
[psi10,~] = wavelets(fb10);
```

Use the `scales` function to find the mother wavelet for each filter bank.

```
sca60 = scales(fb60);
sca10 = scales(fb10);
[~,idx60] = min(abs(sca60-1));
[~,idx10] = min(abs(sca10-1));
m60 = psi60(idx60,:);
m10 = psi10(idx10,:);
```

Since the time-bandwidth product is larger for the `fb60` filter bank, verify the `m60` wavelet has more oscillations under its envelope than the `m10` wavelet.

```
subplot(2,1,1)
plot(t,abs(m60))
grid on
hold on
plot(t,real(m60))
plot(t,imag(m60))
xlim([-30 30])
legend('abs(m60)', 'real(m60)', 'imag(m60)')
title('TimeBandwidth = 60')
subplot(2,1,2)
plot(t,abs(m10))
grid on
hold on
plot(t,real(m10))
plot(t,imag(m10))
xlim([-30 30])
legend('abs(m10)', 'real(m10)', 'imag(m10)')
title('TimeBandwidth = 10')
```



Align the peaks of the m60 and m10 magnitude frequency responses. Verify the frequency response of the m60 wavelet is more narrow than the frequency response for the m10 wavelet.

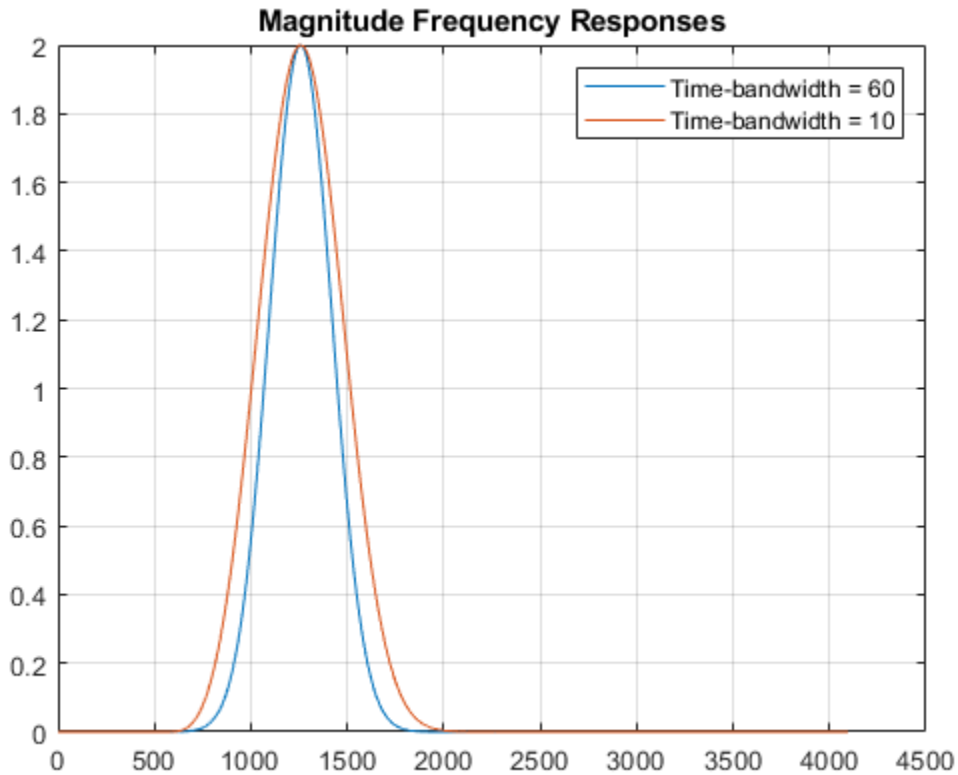
```
cf60 = centerFrequencies(fb60);
cf10 = centerFrequencies(fb10);

m60cFreq = cf60(id60);
m10cFreq = cf10(id10);

freqShift = 2*pi*(m60cFreq-m10cFreq);
x10 = m10.*exp(1j*freqShift*(-sigLen/2:sigLen/2-1));

figure
```

```
plot([abs(fft(m60)).' abs(fft(x10)).'])  
grid on  
legend('Time-bandwidth = 60','Time-bandwidth = 10')  
title('Magnitude Frequency Responses')
```

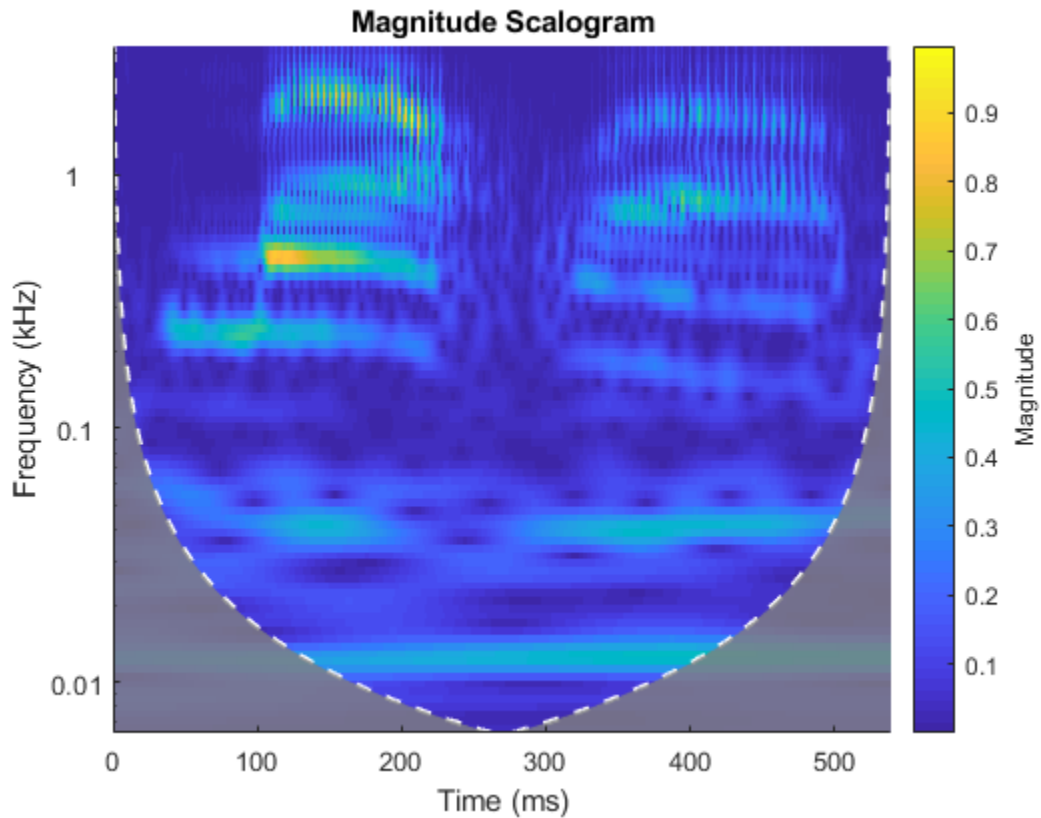


Plot CWT Scalogram in Subplot

This example shows how to plot the CWT scalogram in a figure subplot.

Load the speech sample. The data is sampled at 7418 Hz. Plot the default CWT scalogram.

```
load mtlb  
cwt(mtlb,Fs)
```



Obtain the continuous wavelet transform of the signal, and the frequencies of the CWT.

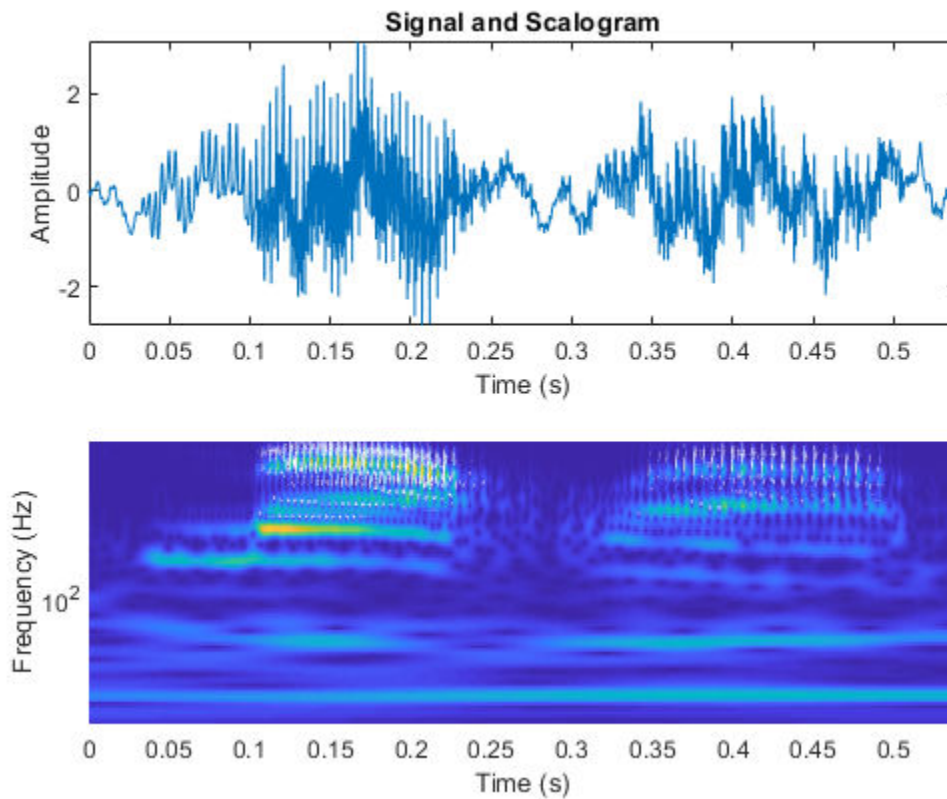
```
[cfs, frq] = cwt(mtlb, Fs);
```

The `cwt` function sets the time and frequency axes in the scalogram. Create a vector representing the sample times.

```
tms = (0: numel(mtlb)-1)/Fs;
```

In a new figure, plot the original signal in the upper subplot and the scalogram in the lower subplot. Plot the frequencies on a logarithmic scale.

```
figure
subplot(2,1,1)
plot(tms,mtlb)
axis tight
title('Signal and Scalogram')
xlabel('Time (s)')
ylabel('Amplitude')
subplot(2,1,2)
surface(tms, frq, abs(cfs))
axis tight
shading flat
xlabel('Time (s)')
ylabel('Frequency (Hz)')
set(gca, 'yscale', 'log')
```



Input Arguments

x — Input signal

real- or complex-valued vector | timetable

Input signal, specified as a double-precision real- or complex-valued vector or a single-variable regularly sampled timetable. **x** must have at least four samples.

wname — Analytic wavelet

'morse' (default) | 'amor' | 'bump'

Analytic wavelet used to compute the CWT, specified as 'morse', 'amor', or 'bump'. These character vectors specify the analytic Morse, Morlet (Gabor), and bump wavelet, respectively.

The default Morse wavelet has symmetry parameter (γ) equal to 3 and time-bandwidth product equal to 60.

fs — Sampling frequency

positive scalar

Sampling frequency, in Hz, specified as a positive scalar. If you specify **fs**, then you cannot specify **ts**.

ts — Sampling period

scalar duration

Sampling period, also known as the time duration, specified as a scalar duration. Valid durations are *years*, *days*, *hours*, *minutes*, and *seconds*. You cannot use calendar durations. If you specify **ts**, then you cannot specify **fs**.

Example: `wt = cwt(x, hours(12))`

Name-Value Pair Arguments

Specify optional comma-separated pairs of **Name**, **Value** arguments. **Name** is the argument name and **Value** is the corresponding value. **Name** must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: 'ExtendSignal', false indicates that the signal is not extended.

ExtendSignal — Extend input signal symmetrically

true (default) | false

Option to extend the input signal symmetrically by reflection, specified as the comma-separated pair consisting of 'ExtendSignal' and either true or false. Extending the signal symmetrically can mitigate boundary effects. If you specify true, then the signal is extended. If you specify false, then the signal is not extended.

FrequencyLimits — Frequency limits

two-element scalar vector

Frequency limits to use in the CWT, specified as a two-element vector with positive strictly increasing entries. The first element specifies the lowest peak passband frequency and must be greater than or equal to the product of the wavelet peak frequency in hertz and two time standard deviations divided by the signal length. The second element specifies the highest peak passband frequency and must be less than or equal to the Nyquist frequency. The base 2 logarithm of the ratio of the maximum frequency to the minimum frequency must be greater than or equal to $1/NV$ where NV is the number of voices per octave.

If you specify frequency limits outside the permissible range, cwt truncates the limits to the minimum and maximum valid values. Use `cwtfreqbounds` to determine frequency limits for different parameterizations of the CWT. For complex-valued signals, $(-1)*flimits$ is used for the anti-analytic part, where `flimits` is the vector specified by `FrequencyLimits`.

Example: 'FrequencyLimits', [0.1 0.3]

Data Types: double

PeriodLimits — Period limits

two-element duration array

Period limits to use in the CWT, specified as a two-element duration array with strictly increasing positive entries. The first element must be greater than or equal to $2*ts$ where ts is the sampling period. The base 2 logarithm of the ratio of the minimum period to the maximum period must be less than or equal to $-1/NV$ where NV is the number of voices per octave. The maximum period cannot exceed the signal length divided by the product of two time standard deviations of the wavelet and the wavelet peak frequency.

If you specify period limits outside the permissible range, `cwt` truncates the limits to the minimum and maximum valid values. Use `cwtfreqbounds` to determine period limits for different parameterizations of the wavelet transform. For complex-valued signals, `(-1)*plimits` is used for the anti-analytic part, where `plimits` is the vector specified by `PeriodLimits`.

Example: `'PeriodLimits',[seconds(0.2) seconds(1)]`

Data Types: `duration`

VoicesPerOctave — Number of voices per octave

10 (default) | even integer from 4 to 48

Number of voices per octave to use for the CWT, specified as the comma-separated pair consisting of `'VoicesPerOctave'` and an even integer from 4 to 48. The CWT scales are discretized using the specified number of voices per octave. The energy spread of the wavelet in frequency and time automatically determines the minimum and maximum scales.

TimeBandwidth — Time-bandwidth product of the Morse wavelet

60 (default) | scalar greater than 3 and less than or equal to 120

Time-bandwidth product of the Morse wavelet, specified as the comma-separated pair consisting of `'TimeBandwidth'` and a scalar greater than 3 and less than or equal to 120. The symmetry parameter, γ , is fixed at 3. Wavelets with larger time-bandwidth products have larger spreads in time and narrower spreads in frequency. The standard deviation of the Morse wavelet in time is approximately $\sqrt{\text{TimeBandwidth}/2}$. The standard deviation of the Morse wavelet in frequency is approximately $1/2*\sqrt{2/\text{TimeBandwidth}}$.

If you specify `'TimeBandwidth'`, you cannot specify `'WaveletParameters'`. To specify both the symmetry and time-bandwidth product, use `'WaveletParameters'` instead.

In the notation of “Morse Wavelets”, `TimeBandwidth` is P^2 .

WaveletParameters — Symmetry and time-bandwidth product of the Morse wavelet

(3,60) (default) | two-element vector of scalars

Symmetry and time-bandwidth product of the Morse wavelet, specified as the comma-separated pair consisting of `'WaveletParameters'` and a two-element vector of scalars.

The first element is the symmetry, γ , which must be greater than or equal to 1. The second element is the time-bandwidth product, which must be strictly greater than γ . The ratio of the time-bandwidth product to γ cannot exceed 40.

When γ is equal to 3, the Morse wavelet is perfectly symmetric in the frequency domain and the skewness is 0. When γ is greater than 3, the skewness is positive. When γ is less than 3, the skewness is negative.

If you specify 'WaveletParameters', you cannot specify 'TimeBandwidth'.

NumOctaves — Number of octaves

positive integer

Number of octaves, specified as the comma-separated pair consisting of 'NumOctaves' and a positive integer. The number of octaves cannot exceed $\log_2(f_{\max}/f_{\min})$ where f_{\max} and f_{\min} are the maximum and minimum CWT frequencies (or periods) as determined by the signal length, sampling frequency, and wavelet. See `cwtfreqbounds` for details.

The 'NumOctaves' name-value pair is not recommended and will be removed in a future release. The recommended way to modify the frequency or period range of the CWT is with the 'FrequencyLimits' or 'PeriodLimits' name-value pairs. You cannot specify both the 'NumOctaves' and 'FrequencyLimits' or 'PeriodLimits' name-value pairs.

FilterBank — CWT filter bank

CWT filter bank object

CWT filter bank to use to compute the CWT, specified as a CWT filter bank object. If you use the 'FilterBank' name-value pair, you cannot specify any other options. All options for the computation of the CWT are defined as properties of `fb`.

If `x` is a timetable, the sampling frequency or sampling period in `fb` must agree with the sampling frequency or sampling period determined by the `RowTimes` of the timetable.

Example: `wt = cwt(x,'FilterBank',fb)`

Output Arguments

wt — Continuous wavelet transform

matrix

Continuous wavelet transform, returned as a matrix of complex values. By default, `cwt` uses the analytic Morse (3,60) wavelet, where 3 is the symmetry and 60 is the time-bandwidth product. `cwt` uses 10 voices per octave. If `x` is real-valued, `wt` is an N_a -by- N matrix, where N_a is the number of scales, and N is the number of samples in `x`. If `x` is complex-valued, `wt` is a 3-D matrix, where the first page is the CWT for the positive scales (analytic part or counterclockwise component) and the second page is the CWT for the negative scales (anti-analytic part or clockwise component). The minimum and maximum scales are determined automatically based on the energy spread of the wavelet in frequency and time. See “Algorithms” on page 1-168 for information on how the scales are determined.

Data Types: `double`

f — Frequencies

vector

Frequencies of the CWT, returned as a vector. If you specify a sampling frequency, `fs`, then `f` is in Hz. If you do not specify `fs`, `cwt` returns `f` in cycles per sample.

period — Time periods

array

Time periods, returned as an array of durations. The durations are in the same format as `ts`. Each row corresponds to a period.

coi — Cone of influence

array of doubles |array of durations

Cone of influence for the CWT, returned as either an array of doubles or array of durations. The cone of influence indicates where edge effects occur in the CWT. If you specify a sampling frequency, `fs`, the cone of influence is in Hz. If you specify a scalar duration, `ts`, the cone of influence is in periods. Due to the edge effects, give less credence to areas that are outside or overlap the cone of influence.

For additional information, see “Boundary Effects and the Cone of Influence”.

fb — CWT filter bank

CWT filter bank object

CWT filter bank used in the CWT, returned as a CWT filter bank object. See `cwtfilterbank`.

scalingcfs — Scaling coefficients

real- or complex-valued vector

Scaling coefficients for the CWT if the analyzing wavelet is 'morse' or 'amor', returned as a real- or complex-valued vector. The length of `scalingcfs` is equal to the length of the input `x`.

Scaling coefficients are not supported for the bump wavelet.

Definitions

Analytic Wavelets

Analytic wavelets are complex-valued wavelets whose Fourier transform vanish for negative frequencies. Analytic wavelets are a good choice when doing time-frequency analysis with the CWT. Because the wavelet coefficients are complex-valued, the coefficients provide phase and amplitude information of the signal being analyzed. Analytic wavelets are well suited for studying how the frequency content in real world nonstationary signals evolves as a function of time.

Analytic wavelets are almost exclusively based on rapidly decreasing functions. If $\psi(t)$ is an analytic rapidly decreasing function in time, then its Fourier transform $\hat{\psi}(\omega)$ is a rapidly decreasing function in frequency and is small outside of some interval $\alpha < \omega < \beta$ where $0 < \alpha < \beta$. Orthogonal and biorthogonal wavelets are typically designed to have compact support in time. Wavelets with compact support in time have relatively poorer energy concentration in frequency than wavelets which rapidly decrease in time. Most orthogonal and biorthogonal wavelets are not symmetric in the Fourier domain.

If your goal is to obtain a joint time-frequency representation of your signal, we recommend you use `cwt` or `cwtfilterbank`. Both functions support the following analytic wavelets:

- Morse Wavelet Family (default)
- Analytic Morlet (Gabor) Wavelet
- Bump

If you want to do time-frequency analysis using orthogonal or biorthogonal wavelets, we recommend `modwpt`.

When using wavelets for time-frequency analysis, you usually convert scales to frequencies or periods to interpret results. `cwt` and `cwtfilterbank` do the conversion. You can obtain the corresponding scales associated by using `scales` on the optional `cwt` output argument `fb`.

For more information regarding Morse wavelets, see “Morse Wavelets”. For guidance on how to choose the wavelet that is right for your application, see “Choose a Wavelet”.

GPU Code Generation

The `cwt` supports CUDA[®] code generation. You must have MATLAB Coder™ and GPU Coder™ to generate GPU code.

Usage notes and limitations:

- Timetable input signal is not supported.
- All inputs must be constant and specified at compilation time.
- Only analytic Morse ('morse') and Morlet ('amor') wavelets are supported.
- The following input arguments are not supported: `ts` (sampling period), `PeriodLimits` name-value pair, `NumOctave` name-value pair, and `FilterBank` name-value pair.
- Scaling coefficient output and filter bank output are not supported.
- Plotting is not supported.

Tips

- If you are taking the CWT of multiple time series, it is more efficient to precompute and use a CWT filter bank than repeatedly apply the `cwt` function. See “Using CWT Filter Bank on Multiple Time Series” on page 1-149.

Algorithms

Minimum Scale

To determine the minimum scale, find the peak frequency ω_x of the base wavelet. For Morse wavelets, dilate the wavelet so that the Fourier transform of the wavelet at π

radians is equal to 10% of the peak frequency. The smallest scale occurs at the largest frequency:

$$s_0 = \frac{\omega'_x}{\pi}$$

As a result, the smallest scale is the minimum of $(2, s_0)$. For Morse wavelets, the smallest scale is usually s_0 . For the Morlet wavelet, the smallest scale is usually 2.

Maximum Scale

Both the minimum and maximum scales of the CWT are determined automatically based on the energy spread of the wavelet in frequency and time. To determine the maximum scale, CWT uses the following algorithm.

The standard deviation of the Morse wavelet in time, σ_t , is approximately $\sqrt{\frac{P^2}{2}}$, where P^2 is the time-bandwidth product. The standard deviation in frequency, σ_f , is approximately $\frac{1}{2}\sqrt{\frac{2}{P^2}}$. If you scale the wavelet by some $s > 1$, the time duration changes to $2s\sigma_t = N$, which is the wavelet stretched to equal the full length (N samples) of the input. You cannot translate this wavelet or stretch it further without causing it to wrap, so the largest scale is $\text{floor}\left(\frac{N}{2\sigma_t}\right)$.

Wavelet transform scales are powers of 2 and are denoted by $s_0\left(2^{\frac{1}{NV}}\right)^j$. NV is the number of voices per octave, and j ranges from 0 to the largest scale. For a specific small scale, s_0 :

$$s_0\left(2^{\frac{1}{NV}}\right)^j \leq \frac{N}{2\sigma_t}$$

Converting to log2:

$$j\log_2\left(2^{\frac{1}{NV}}\right) \leq \log_2\left(\frac{N}{2\sigma_t s_0}\right)$$

$$j \leq NV\log_2\left(\frac{N}{2\sigma_t s_0}\right)$$

Therefore, the maximum scale is

$$s_0 \left(\frac{1}{2^{NV}} \right)^{\text{floor} \left(NV \log_2 \left(\frac{N}{2\sigma_t s_0} \right) \right)}$$

L1 Norm for CWT

In integral form, the CWT preserves energy. However, when you implement the CWT numerically, energy is not preserved. In this case, regardless of the normalization you use, the CWT is not an orthonormal transform. The `cwt` function uses L1 normalization.

Wavelet transforms commonly use L2 normalization of the wavelet. For the L2 norm, dilating a signal by $1/s$, where s is greater than 0, is defined as follows:

$$\left\| x \left(\frac{t}{s} \right) \right\|_2^2 = s \|x(t)\|_2^2$$

The energy is now s times the original energy. When included in the Fourier transform, multiplying by $1/\sqrt{s}$ produces different weights being applied to different scales, so that the peaks at higher frequencies are reduced more than the peaks at lower frequencies.

In many applications, L1 normalization is better. The L1 norm definition does not include squaring the value, so the preserving factor is $1/s$ instead of $1/\sqrt{s}$. Instead of high-frequency amplitudes being reduced as in the L2 norm, for L1 normalization, all frequency amplitudes are normalized to the same value. Therefore, using the L1 norm shows a more accurate representation of the signal. See example “Continuous Wavelet Transform of Two Complex Exponentials” on page 1-140.

Compatibility Considerations

'NumOctaves' name-value pair will be removed

Not recommended starting in R2018a

The 'NumOctaves' name-value pair argument will be removed in a future release. Use either:

- Name-value pair argument 'FrequencyLimits' to modify the frequency range of the CWT.
- Name-value pair argument 'PeriodLimits' to modify the period range of the CWT.

See `cwtfreqbounds` for additional information.

References

- [1] Lilly, J. M., and S. C. Olhede. "Generalized Morse Wavelets as a Superfamily of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 60, No. 11, 2012, pp. 6036-6041.
- [2] Lilly, J. M., and S. C. Olhede. "Higher-Order Properties of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 57, No. 1, 2009, pp. 146-160.
- [3] Lilly, J. M. *jLab: A data analysis package for Matlab*, version 1.6.2. 2016. <http://www.jmlilly.net/jmlsoft.html>.
- [4] Lilly, J. M. "Element analysis: a wavelet-based method for analysing time-localized events in noisy time series." *Proceedings of the Royal Society A*. Volume 473: 20160776, 2017, pp. 1-28. [dx.doi.org/10.1098/rspa.2016.0776](https://doi.org/10.1098/rspa.2016.0776).

See Also

`cwtfilterbank` | `cwtfreqbounds` | `icwt`

Topics

"Continuous and Discrete Wavelet Transforms"

"Time-Frequency Analysis with the Continuous Wavelet Transform"

"Boundary Effects and the Cone of Influence"

"Morse Wavelets"

Introduced in R2016b

cwtfilterbank

Continuous wavelet transform filter bank

Description

Use `cwtfilterbank` to create a continuous wavelet transform (CWT) filter bank. The default wavelet used in the filter bank is the analytic Morse (3,60) wavelet. You can vary the time-bandwidth and symmetry parameters for the Morse wavelets, to tune the Morse wavelet for your needs. You can also use the analytic Morlet (Gabor) wavelet or bump wavelet. When analyzing multiple signals in time-frequency, for improved computational efficiency, you can precompute the filters once and then pass the filter bank as input to `cwt`. With the filter bank, you can visualize wavelets in time and frequency. You can also create filter banks with specific frequency or period ranges, and measure 3-dB bandwidths. You can determine the quality factor for the wavelets in the filter bank.

Creation

Syntax

```
fb = cwtfilterbank  
fb = cwtfilterbank(Name,Value)
```

Description

`fb = cwtfilterbank` creates a continuous wavelet transform (CWT) filter bank `fb`. The filters are normalized so that the peak magnitudes for all passbands are approximately equal to 2. The default filter bank is designed for a signal with 1024 samples. The default filter bank uses the analytic Morse (3,60) wavelet. The filter bank uses the default scales: approximately 10 wavelet bandpass filters per octave (10 voices per octave). The highest-frequency passband is designed so that the magnitude falls to half the peak value at the Nyquist frequency.

As implemented, the CWT uses L1 normalization. With L1 normalization, if you have equal amplitude oscillatory components in your data at different scales, they will have equal

magnitude in the CWT. Using L1 normalization shows a more accurate representation of the signal. See “L1 Norm for CWT” on page 1-170. The wavelet filters are normalized with peak magnitudes approximately equal to 2 so that the amplitudes of the oscillatory components agree with the amplitudes of the corresponding wavelet coefficients. See “Sinusoid and Wavelet Coefficient Amplitudes” on page 1-183.

`fb` can be used as input for `cwt`.

`fb = cwtfilterbank(Name, Value)` creates a CWT filter bank `fb` with properties specified by one or more `Name, Value` pair arguments. Properties can be specified in any order as `Name1, Value1, . . . , NameN, ValueN`. Enclose each property name in quotes.

Note You cannot change a property value of an existing filter bank. For example, if you have a filter bank `fb` with a `SignalLength` of 2000, you must create a second filter bank `fb2` to process a signal with 2001 samples. You cannot assign a different `SignalLength` to `fb`.

Properties

SignalLength — Length of the signal

1024 (default) | positive integer

Length of the signal, specified as a positive integer. The signal must have at least four samples.

Example: `'SignalLength', 1700`

Data Types: `double`

Wavelet — Analysis wavelet

'Morse' (default) | 'amor' | 'bump'

Analysis wavelet used in the filter bank, specified as `'Morse'`, `'amor'`, or `'bump'`. These strings specify the analytic Morse, Morlet (Gabor), and bump wavelet, respectively. The default wavelet is the analytic Morse (3,60) wavelet.

By default, for Morse wavelets, the frequency response decays to 50% of the peak magnitude at the Nyquist. For the Morlet and bump wavelets, the frequency response decays to 10% of the peak magnitude. You can change the decay percentage by setting the filter bank `FrequencyLimits` property. See `cwtfreqbounds`.

For Morse wavelets, you can also parametrize the wavelet using the `TimeBandwidth` or `WaveletParameters` properties.

Example: `'Wavelet', 'bump'`

VoicesPerOctave — Number of voices per octave

10 (default) | even integer between 4 and 48

Number of voices per octave to use for the CWT, specified as an even integer from 4 to 48. The CWT scales are discretized using the specified number of voices per octave. The energy spread of the wavelet in frequency and time automatically determines the minimum and maximum scales.

You can use `cwtfreqbounds` to determine the frequency limits of the wavelet filter bank. The frequency limits depend on parameters such as the energy spread of the wavelet, number of voices per octave, signal length, and sampling frequency.

Example: `'VoicesPerOctave', 20`

Data Types: `single` | `double`

SamplingFrequency — Sampling frequency in hertz

1 (default) | positive scalar

Sampling frequency in hertz, specified as a positive scalar. If unspecified, frequencies are in cycles/sample and the Nyquist frequency is $\frac{1}{2}$. To specify scales in periods, use the `SamplingPeriod` and `PeriodLimits` properties.

You cannot specify both the `SamplingFrequency` and `SamplingPeriod` properties.

Example: `'SamplingFrequency', 5`

Data Types: `single` | `double`

FrequencyLimits — Frequency limits

two-element scalar vector

Frequency limits of the wavelet filter bank, specified as a two-element vector with positive strictly increasing entries. The first element specifies the lowest peak passband frequency and must be greater than or equal to the product of the wavelet peak frequency in hertz and two time standard deviations divided by the signal length. The base 2 logarithm of the ratio of maximum frequency to minimum frequency must be greater than or equal to $1/NV$ where NV is the number of voices per octave. The high frequency limit must be less than or equal to the Nyquist.

If you specify frequency limits outside the permissible range, `cwtfilterbank` truncates the limits to the minimum and maximum values. Use `cwtfreqbounds` to determine frequency limits for different parametrizations of the wavelet transform.

If using a sampling period in the filter bank, you cannot specify the `FrequencyLimits` property.

Example: `'SamplingFrequency',20,'FrequencyLimits',[1 5]`

Data Types: `double`

SamplingPeriod — Sampling period

duration scalar

Sampling period, specified as a scalar duration. You cannot specify both the `SamplingFrequency` and `SamplingPeriod` properties.

Example: `'SamplingPeriod',seconds(0.5)`

Data Types: `duration`

PeriodLimits — Period limits

two-element duration array

Period limits of the wavelet filter bank, specified as a two-element `duration` array with positive strictly increasing entries. The first element of `PeriodLimits` specifies the largest peak passband frequency and must be greater than or equal to twice the `SamplingPeriod`. The base 2 logarithm of the ratio of the minimum period to the maximum period must be less than or equal to $-1/NV$, where NV is the number of voices per octave. The maximum period cannot exceed the signal length divided by the product of two time standard deviations of the wavelet and the wavelet peak frequency.

If you specify period limits outside the permissible range, `cwtfilterbank` truncates the limits to the minimum and maximum values. Use `cwtfreqbounds` to determine period limits for different parametrizations of the wavelet transform.

If using a sampling frequency in the filter bank, you cannot specify the `PeriodLimits` property.

Example: `'SamplingPeriod',seconds(0.1),'PeriodLimits',[seconds(0.2) seconds(1)]`

Data Types: `duration`

TimeBandwidth — Time-bandwidth product for Morse wavelets

60 (default) | positive scalar

Time-bandwidth product for Morse wavelets, specified as a positive scalar. This property is only valid when the `Wavelet` property is `'morse'`. This property specifies the time-bandwidth product of the Morse wavelet with the symmetry parameter (γ) fixed at 3. `TimeBandwidth` is a positive number strictly greater than 3 and less than or equal to 120.

The larger the time-bandwidth product, the more spread out the wavelet is in time and narrower the wavelet is in frequency. The standard deviation of the Morse wavelet in time is approximately $\sqrt{\text{TimeBandwidth}/2}$. The standard deviation in frequency is approximately $1/2 \cdot \sqrt{2/\text{TimeBandwidth}}$. See “Generalized Morse and Analytic Morlet Wavelets” on page 1-187.

The `TimeBandwidth` and `WaveletParameters` properties cannot both be specified.

In the notation of “Morse Wavelets”, `TimeBandwidth` is P^2 .

Example: `'TimeBandwidth', 20`

Data Types: `double`

WaveletParameters — Morse wavelet parameters

(3, 60) (default) | two-element vector of scalars

Morse wavelet parameters, specified as a two-element vector. The first element is the symmetry parameter (γ), which must be greater than or equal to 1. The second element is the time-bandwidth product, which must be strictly greater than γ . The ratio of the time-bandwidth product to γ cannot exceed 40.

When γ is equal to 3, the Morse wavelet is perfectly symmetric in the frequency domain. The skewness is equal to 0. Values of γ greater than 3 result in positive skewness, while values of γ less than 3 result in negative skewness.

`WaveletParameters` is only valid if the `Wavelet` property is set to `'Morse'`.

The `WaveletParameters` and `TimeBandwidth` properties cannot both be specified.

Example: `'WaveletParameters', [4, 20]`

Boundary — Boundary extension

'reflection' (default) | 'periodic'

Boundary extension of signal, specified as either 'reflection' or 'periodic'. Determines how the data is treated at the boundary.

Example: 'Boundary', 'periodic'

Object Functions

wt	Continuous wavelet transform with a filter bank
freqz	CWT filter bank frequency responses
wavelets	CWT filter bank time-domain wavelets
scales	CWT filter bank scales
waveletsupport	CWT filter bank time supports
qfactor	CWT filter bank quality factor
powerbw	CWT filter bank 3-dB bandwidths
centerFrequencies	CWT filter bank bandpass center frequencies
centerPeriods	CWT filter bank bandpass center periods

Examples

Continuous Wavelet Transform Filter Bank

Create a continuous wavelet transform filter bank.

```
fb = cwtfilterbank
```

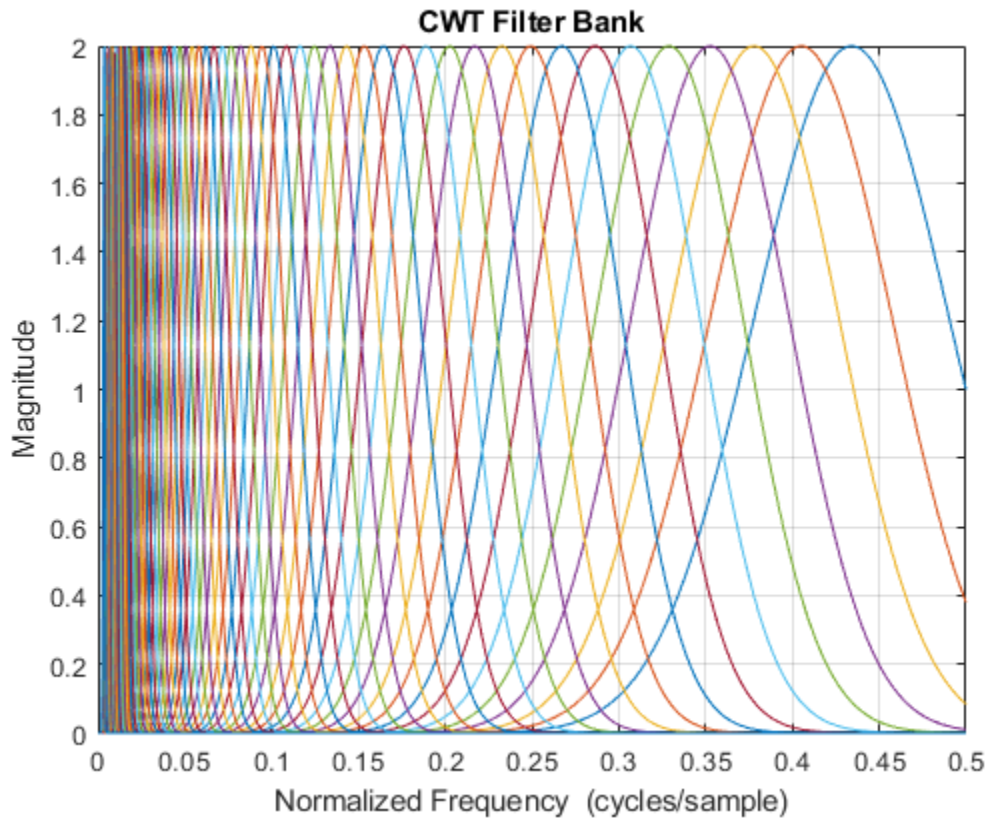
```
fb =
```

```
cwtfilterbank with properties:
```

```
    VoicesPerOctave: 10
           Wavelet: 'Morse'
    SamplingFrequency: 1
           SamplingPeriod: []
           PeriodLimits: []
           SignalLength: 1024
           FrequencyLimits: []
           TimeBandwidth: 60
    WaveletParameters: []
           Boundary: 'reflection'
```

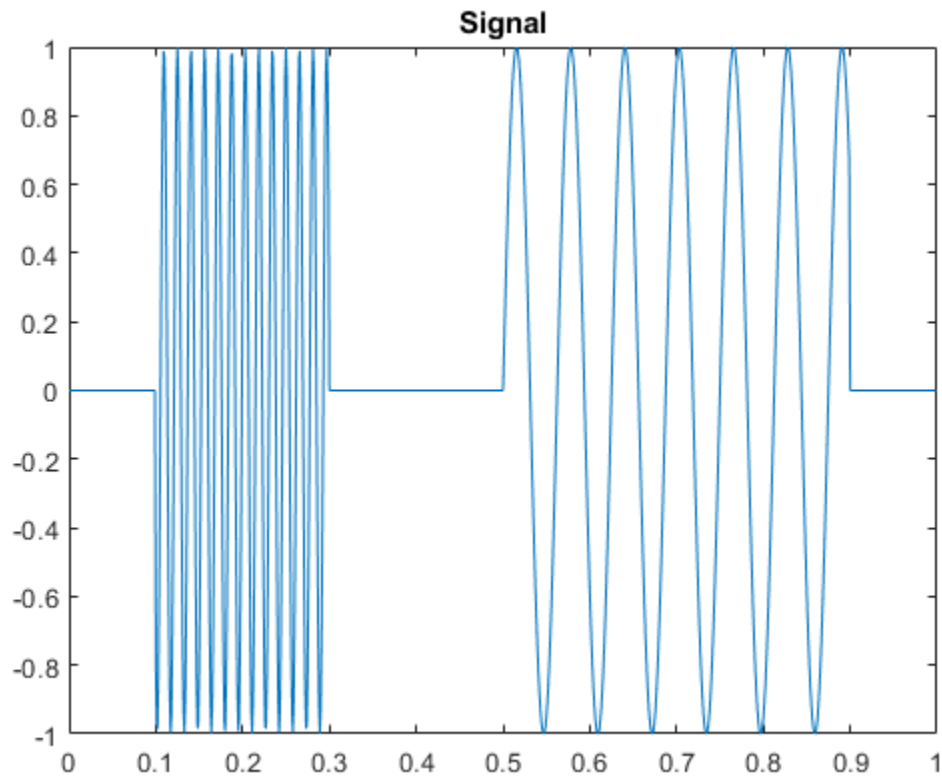
Plot the magnitude frequency response.

freqz(fb)

**Frequency Resolution of Continuous Wavelet Transform Filter Banks**

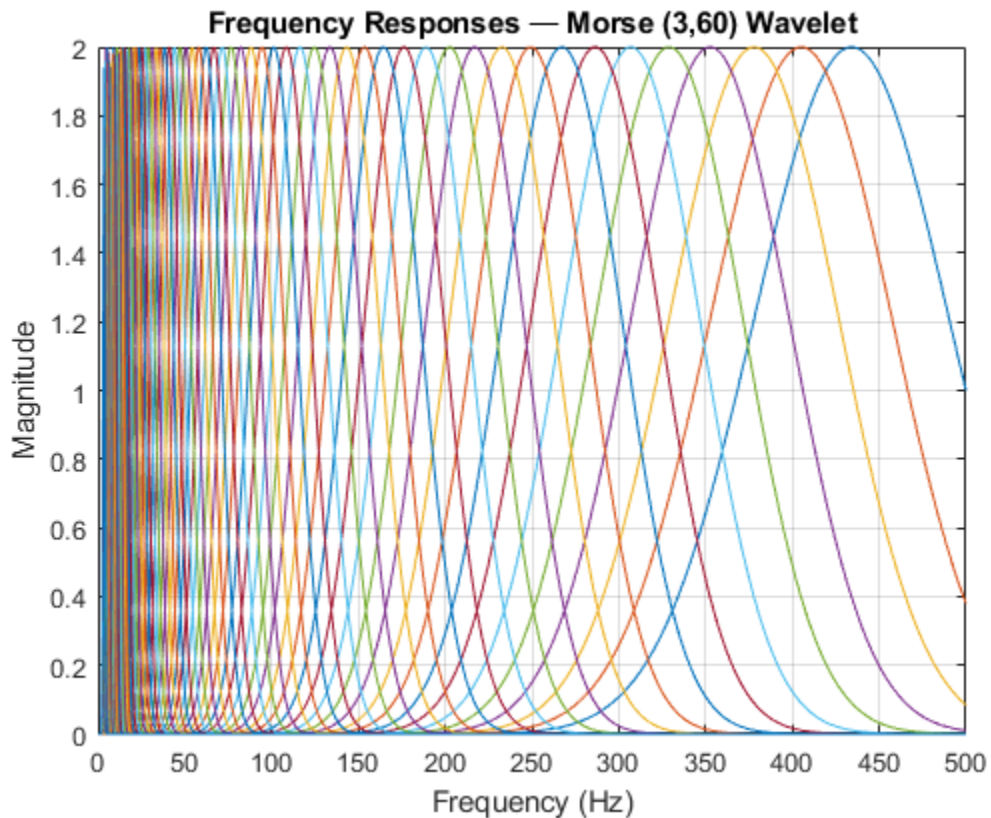
Create two sine waves with frequencies of 16 and 64 Hz. The data is sampled at 1000 Hz. Plot the signal.

```
Fs = 1e3;  
t = 0:1/Fs:1-1/Fs;  
x = cos(2*pi*64*t).*(t>=0.1 & t<0.3)+sin(2*pi*16*t).*(t>=0.5 & t<0.9);  
plot(t,x)  
title('Signal')
```

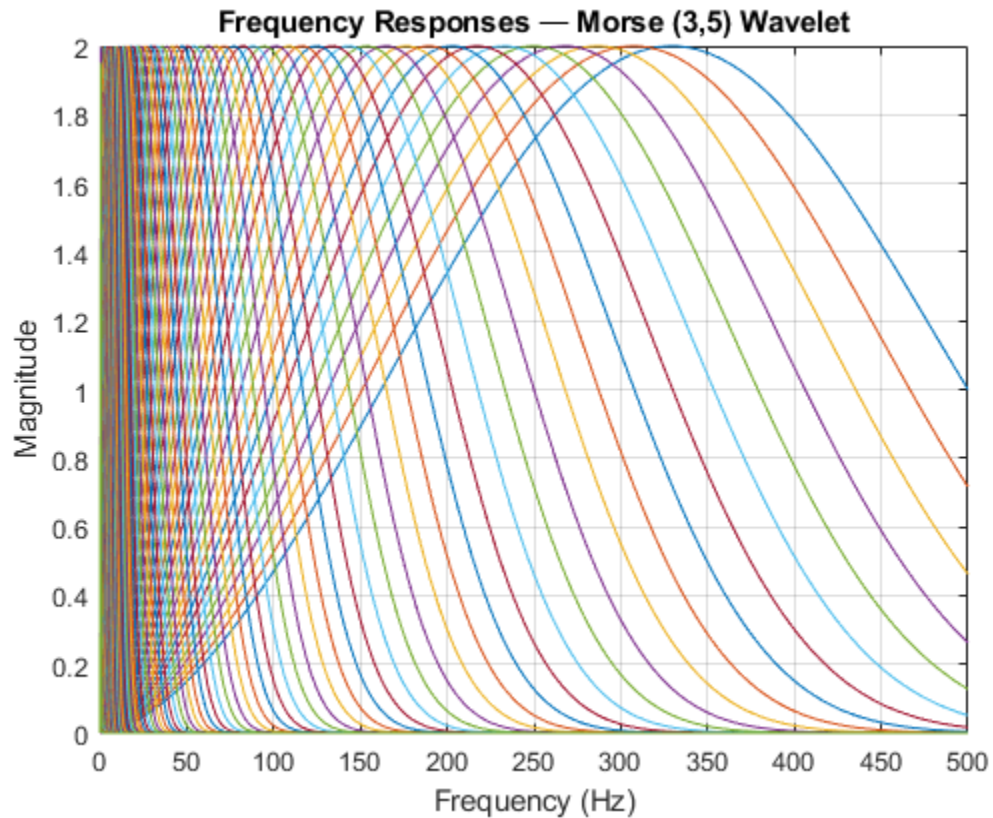
Create a CWT filter bank for the signal. Plot the frequency responses of the wavelets in the filter bank.

```
fb = cwtfilterbank('SignalLength', numel(t), 'SamplingFrequency', Fs);  
freqz(fb)  
title('Frequency Responses – Morse (3,60) Wavelet')
```



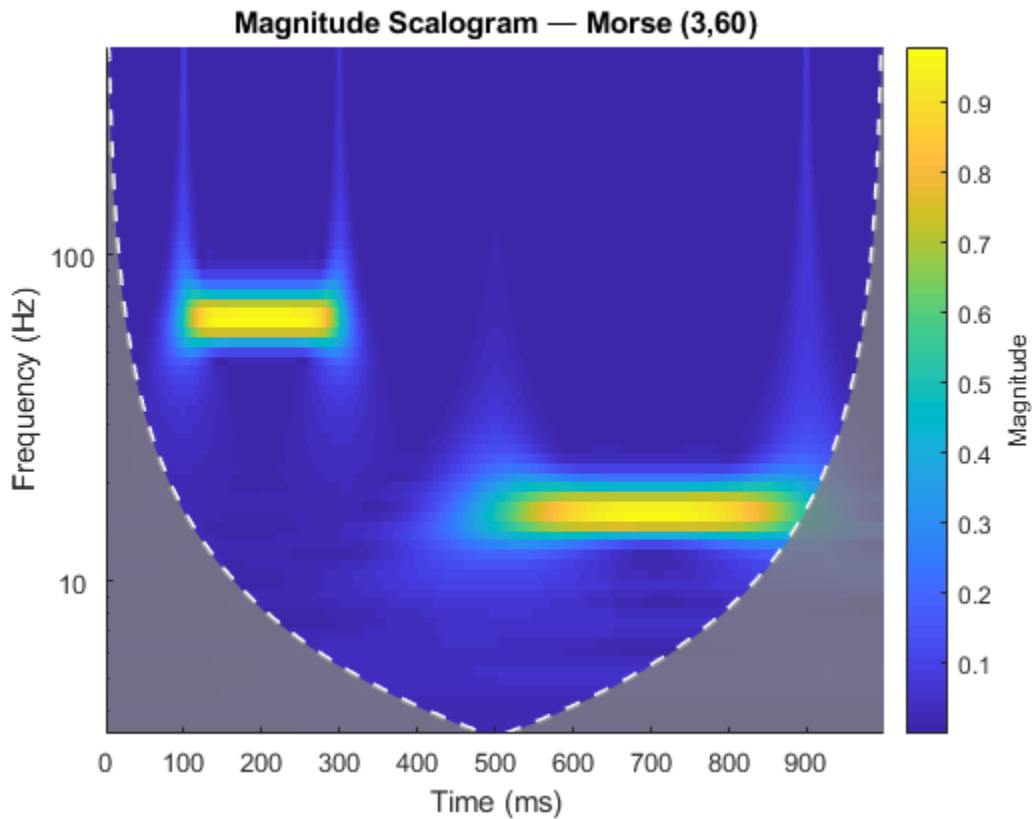
The analytic Morse (3,60) wavelet is the default wavelet in the filter bank. The wavelet has a time-bandwidth product equal to 60. Create a second filter bank identical to the first filter bank but instead uses the analytic Morse (3,5) wavelet. Plot the frequency responses of the wavelets in the second filter bank.

```
fb3x5 = cwtfilterbank('SignalLength', numel(t), 'SamplingFrequency', Fs, ...
    'TimeBandwidth', 5);
figure
freqz(fb3x5)
title('Frequency Responses - Morse (3,5) Wavelet')
```

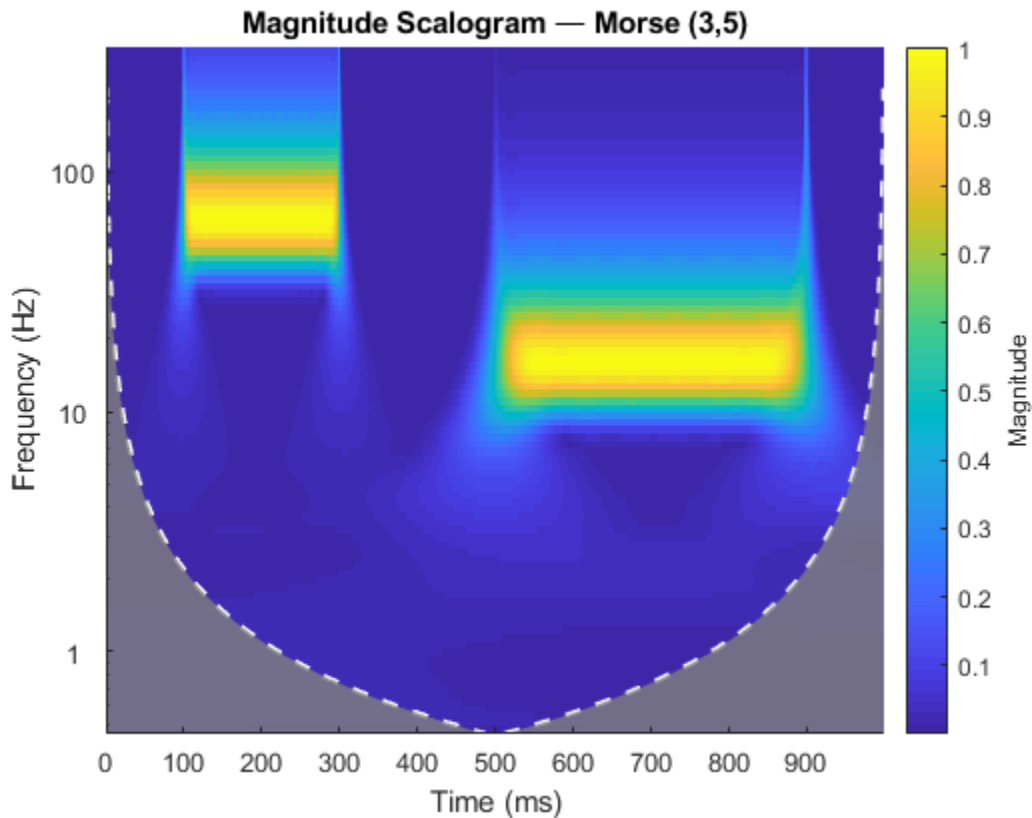


Observe that the frequency responses are wider than in the first filter bank. The Morse (3,60) wavelet is better localized in frequency than the Morse (3,5) wavelet. Apply each filter bank to the signal and plot the resulting scalograms. Observe that the Morse (3,60) wavelet has better frequency resolution than the Morse (3,5) wavelet.

```
figure
cwt(x,'FilterBank',fb)
title('Magnitude Scalogram - Morse (3,60)')
```



```
figure  
cwt(x,'FilterBank',fb3x5)  
title('Magnitude Scalogram - Morse (3,5)')
```

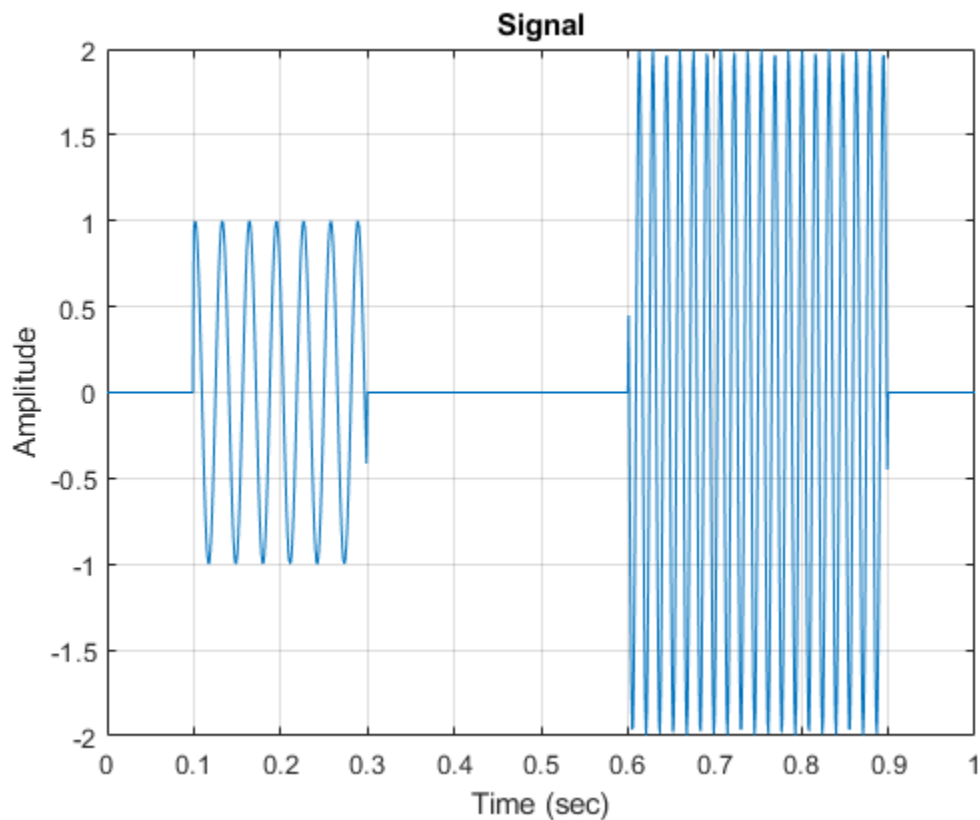


Sinusoid and Wavelet Coefficient Amplitudes

This example shows that the amplitudes of oscillatory components in a signal agree with the amplitudes of the corresponding wavelet coefficients.

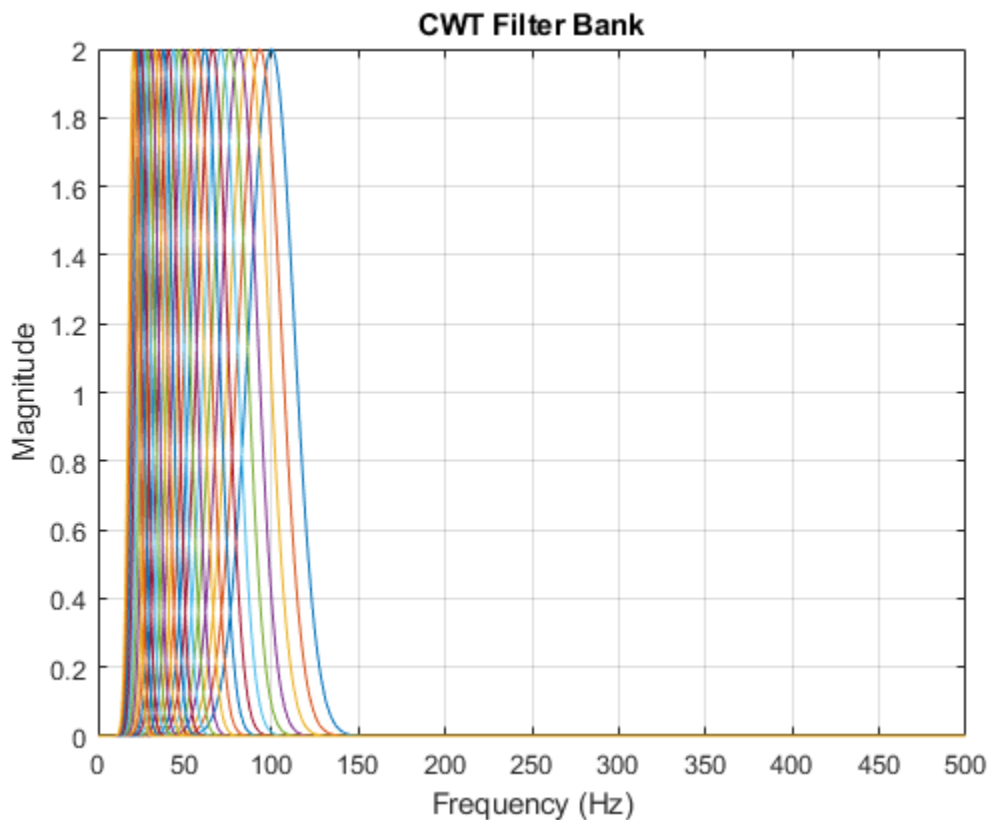
Create a signal composed of two sinusoids with disjoint support in time. One sinusoid has a frequency of 32 Hz and amplitude equal to 1. The other sinusoid has a frequency of 64 Hz and amplitude equal to 2. The signal is sampled for one second at 1000 Hz. Plot the signal.

```
frq1 = 32;  
amp1 = 1;  
frq2 = 64;  
amp2 = 2;  
  
Fs = 1e3;  
t = 0:1/Fs:1;  
x = amp1*sin(2*pi*frq1*t).*(t>=0.1 & t<0.3)+amp2*sin(2*pi*frq2*t).*(t>0.6 & t<0.9);  
  
plot(t,x)  
grid on  
xlabel('Time (sec)')  
ylabel('Amplitude')  
title('Signal')
```



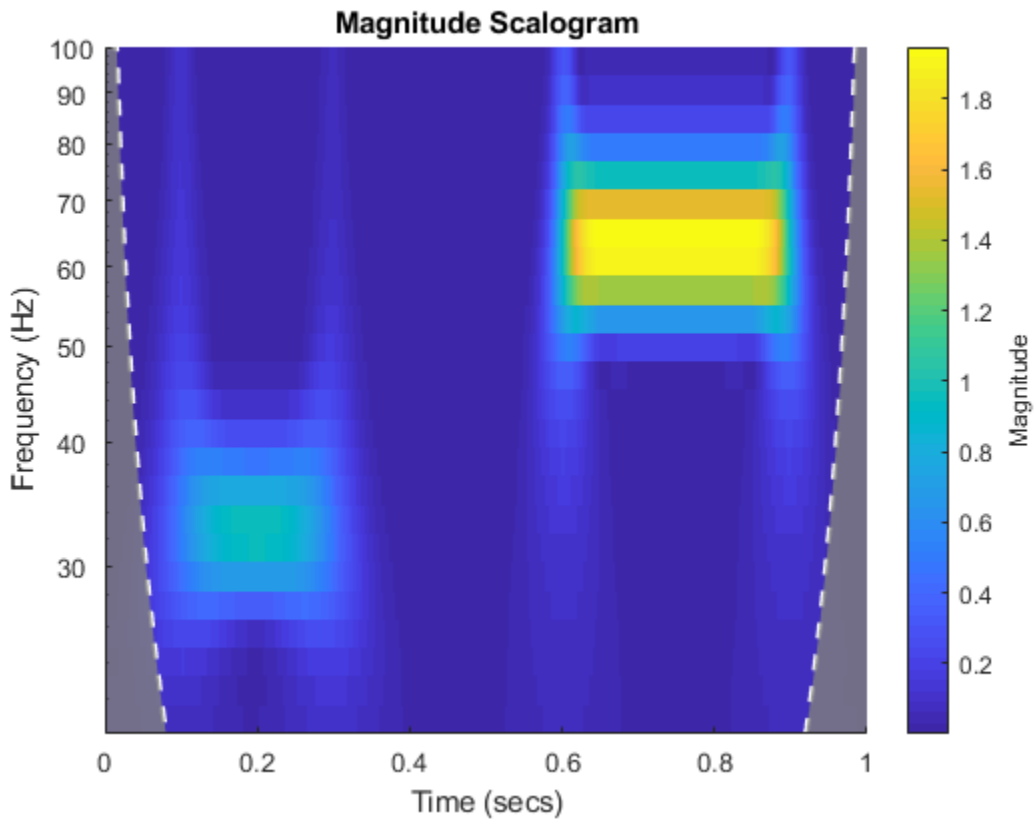
Create a CWT filter bank that can be applied to the signal. Since the signal component frequencies are known, set the frequency limits of the filter bank to a narrow range that includes the known frequencies. To confirm the range, plot the magnitude frequency responses for the filter bank.

```
fb = cwtfilterbank('SignalLength',numel(x),'SamplingFrequency',Fs,...  
    'FrequencyLimits',[20 100]);  
figure  
freqz(fb)
```

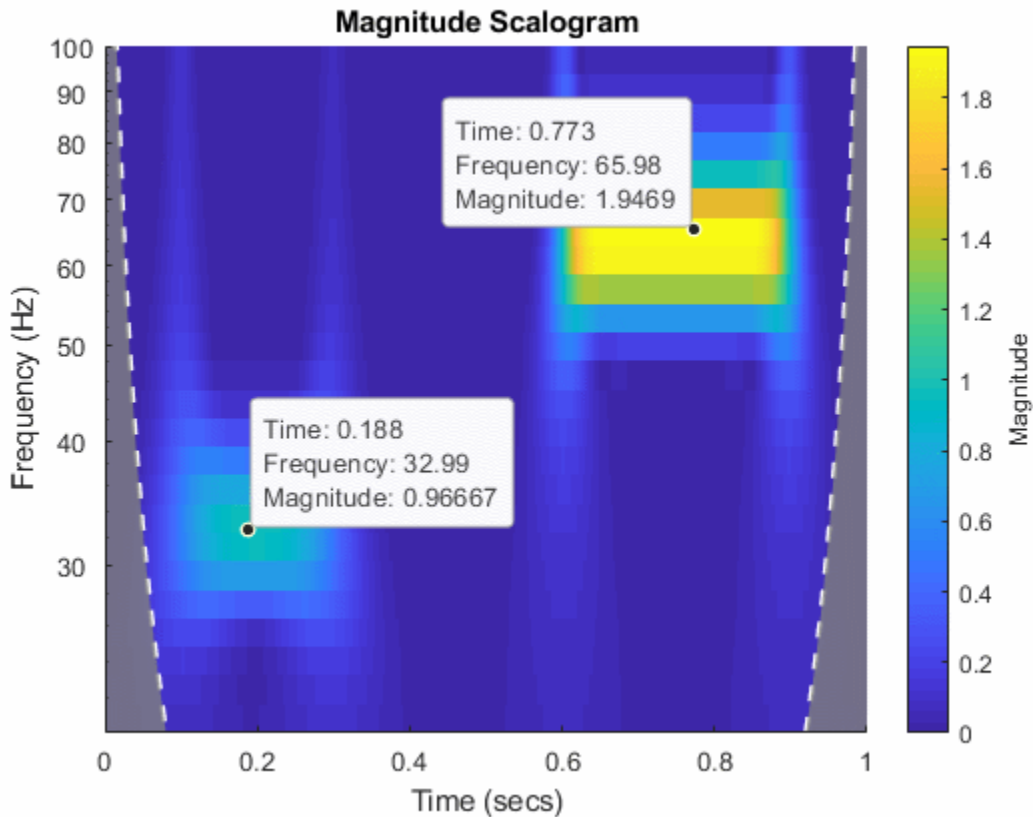


Use `cwt` and the filter bank to plot the scalogram of the signal.

```
figure  
cwt(x,'FilterBank',fb)
```



Execute this script and use a data cursor to confirm that the amplitudes of the wavelet coefficients are essentially equal to the amplitudes of the sinusoidal components.



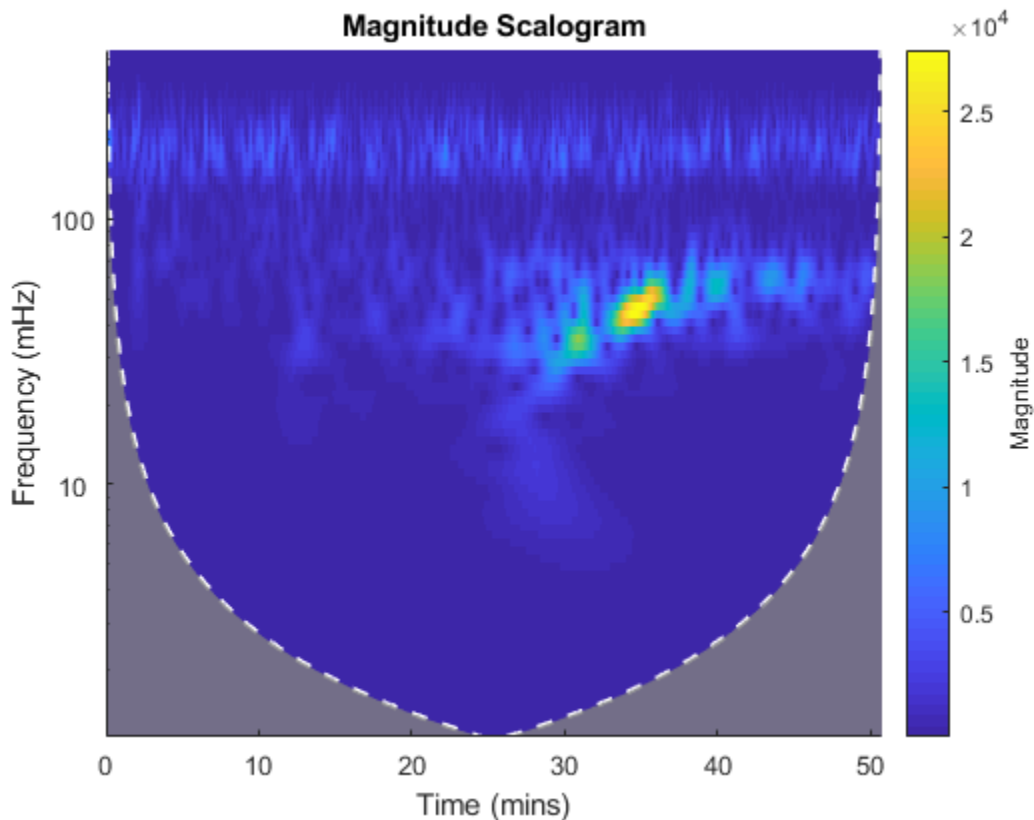
Generalized Morse and Analytic Morlet Wavelets

This example shows how to vary the time-bandwidth parameter of the generalized Morse wavelet to approximate the analytic Morlet wavelet.

Generalized Morse wavelets are a family of exactly analytic wavelets. Morse wavelets have two parameters, symmetry and time-bandwidth product. You can vary these parameters to obtain analytic wavelets with different properties and behaviors. For additional information, see “Morse Wavelets” and the references therein.

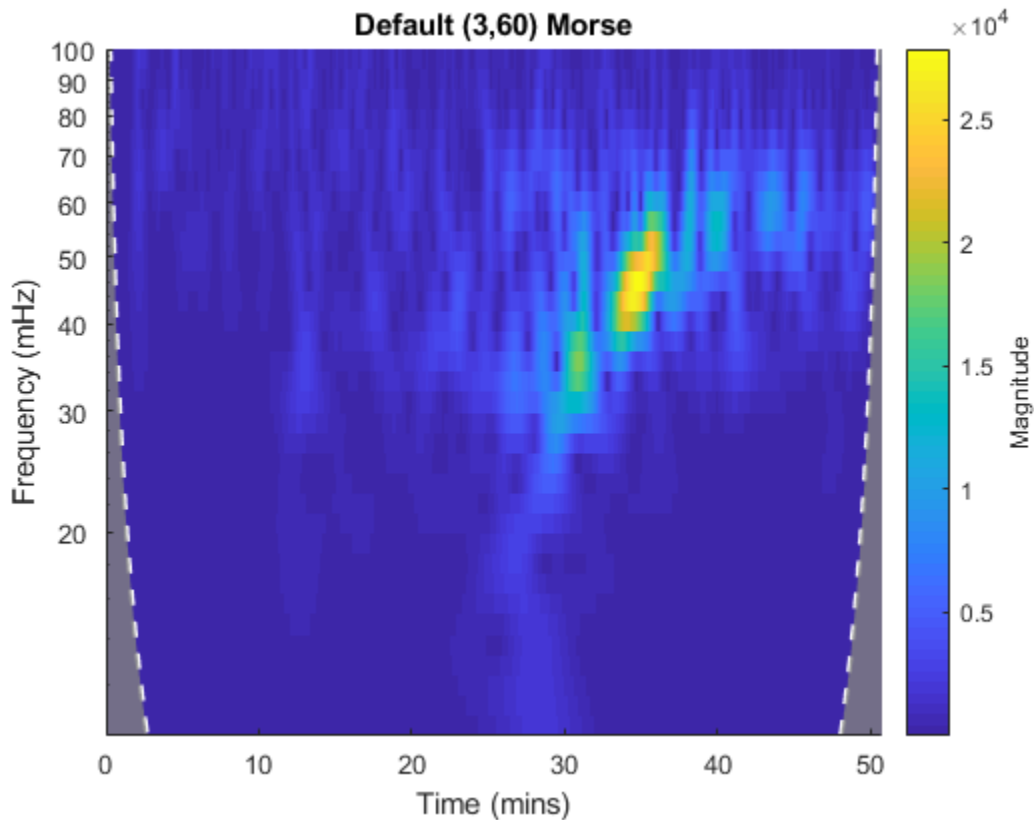
Load the seismograph data recorded during the 1995 Kobe earthquake. The data are seismograph (vertical acceleration, nm/sq.sec) measurements recorded at Tasmania University, Hobart, Australia on 16 January 1995 beginning at 20:56:51 (GMT) and continuing for 51 minutes at 1 second intervals. Create a CWT filter bank with default settings that can be applied to the data. Use the filter bank to generate the scalogram.

```
load kobe
fb = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1);
cwt(kobe, 'FilterBank', fb)
```



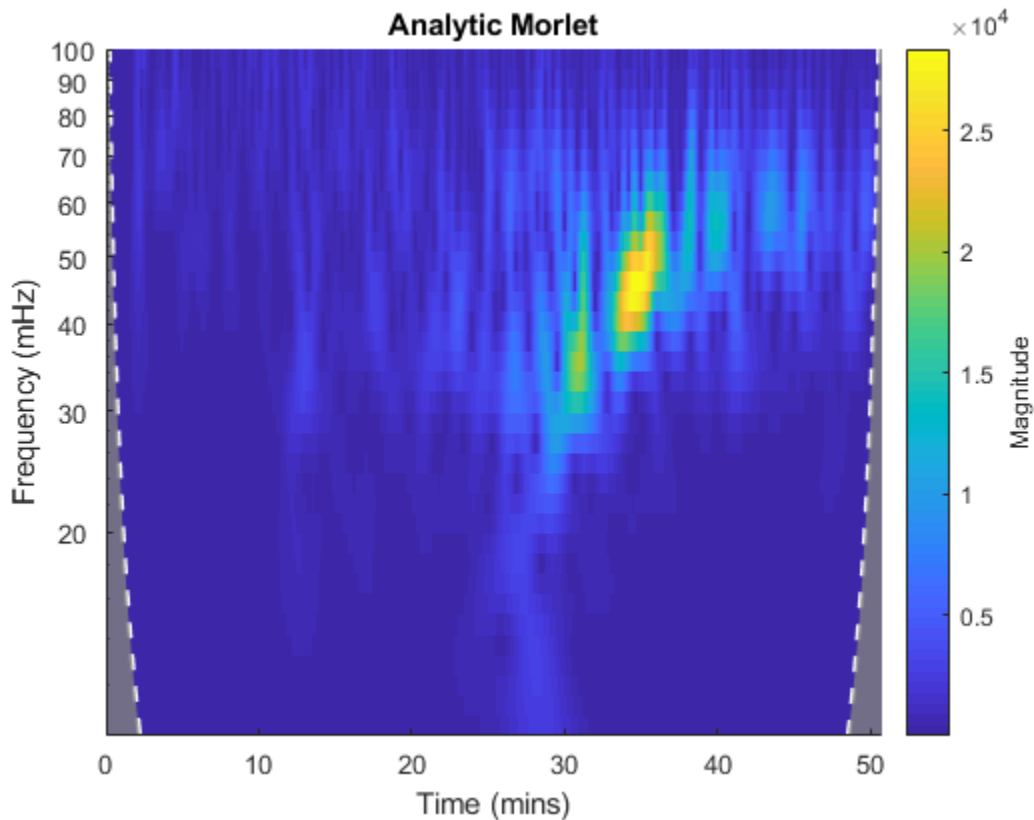
The magnitude of the wavelet coefficients is large in the frequency range from 10 mHz to 100 mHz. Create a new filter bank with frequency limits set to these values. Generate the scalogram.

```
fb2 = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1, ...
    'FrequencyLimits', [1e-2 1e-1]);
cwt(kobe, 'FilterBank', fb2)
title('Default (3,60) Morse')
```



By default, `cwtfilterbank` uses the (3,60) Morse wavelet. Create a filter bank using the analytic Morlet wavelet with the same frequency limits. Generate a scalogram and compare with the scalogram generated by the (3,60) Morse wavelet.

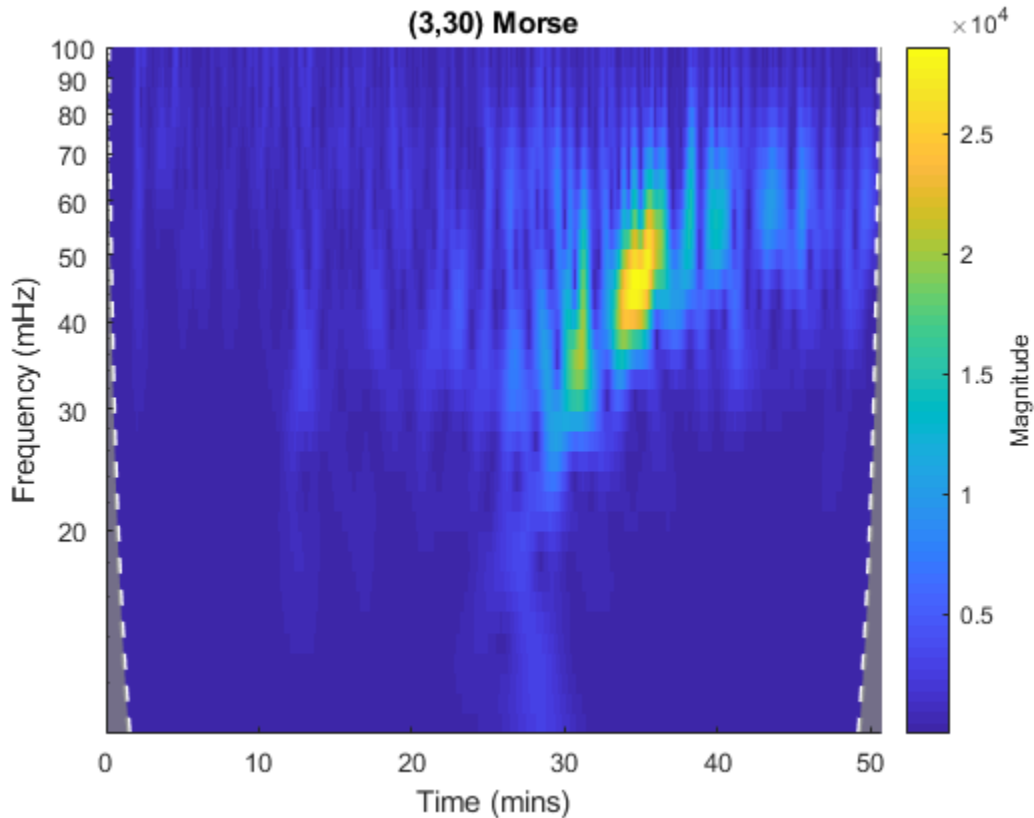
```
fbMorlet = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1, ...
    'FrequencyLimits', [1e-2 1e-1], ...
    'Wavelet', 'amor');
cwt(kobe, 'FilterBank', fbMorlet)
title('Analytic Morlet')
```



The Morlet wavelet is not as well localized in frequency as the (3,60) Morse wavelet. However, by varying the time-bandwidth product, you can create a Morse wavelet with properties similar to the Morlet wavelet.

Create a filter bank using the Morse wavelet with a time-bandwidth value of 30 [2], with frequency limits as above. Generate the scalogram of the seismograph data. Note there is smearing in frequency nearly identical to the Morlet results.

```
fbMorse = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1, ...
    'FrequencyLimits', [1e-2 1e-1], ...
    'TimeBandwidth', 30);
cwt(kobe, 'FilterBank', fbMorse)
title('(3,30) Morse')
```



Now examine the wavelets associated with the `fbMorlet` and `fbMorse` filter banks. From both filter banks, obtain the wavelet center frequencies, filter frequency responses, and time-domain wavelets. Confirm the center frequencies are nearly identical.

```
cfMorlet = centerFrequencies(fbMorlet);
[frMorlet, fMorlet] = freqz(fbMorlet);
[wvMorlet, tMorlet] = wavelets(fbMorlet);
cfMorse = centerFrequencies(fbMorse);
[frMorse, fMorse] = freqz(fbMorse);
[wvMorse, tMorse] = wavelets(fbMorse);

disp(['Number of Center Frequencies: ', num2str(length(cfMorlet))]);

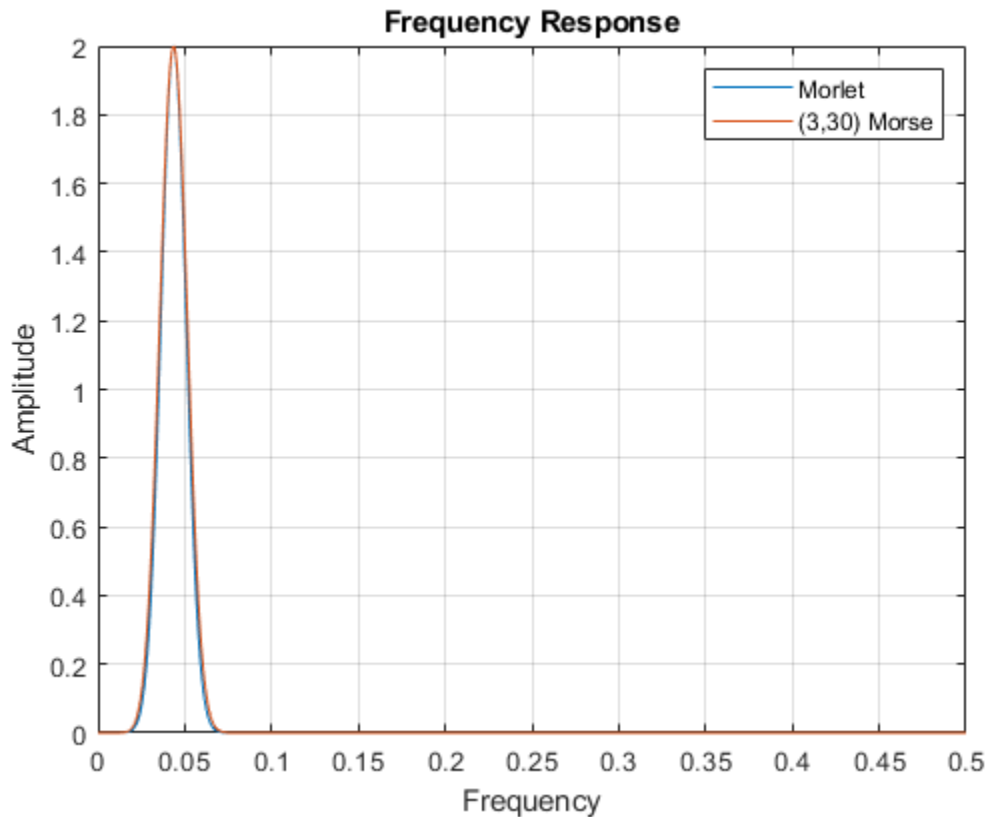
Number of Center Frequencies: 34
```

```
disp(['Maximum difference: ', num2str(max(abs(cfMorlet-cfMorse)))]);
```

```
Maximum difference: 2.7756e-17
```

Each filter bank contains the same number of wavelets. Choose a center frequency, and plot the frequency response of the associated filter from each filter bank. Confirm the responses are nearly identical.

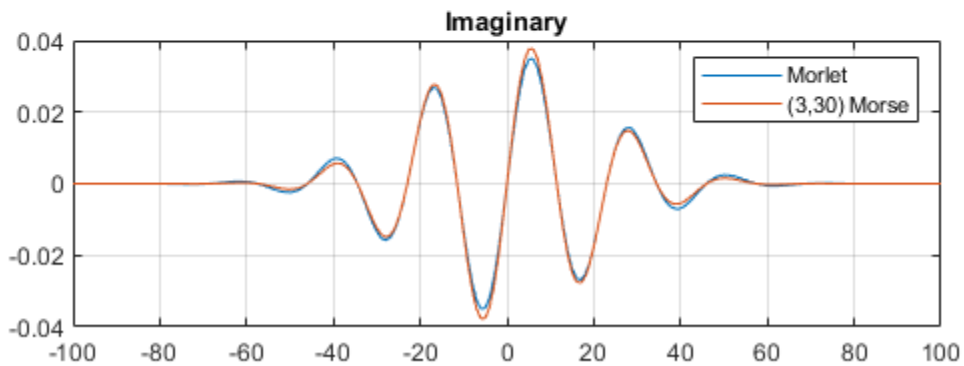
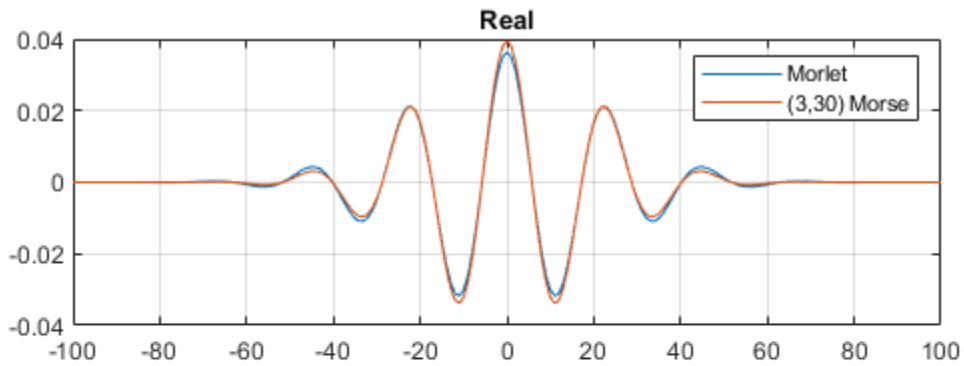
```
wv = 13;
figure
plot(fMorlet, frMorlet(wv, :));
hold on
plot(fMorse, frMorse(wv, :));
grid on
title('Frequency Response')
xlabel('Frequency')
ylabel('Amplitude')
legend('Morlet', '(3,30) Morse')
```



Plot the time-domain wavelets associated with the same center frequency. Confirm they are nearly identical.

```
figure
subplot(2,1,1)
plot(tMorlet,real(wvMorlet(wv,:)))
hold on
plot(tMorse,real(wvMorse(wv,:)))
grid on
title('Real')
legend('Morlet','(3,30) Morse')
xlim([-100 100])
subplot(2,1,2)
plot(tMorlet,imag(wvMorlet(wv,:)))
```

```
hold on
plot(tMorse, imag(wvMorse(wv, :)))
grid on
title('Imaginary')
legend('Morlet', '(3,30) Morse')
xlim([-100 100])
```



Changing the Time-bandwidth Product

This example shows that increasing the time-bandwidth product P^2 of the Morse wavelet creates a wavelet with more oscillations under its envelope. All other things being equal, increasing P^2 makes the wavelet more frequency-localized as a result.

Create two filter banks. One filter bank has the default TimeBandwidth value of 60. The second filter bank has a TimeBandwidth value of 10. The SignalLength for both filter banks is 4096 samples.

```
sigLen = 4096;
fb60 = cwtfilterbank('SignalLength',sigLen);
fb10 = cwtfilterbank('SignalLength',sigLen,'TimeBandwidth',10);
```

Obtain the time-domain wavelets for the filter banks.

```
[psi60,t] = wavelets(fb60);
[psi10,~] = wavelets(fb10);
```

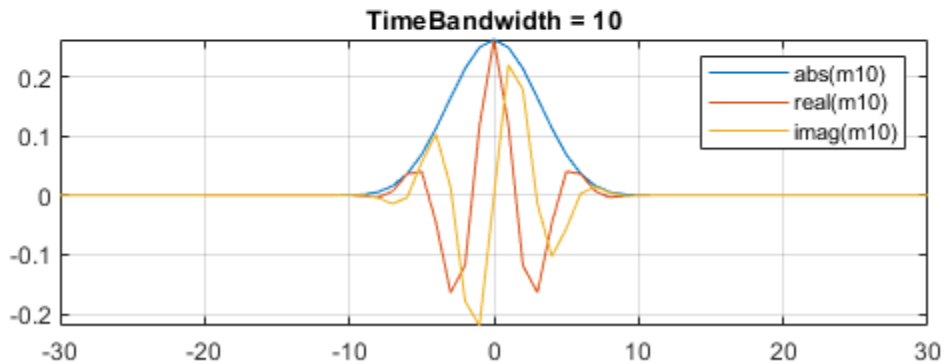
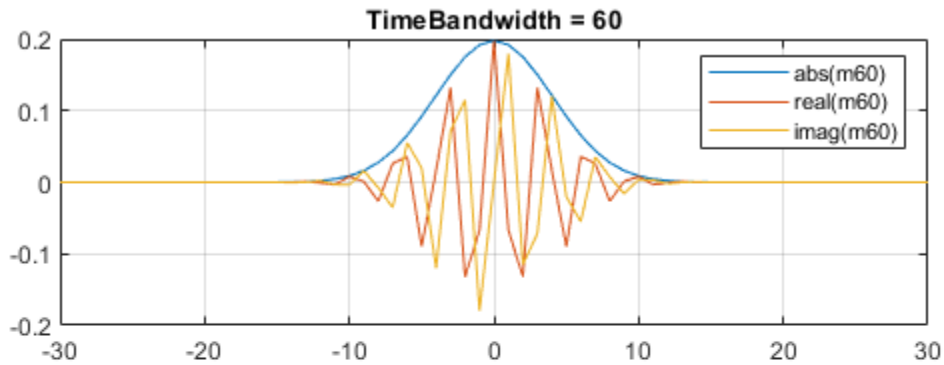
Use the scales function to find the mother wavelet for each filter bank.

```
sca60 = scales(fb60);
sca10 = scales(fb10);
[~,idx60] = min(abs(sca60-1));
[~,idx10] = min(abs(sca10-1));
m60 = psi60(idx60,:);
m10 = psi10(idx10,:);
```

Since the time-bandwidth product is larger for the fb60 filter bank, verify the m60 wavelet has more oscillations under its envelope than the m10 wavelet.

```
subplot(2,1,1)
plot(t,abs(m60))
grid on
hold on
plot(t,real(m60))
plot(t,imag(m60))
xlim([-30 30])
legend('abs(m60)','real(m60)','imag(m60)')
title('TimeBandwidth = 60')
subplot(2,1,2)
plot(t,abs(m10))
grid on
hold on
```

```
plot(t,real(m10))
plot(t,imag(m10))
xlim([-30 30])
legend('abs(m10)','real(m10)','imag(m10)')
title('TimeBandwidth = 10')
```

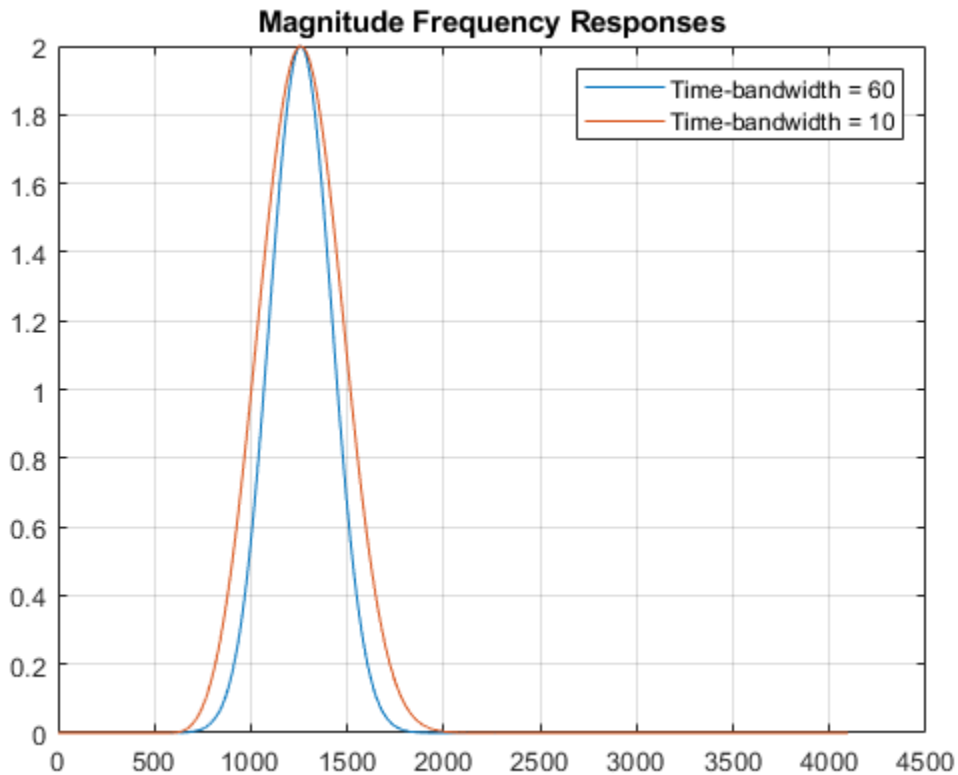


Align the peaks of the $m60$ and $m10$ magnitude frequency responses. Verify the frequency response of the $m60$ wavelet is more narrow than the frequency response for the $m10$ wavelet.

```
cf60 = centerFrequencies(fb60);
cf10 = centerFrequencies(fb10);

m60cFreq = cf60(idx60);
```

```
m10cFreq = cf10(idx10);  
  
freqShift = 2*pi*(m60cFreq-m10cFreq);  
x10 = m10.*exp(1j*freqShift*(-sigLen/2:sigLen/2-1));  
  
figure  
plot([abs(fft(m60)).' abs(fft(x10)).'])  
grid on  
legend('Time-bandwidth = 60','Time-bandwidth = 10')  
title('Magnitude Frequency Responses')
```



Using CWT Filter Bank on Multiple Time Series

This example shows how using a CWT filter bank improves computational efficiency when taking the CWT of multiple time series.

Load the seismograph data recorded during the 1995 Kobe earthquake. The data are seismograph (vertical acceleration, nm/sq.sec) measurements recorded at Tasmania University, Hobart, Australia on 16 January 1995 beginning at 20:56:51 (GMT) and continuing for 51 minutes at 1 second intervals. Create a CWT filter bank that can be applied to the data.

```
load kobe
fb = cwtfilterbank('SignalLength', numel(kobe), 'SamplingFrequency', 1);
```

Use the `cwt` function and take the CWT of the data 250 times. Display the elapsed time used.

```
num = 250;
tic;
for k=1:num
    cfs = cwt(kobe);
end
toc
```

Elapsed time is 6.551628 seconds.

Now use the `wt` object function of the filter bank to take the CWT of the data. Confirm using the filter bank is faster.

```
tic;
for k=1:num
    cfs = wt(fb, kobe);
end
toc
```

Elapsed time is 3.782376 seconds.

Compatibility Considerations

BPfrequencies and BPperiods will be removed

Not recommended starting in R2018b

The `BPfrequencies` and `BPperiods` object functions of `cwtfilterbank` have been renamed `centerFrequencies` and `centerPeriods`, respectively. The functionality remains unchanged. `BPfrequencies` and `BPperiods` will be removed in a future release.

References

- [1] Lilly, J. M., and S. C. Olhede. "Generalized Morse Wavelets as a Superfamily of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 60, No. 11, 2012, pp. 6036-6041.
- [2] Lilly, J. M., and S. C. Olhede. "Higher-Order Properties of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 57, No. 1, 2009, pp. 146-160.
- [3] Lilly, J. M. *jLab: A data analysis package for Matlab*, version 1.6.2. 2016. <http://www.jmlilly.net/jmlsoft.html>.
- [4] Lilly, J. M. "Element analysis: a wavelet-based method for analysing time-localized events in noisy time series." *Proceedings of the Royal Society A*. Volume 473: 20160776, 2017, pp. 1-28. [dx.doi.org/10.1098/rspa.2016.0776](https://doi.org/10.1098/rspa.2016.0776).

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

The `cwtfilterbank` object supports C/C++ code generation. You must have MATLAB Coder to generate C/C++ code.

Usage notes and limitations:

- The following properties are not supported: `SamplingPeriod`, `PeriodLimits`.
- The following object functions support C/C++ code generation, with limitations as noted:
 - `wt`

- `freqz`: Plotting is not supported.
- `wavelets`
- `scales`
- `qfactor`
- `centerFrequencies`

See Also

`cwt` | `cwtfreqbounds` | `icwt`

Topics

“Boundary Effects and the Cone of Influence”

“Morse Wavelets”

Introduced in R2018a

cwtfreqbounds

CWT maximum and minimum frequency or period

Syntax

```
[minfreq,maxfreq] = cwtfreqbounds(N)
[minfreq,maxfreq] = cwtfreqbounds(N,Fs)
[maxperiod,minperiod] = cwtfreqbounds(N,Ts)
[ ___ ] = cwtfreqbounds( ___ ,Name,Value)
```

Description

[minfreq,maxfreq] = cwtfreqbounds(N) returns the minimum and maximum wavelet bandpass frequencies in cycles/sample for a signal of length N. The minimum and maximum frequencies are determined for the default Morse (3,60) wavelet. The minimum frequency is determined so that two time standard deviations of the default wavelet span the N-point signal at the coarsest scale. The maximum frequency is such that the highest frequency wavelet bandpass filter drops to ½ of its peak magnitude at the Nyquist frequency.

[minfreq,maxfreq] = cwtfreqbounds(N,Fs) returns the bandpass frequencies in hertz for the sampling frequency Fs.

[maxperiod,minperiod] = cwtfreqbounds(N,Ts) returns the bandpass periods for the sampling period Ts. maxperiod and minperiod are scalar durations with the same format as Ts. If the number of standard deviations is set so that $\log_2(\text{maxperiod}/\text{minperiod}) < 1/NV$ where NV is the number of voices per octave, maxperiod is adjusted to $\text{minperiod} * 2^{(1/NV)}$.

[___] = cwtfreqbounds(___ ,Name,Value) returns the minimum and maximum wavelet bandpass frequencies or periods with additional options specified by one or more Name,Value pair arguments.

Examples

Wavelet Bandpass Frequencies Using Default Values

Obtain the minimum and maximum wavelet bandpass frequencies for a signal with 1000 samples using the default values.

```
[minfreq,maxfreq] = cwtfreqbounds(1000)
minfreq = 0.0033
maxfreq = 0.4341
```

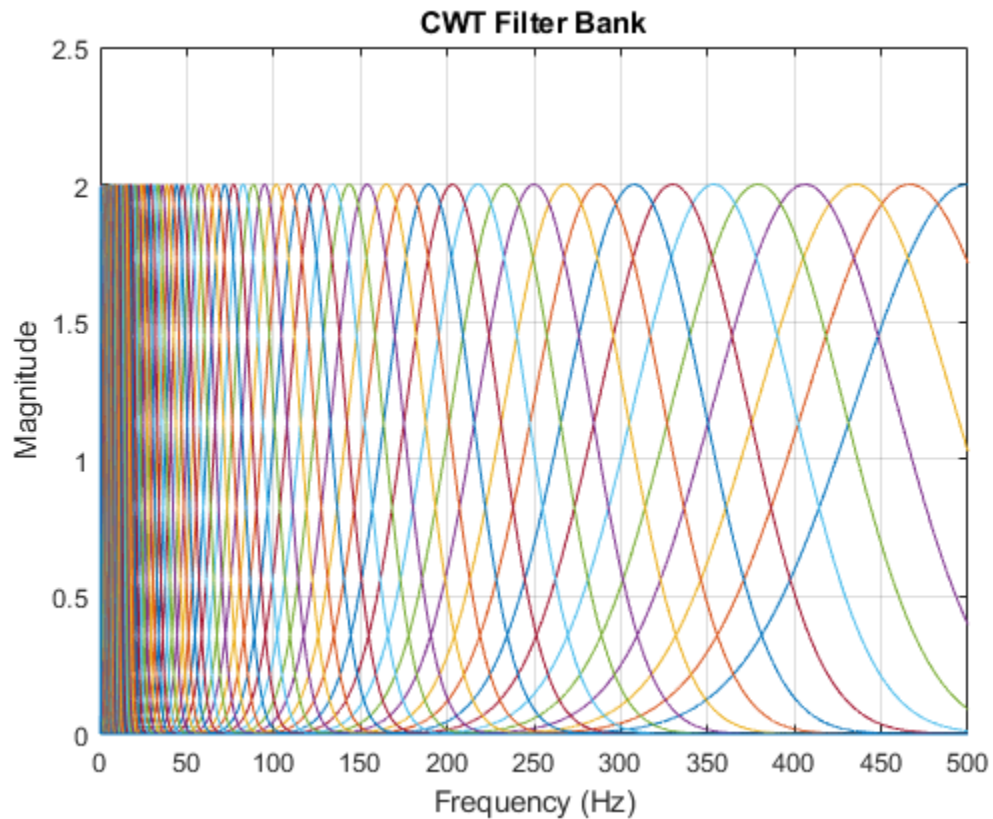
Construct CWT Filter Bank With Peak Magnitude at Nyquist

Obtain the minimum and maximum wavelet bandpass frequencies for the default Morse wavelet for a signal of length 10,000 and a sampling frequency of 1kHz. Set the cutoff to 100% so that the highest frequency wavelet bandpass filter peaks at the Nyquist.

```
sigLength = 10000;
Fs = 1e3;
[minfreq,maxfreq] = cwtfreqbounds(sigLength,Fs,'cutoff',100);
```

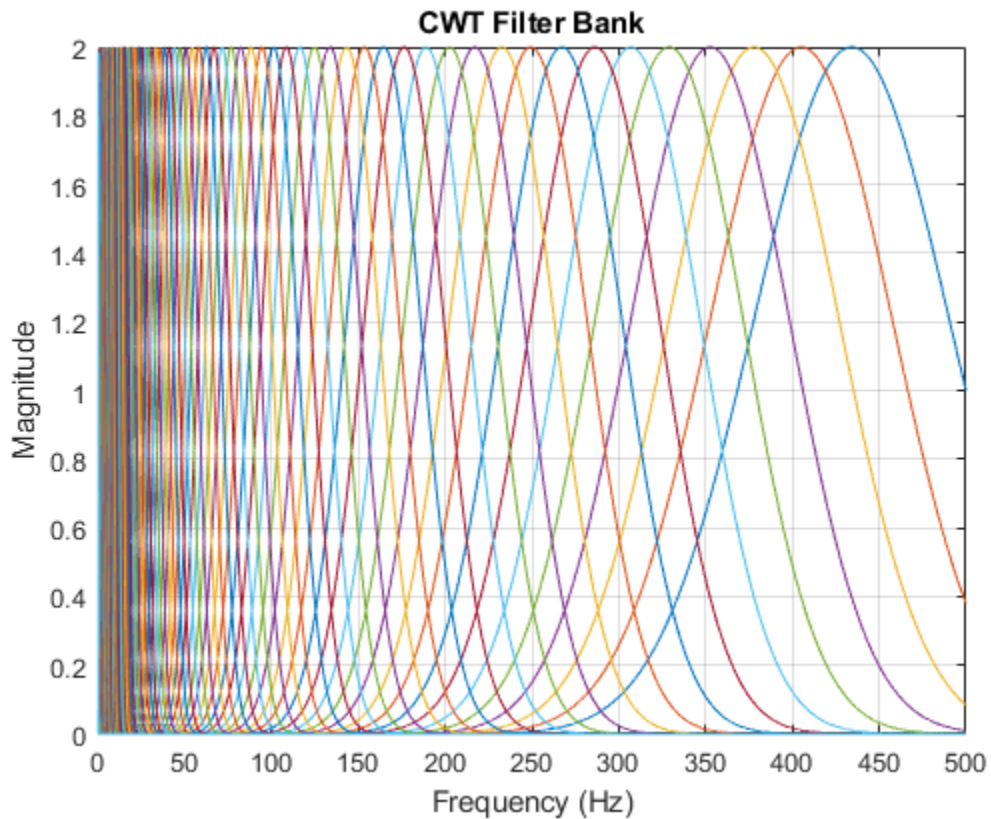
Construct the filter bank using the values returned by `cwtfreqbounds`. Plot the frequency response. Note that the highest frequency wavelet bandpass filter peaks at the Nyquist frequency of 500 Hz.

```
fb = cwtfilterbank('SignalLength',sigLength,'SamplingFrequency',Fs,'FrequencyLimits',[
freqz(fb)
```

Create a second frequency bank identical to the first, but instead use the default frequency limits. Plot the frequency response. In the case of the Morse wavelet, the CWT filter bank uses a default cutoff of 50% at the Nyquist.

```
fb2 = cwtfilterbank('SignalLength', sigLength, 'SamplingFrequency', Fs);  
figure  
freqz(fb2)
```



Input Arguments

N — Signal length

positive integer

Signal length, specified as a positive integer greater than or equal to 4.

Fs — Sampling frequency

positive scalar

Sampling frequency in hertz, specified as a positive scalar.

Data Types: double

Ts — Sampling period

scalar duration

Sampling period, specified as a positive scalar duration.

Data Types: duration

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

Example: `'Wavelet', 'amor', 'CUTOFF', 75`

Wavelet — Analysis wavelet

`'Morse'` (default) | `'amor'` | `'bump'`

Analysis wavelet used to determine the minimum and maximum frequencies or periods, specified as `'Morse'`, `'amor'`, or `'bump'`. These strings specify the analytic Morse, Morlet, and bump wavelet, respectively. The default wavelet is the analytic Morse (3,60) wavelet.

For Morse wavelets, you can also parametrize the wavelet using the `TimeBandwidth` or `WaveletParameters` name-value pairs.

Example: `'Wavelet', 'bump'`

CUTOFF — Percentage of the peak magnitude

50 for the Morse wavelet | 10 for the analytic Morlet and bump wavelets

Percentage of the peak magnitude at the Nyquist, specified as a scalar between 0 and 100. Setting `CUTOFF` to 0 indicates that the wavelet frequency response decays to 0 at the Nyquist. Setting `CUTOFF` to 100 indicates that the value of the wavelet bandpass filters peaks at the Nyquist.

Data Types: double

StandardDeviations — Number of time standard deviations

2 (default)

Number of time standard deviations used to determine the minimum frequency (longest scale), specified as a positive integer greater than or equal to 2. For the Morse, analytic Morlet, and bump wavelets, four standard deviations generally ensures that the wavelet decays to zero at the ends of the signal support. Incrementing 'StandardDeviations' by multiples of 4, for example $4 \cdot M$, ensures that M whole wavelets fit within the signal length. If unspecified, 'StandardDeviations' defaults to 2. If the number of standard deviations is set so that $\log_2(\text{minfreq}/\text{maxfreq}) > -1/NV$ where NV is the number of voices per octave, minfreq is adjusted to $\text{maxfreq} \cdot 2^{(-1/NV)}$.

Data Types: double

TimeBandwidth — Time-bandwidth for Morse wavelets

positive scalar

This property is read-only.

Time-bandwidth for Morse wavelets, specified as a positive scalar. This property is only valid when the `Wavelet` property is 'morse'. This property specifies the time-bandwidth parameter of the Morse wavelet with the symmetry parameter (γ) fixed at 3. `TimeBandwidth` is a positive number strictly greater than 3 and less than or equal to 120. The larger the time-bandwidth parameter, the more spread out the wavelet is in time and narrower the wavelet is in frequency. The standard deviation of the Morse wavelet in time is approximately $\sqrt{\text{TimeBandwidth}/2}$. The standard deviation in frequency is approximately $1/2 \cdot \sqrt{2/\text{TimeBandwidth}}$.

The `TimeBandwidth` and `WaveletParameters` properties cannot both be specified.

Example: 'TimeBandwidth',4

Data Types: double

WaveletParameters — Morse wavelet parameters

(3,60) (default) | two-element vector of scalars

This property is read-only.

Morse wavelet parameters, specified as a two-element vector. The first element is the symmetry parameter (γ), which must be greater than or equal to 1. The second element is the time-bandwidth parameter, which must be strictly greater than γ . The ratio of the time-bandwidth parameter to γ cannot exceed 40. When γ is equal to 3, the Morse wavelet is perfectly symmetric in the frequency domain. The skewness is equal to 0. Values of γ greater than 3 result in positive skewness, while

values of gamma less than 3 result in negative skewness. `WaveletParameters` is only valid if the `Wavelet` property is set to 'Morse'.

The `WaveletParameters` and `TimeBandwidth` properties cannot both be specified.

Example: `'WaveletParameters', [4,20]`

VoicesPerOctave — Number of voices per octave

10 (default) | even integer between 4 and 48

Number of voices per octave to use in determining the necessary separation between the minimum and maximum scales, specified as an even integer between 4 and 48. The minimum and maximum scales are equivalent to the minimum and maximum frequencies or maximum and minimum periods respectively.

Example: `'VoicesPerOctave', 20`

Data Types: double

Output Arguments

minfreq — Minimum wavelet bandpass frequency

scalar

Minimum wavelet bandpass frequency, returned as a scalar. `minfreq` is in cycles/sample if `SamplingFrequency` is not specified. Otherwise, `minfreq` is in hertz.

maxfreq — Maximum wavelet bandpass frequency

scalar

Maximum wavelet bandpass frequency, returned as a scalar. `maxfreq` is in cycles/sample if `SamplingFrequency` is not specified. Otherwise, `maxfreq` is in hertz.

maxperiod — Maximum wavelet bandpass period

scalar duration

Maximum wavelet bandpass period, returned as a scalar duration with the same format as `Ts`.

If the number of standard deviations is set so that $\log_2(\text{maxperiod}/\text{minperiod}) < 1/NV$ where `NV` is the number of voices per octave, `maxperiod` is adjusted to $\text{minperiod} * 2^{(1/NV)}$.

minperiod — Minimum wavelet bandpass period

scalar duration

Minimum wavelet bandpass period, returned as a scalar duration with the same format as `Ts`.

If the number of standard deviations is set so that $\log_2(\text{maxperiod}/\text{minperiod}) < 1/NV$ where NV is the number of voices per octave, `maxperiod` is adjusted to $\text{minperiod} * 2^{(1/NV)}$

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

The `cwtfreqbounds` function supports C/C++ code generation. You must have MATLAB Coder to generate C/C++ code.

Usage notes and limitations:

- The sampling period (`Ts`) input argument is not supported.

See Also

`cwt` | `cwtfilterbank`

Introduced in R2018a

cwt

Continuous 1-D wavelet transform

Note This version of `cwt` is no longer recommended. Use the updated `cwt` instead.

Syntax

```

coefs = cwt(x,scales,'wname')
coefs = cwt(x,scales,'wname','plot')
coefs = cwt(x,scales,'wname','coloration')
coefs = cwt(x,scales,'wname','coloration',xlim)
[coefs,sgram] = cwt(x,scales,'wname','scal')
[coefs,sgram] = cwt(x,scales,'wname','scalCNT')
[coefs,frequencies] = cwt(x,scales,wname,samplingperiod)
[coefs,sgram,frequencies] = cwt(x,scales,wname,samplingperiod,'scal')

```

Description

`coefs = cwt(x,scales,'wname')` returns the continuous wavelet transform (CWT) of the real-valued signal `x`. The wavelet transform is computed for the specified scales using the analyzing wavelet `wname`. `scales` is a 1-D vector with positive elements. The character vector or string scalar `wname` denotes a wavelet recognized by `wavemngr`. `coefs` is a matrix with the number of rows equal to the length of `scales` and number of columns equal to the length of the input signal. The `k`-th row of `coefs` corresponds to the CWT coefficients for the `k`-th element in the scales vector.

`coefs = cwt(x,scales,'wname','plot')` plots the continuous wavelet transform coefficients, using default coloration `'absglb'`.

`coefs = cwt(x,scales,'wname','coloration')` uses the specified coloration. See “Definitions” on page 1-212 for coloration options.

`coefs = cwt(x,scales,'wname','coloration',xlim)` colors the coefficients using coloration and `xlim`, where `xlim` is a vector, `[x1 x2]`, with $1 \leq x1 < x2 \leq \text{length}(x)$.

`[coefs, sgram] = cwt(x, scales, 'wname', 'scal')` returns and plots the scalogram. 'scal' produces an image plot of the scalogram.

`[coefs, sgram] = cwt(x, scales, 'wname', 'scalCNT')` displays a contour representation of the scalogram.

`[coefs, frequencies] = cwt(x, scales, wname, samplingperiod)` returns the frequencies in cycles per unit time corresponding to the scales and the analyzing wavelet *wname*. *samplingperiod* is a positive real-valued scalar. If the units of *samplingperiod* are seconds, the frequencies are in hertz.

`[coefs, sgram, frequencies] = cwt(x, scales, wname, samplingperiod, 'scal')` returns the scalogram and the frequencies corresponding to the scales and the analyzing wavelet. If you have at least two elements in *scales*, you can also use the flag 'scalCNT' to output the scalogram. The *samplingperiod* is only used in the conversion of scales to frequencies. Specifying *samplingperiod* does not affect the appearance of plots generated by `cwt`.

Examples

Plot the continuous wavelet transform and scalogram using `sym2` wavelet at all integer scales from 1 to 32, using a fractal signal as input:

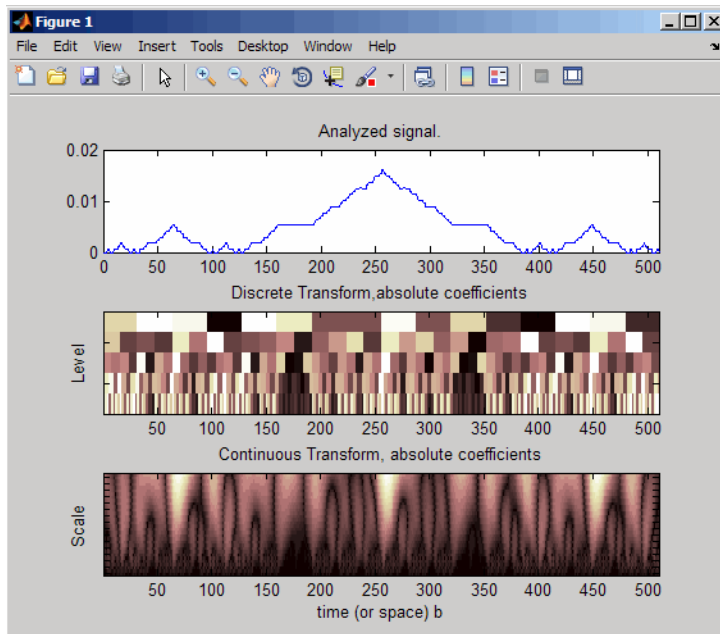
```
load vonkoch
vonkoch=vonkoch(1:510);
len = length(vonkoch);
cw1 = cwt(vonkoch,1:32,'sym2','plot');
title('Continuous Transform, absolute coefficients.')
ylabel('Scale')
[cw1,sc] = cwt(vonkoch,1:32,'sym2','scal');
title('Scalogram')
ylabel('Scale')
```

Compare discrete and continuous wavelet transforms, using a fractal signal as input:

```
load vonkoch
vonkoch=vonkoch(1:510);
len=length(vonkoch);
[c,l]=wavedec(vonkoch,5,'sym2');
% Compute and reshape DWT to compare with CWT.
cfd=zeros(5,len);
for k=1:5
```



```
d=detcoef(c,l,k);
d=d(ones(1,2^k),:);
cfd(k,:)=wkeep(d(:)',len);
end
cfd=cfd(:);
I=find(abs(cfd) <sqrt(eps));
cfd(I)=zeros(size(I));
cfd=reshape(cfd,5,len);
% Plot DWT.
subplot(311); plot(vonkoch); title('Analyzed signal. ');
set(gca,'xlim',[0 510]);
subplot(312);
image(flipud(wcodemat(cfd,255,'row')));
colormap(pink(255));
set(gca,'yticklabel',[]);
title('Discrete Transform, absolute coefficients');
ylabel('Level');
% Compute CWT and compare with DWT
subplot(313);
ccfs=cwt(vonkoch,1:32,'sym2','plot');
title('Continuous Transform, absolute coefficients');
set(gca,'yticklabel',[]);
ylabel('Scale');
```



Definitions

Scale Values

Scale values determine the degree to which the wavelet is compressed or stretched. Low scale values compress the wavelet and correlate better with high frequencies. The low scale CWT coefficients represent the fine-scale features in the input signal vector. High scale values stretch the wavelet and correlate better with the low frequency content of the signal. The high scale CWT coefficients represent the coarse-scale features in the input signal.

Coloration

Coloration is the method used to scale the coefficient values for plotting. Each coefficient is divided by the resulting coloration value.

- 'lvl' — uses maximum value in each scale

- 'glb' — uses maximum value in all scales
- 'abslvl' or 'lvlabs' — uses maximum absolute value in each scale
- 'absglb' or 'glbabs' — uses maximum absolute value in all scales
- 'scal' — produces a scaled image of the scalogram
- 'scalCNT' — produces a contour representation of the scalogram

For 3-D plots (surfaces), use the `coloration` parameter preceded by '3D', such as `coefs = cwt(..., '3Dplot')` or `coefs = cwt(..., '3Dlvl')` ...

Scalogram

Scalograms are plots that represent the percentage energy for each coefficient.

References

Daubechies, I. *Ten Lectures on Wavelets*, Philadelphia, PA: Society for Industrial and Applied Mathematics (SIAM), 1992.

Mallat, S. *A Wavelet Tour of Signal Processing*, San Diego, CA: Academic Press, 1998.

See Also

`cwt` | `dwt` | `wavedec` | `wavefun` | `waveinfo` | `wcodemat`

Topics

"New Wavelet for CWT"

Introduced before R2006a

cwtft

Continuous wavelet transform using FFT algorithm

Note This function is no longer recommended. Use `cwt` instead.

Syntax

```
cwtstruct = cwtft(sig)
cwtstruct = cwtft(sig,Name,Value)
cwtstruct = cwtft(...,'plot')
```

Description

`cwtstruct = cwtft(sig)` returns the continuous wavelet transform (CWT) of the 1-D input signal `sig`. `cwtft` uses an FFT algorithm to compute the CWT. `sig` can be a vector, a structure array, or a cell array. If the sampling interval of your signal is not equal to 1, you must input the sampling period with `sig` in a cell array or a structure array to obtain correct results. If `sig` is a cell array, `sig{1}` is equal to your signal and `sig{2}` is equal to the sampling interval. If `sig` is a structure array, the field `sig.val` contains your signal and `sig.period` contains the sampling interval.

By default, `cwtft` uses the analytic Morlet wavelet. See [More About](#) on page 1-220 for descriptions of valid analyzing wavelets.

For additional default values, see `scales` in “Name-Value Pair Arguments” on page 1-215.

`cwtstruct = cwtft(sig,Name,Value)` returns the continuous wavelet transform (CWT) of the 1-D input signal `sig` with additional options specified by one or more `Name,Value` pair arguments. See “Name-Value Pair Arguments” on page 1-215 for a comprehensive list.

`cwtstruct = cwtft(...,'plot')` plots the continuous wavelet transform. If the analyzing wavelet is real-valued, the original signal along with the CWT coefficient magnitudes and signed CWT coefficients are plotted. If the analyzing wavelet is complex-valued, the original signal is plotted along with the moduli, real parts, imaginary parts,

and angles of the CWT coefficients. You can select the radio button in the bottom left of the plot to superimpose the signal's reconstruction using `icwtft`.

Input Arguments

`sig`

The 1-D input signal. `sig` can be a vector, a structure array, or a cell array. If `sig` is a structure array, `sig` contains two fields: `val` and `period`. `sig.val` is the signal vector and `sig.period` is the sampling period. If `sig` is a cell array, `sig{1}` is the signal vector and `sig{2}` is the sampling period.

If `sig` is a vector, the sampling period defaults to 1.

Note If the sampling interval of your input signal is not 1, you must input the sampling interval with `sig` in a cell array or structure array to obtain correct results. If `sig` is a cell array, `sig{1}` is the 1-D input signal and `sig{2}` is the sampling period. If `sig` is a structure array, the field `sig.val` is the 1-D input signal and `sig.period` is the sampling interval.

Name-Value Pair Arguments

`scales`

Scales over which to compute the CWT. The value of `scales` can be a vector, a structure array, or a cell array. If `scales` is a structure array, it contains at most five fields. The first three fields are mandatory. The last two fields are optional.

- 1 `s0` — The smallest scale. The default `s0` depends on the wavelet. See the More About on page 1-220 for descriptions of the default for each wavelet.
- 2 `nb` — Number of scales. The default `nb` depends on the wavelet. See the More About on page 1-220 for descriptions of the default for each wavelet.
- 3 `ds` — Spacing between scales. The default `ds` depends on the wavelet. See the More About on page 1-220 for descriptions of the default for each wavelet. You can construct a linear or logarithmic scale vector using `ds`. A logarithmic scale vector is constructed by default. Use the `type` field of `scales` to construct a linear scale vector.

- 4** `type` — Type of spacing between scales. `type` can be one of `'pow'` or `'lin'`. The default is `'pow'`. If `type` is equal to `'pow'`, the CWT scales are $s_0 * \text{pow}.^{((\theta : \text{nb} - 1) * ds)}$. This results in a constant spacing of `ds` if you take the logarithm to the base power of the scales vector. If `type` is equal to `'lin'`, the CWT scales are linearly spaced by $s_0 + (\theta : \text{nb} - 1) * ds$.

Use the default power of two spacing to ensure an accurate approximation to the original signal based only on select scales. See the second example in “Examples” on page 1-217 for a signal approximation based on select scales.

- 5** `pow` — The base for `'pow'` spacing. The default is 2. This input is valid only if the `type` argument is `'pow'`.

If `scales` is a cell array, the first three elements of the cell array are identical to the first three elements of the structure array described in the preceding list. The last two elements of the cell array are optional and match the two optional inputs in the structure array described in the preceding list.

wavelet

Analyzing wavelet. To include a parameter for the wavelet, use a cell array. For example, to specify a fourth order derivative of a Gaussian wavelet, use `'wavelet', {'dog', 4}` as in `cwtstruct = (sig, 'wavelet', {'dog', 4})`.

The supported analyzing wavelets are:

- `'dog'` — m -th order derivative of a Gaussian wavelet where m is a positive even integer. The default value of m is 2, which is the Mexican hat or Ricker wavelet..
- `'morl'` — Morlet wavelet. Results in an analytic Morlet wavelet. The Fourier transform of an analytic wavelet is zero for negative frequencies.
- `'morlex'` — non-analytic Morlet wavelet
- `'morl0'` — non-analytic Morlet wavelet with zero mean
- `'mexh'` — Mexican hat wavelet, which is also known as the Ricker wavelet. The Mexican hat wavelet is a special case of the m -th order derivative of a Gaussian wavelet with $m = 2$.
- `'paul'` — Paul wavelet
- `'bump'` — Bump wavelet

See the More About on page 1-220 for formal definitions of the supported analyzing wavelets and associated defaults.

Default: 'morl'

padmode

Signal extension mode. See `dwtmode` for supported extension modes. By default, `cwtft` does not extend the signal prior to computing the CWT. In a Fourier-transform-based CWT algorithm, extending a signal can mitigate wrap-around effects. The number of CWT coefficients in each row of the output matrix `cwtstruct.cfs` is truncated to match the length of the input signal.

Output Arguments

cwtstruct

A structure array with six fields. The fields of the structure array are:

- `cfs` — The CWT coefficient matrix. `cwtstruct.cfs` is an `nb-by-N` matrix where `nb` is the number of scales and `N` is the length of the input signal.
- `scales` — Vector of scales at which the CWT is computed. The length of `cwtstruct.scales` is equal to the row dimension of `cwtstruct.cfs`.
- `frequencies` — Frequencies in cycles per unit time (or space) corresponding to the scales. If the sampling period units are seconds, the frequencies are in hertz. The elements of `frequencies` are in decreasing order to correspond to the elements in the `scales` vector. Use this field to examine the CWT in the time-frequency plane.
- `omega` — Vector of angular frequencies used in the Fourier transform of the wavelet. This field is used in `icwtft` and `icwtlin` for the inversion of the CWT for all wavelets except the bump wavelet.
- `meanSIG` — Mean of the analyzed signal
- `dt` — The sampling interval of the 1-D input signal
- `wav` — Analyzing wavelet

Examples

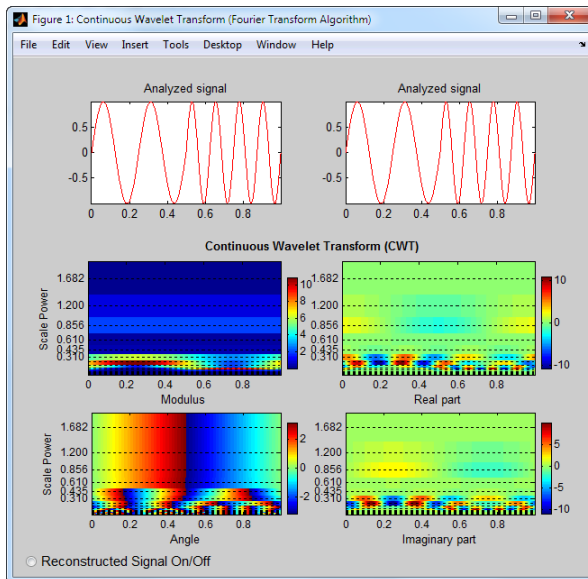
Compute and display the CWT of sine waves with disjoint support. The sampling interval is $1/1023$.

```
N = 1024;
% Sampling interval is 1/1023
```

```

t = linspace(0,1,N);
y = sin(2*pi*4*t).*(t<=0.5)+sin(2*pi*8*t).*(t>0.5);
% Because the sampling interval differs from the default
% you must input it along with the signal
% Using cell array input
sig = {y,1/1023};
cwtS1 = cwtft(sig,'plot');

```



You can display or hide the reconstructed signal using the radio button at the bottom left of the figure. When you select the radio button, the maximum and quadratic relative errors are computed and displayed along with the reconstructed signal.

Reconstruct an approximation to a sum of disjoint sine waves in noise using `cwtft` to decompose the signal and `icwtft` to reconstruct the approximation. Use the CWT coefficients to identify the scales isolating the sinusoidal components. Reconstruct an approximation to the signal based on those scales using the inverse CWT. To ensure an accurate approximation to the based on select scales, use the default power of two spacing in the CWT.

```

rng default % Reset random number generator for reproducible results
N = 1024;
% Sampling interval is 1/1023
t = linspace(0,1,N);

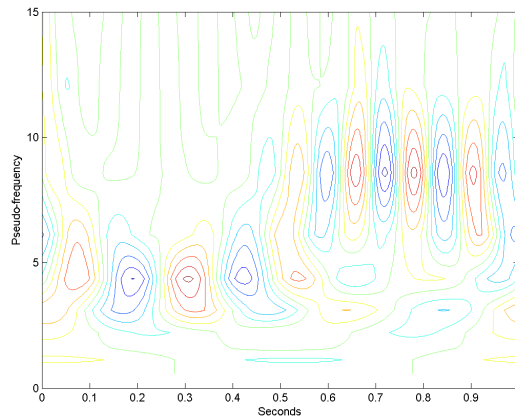
```



```

y = sin(2*pi*4*t).*(t<=0.5)+sin(2*pi*8*t).*(t>0.5);
ynoise = y+randn(size(t));
% Because the sampling interval differs from the default
% you must input it along with the signal
% Using structure array input
sig = struct('val',ynoise,'period',1/1023);
cwtS1 = cwtft(sig);
scales = cwtS1.scales;
MorletFourierFactor = 4*pi/(6+sqrt(2+6^2));
freq = 1./(scales.*MorletFourierFactor);
contour(t,freq,real(cwtS1.cfs));
xlabel('Seconds'); ylabel('Pseudo-frequency');
axis([0 t(end) 0 15]);

```



Extract the scales dominated by energy from the two sine waves and reconstruct a signal approximation using the inverse CWT.

```

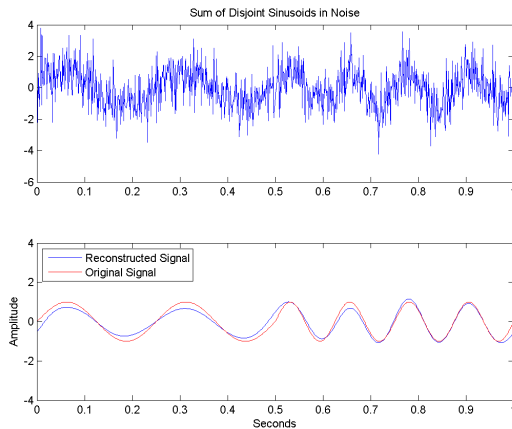
cwtS2 = cwtS1;
cwtS2.cfs = zeros(size(cwtS1.cfs));
cwtS2.cfs(13:15,:) = cwtS1.cfs(13:15,:);
xrec = icwtft(cwtS2);
subplot(2,1,1);
plot(t,ynoise);
title('Sum of Disjoint Sinusoids in Noise');
subplot(2,1,2);
plot(t,xrec,'b'); hold on; axis([0 1 -4 4]);
plot(t,y,'r');

```

```

legend('Reconstructed Signal','Original Signal',...
      'Location','NorthWest');
xlabel('Seconds'); ylabel('Amplitude');

```



Definitions

Morlet Wavelet

Both non-analytic and analytic Morlet wavelets are supported. The analytic Morlet wavelet, 'morl', is defined in the Fourier domain by:

$$\widehat{\psi}(s\omega) = \pi^{-1/4} e^{-(s\omega - \omega_0)^2/2} U(s\omega)$$

where $U(\omega)$ is the Heaviside step function [5] on page 1-223.

The non-analytic Morlet wavelet, 'morlex', is defined in the Fourier domain by:

$$\widehat{\psi}(s\omega) = \pi^{-1/4} e^{-(s\omega - \omega_0)^2/2}$$

'morl0' defines a non-analytic Morlet wavelet in the Fourier domain with exact zero mean:

$$\widehat{\psi}(s\omega) = \pi^{-1/4} \left\{ e^{-(s\omega - \omega_0)^2/2} - e^{-\omega_0^2/2} \right\}$$

The default value of ω_0 is 6.

The default smallest scale for the Morlet wavelets is $s_0 = 2*dt$ where dt is the sampling period.

The default spacing between scales for the Morlet wavelets is $ds=0.4875$.

The default number of scales for the Morlet wavelets is $NbSc = \text{fix}(\log_2(\text{length}(\text{sig})*dt/s_0)/ds)$.

The default scales for the Morlet wavelet are $s_0*2.^{((0:NbSc-1)*ds)}$.

***m*-th Order Derivative of Gaussian Wavelets**

In the Fourier domain, the m -th order derivative of Gaussian wavelets, 'dog', are defined by:

$$\widehat{\Psi}(s\omega) = -\frac{1}{\sqrt{\Gamma(m+1/2)}}(js\omega)^m e^{-(s\omega)^2/2}$$

where $\Gamma(\cdot)$ denotes the gamma function [5] on page 1-223.

The derivative must be an even order. The default order of the derivative is 2, which is also known as the Mexican hat wavelet.

The default smallest scale for the DOG wavelet is $s_0 = 2*dt$ where dt is the sampling period.

The default spacing between scales for the DOG wavelet is $ds=0.4875$.

The default number of scales for the DOG wavelet is $NbSc = \text{fix}(\log_2(\text{length}(\text{sig})*dt/s_0)/ds)$.

The default scales for the DOG wavelet are $s_0*2.^{((0:NbSc-1)*ds)}$.

Paul Wavelet

The Fourier transform of the analytic Paul wavelet, 'paul', of order m is:

$$\widehat{\Psi}(s\omega) = \frac{2^m}{\sqrt{m(2m-1)!}}(s\omega)^m e^{-s\omega} U(s\omega)$$

where $U(\omega)$ is the Heaviside step function [5] on page 1-223.

The default order of the Paul wavelet is 4.

The default smallest scale for the Paul wavelet is $s_0 = 2*dt$ where dt is the sampling period.

The default spacing between scales for the Paul wavelet is $ds=0.4875$.

The default number of scales for the Paul wavelet is $NbSc = \text{fix}(\log_2(\text{length}(\text{sig}) * dt / s_0) / ds)$.

The default scales for the Paul wavelet are $s_0 * 2.^{(0:NbSc-1) * ds}$.

Bump wavelet

The Fourier transform of the analytic bump wavelet, 'bump', with parameters μ and σ is

$$\widehat{\psi}(s\omega) = e^{\left(1 - \frac{1}{1 - (s\omega - \mu)^2 / \sigma^2}\right)} 1_{[(\mu - \sigma)/s, (\mu + \sigma)/s]}$$

where $1_{[(\mu - \sigma)/s, (\mu + \sigma)/s]}$ is the indicator function for the interval $(\mu - \sigma)/s \leq \omega \leq (\mu + \sigma)/s$.

Valid values for μ are [3,6]. Valid values for σ are [0.1, 1.2]. Smaller values of σ result in a wavelet with superior frequency localization but poorer time localization. Larger values of σ produce a wavelet with better time localization and poorer frequency localization.

The default values for μ and σ are 5 and 0.6 respectively.

The default smallest scale for the bump wavelet is $s_0 = 2*dt$ where dt is the sampling period.

The default spacing between scales for the bump wavelet is $ds=1/10$.

The default number of scales for the bump wavelet is $NbSc = \text{fix}(\log_2(\text{length}(\text{sig}) * dt / s_0) / ds)$.

The default scales for the bump wavelet are $s_0 * 2.^{(0:NbSc-1) * ds}$.

Algorithms

`cwtft` implements the following algorithm:

- Obtain the discrete Fourier transform (DFT) of the signal using `fft`.
- Obtain the DFT of the analyzing wavelet at the appropriate angular frequencies. Scale the DFT of the analyzing wavelet at different scales to ensure different scales are directly comparable.
- Take the product of the signal DFT and the wavelet DFT over all the scales. Invert the DFT to obtain the CWT coefficients.

For a mathematical motivation for the FFT-based algorithm see “Continuous Wavelet Transform and Scale-Based Analysis”.

Alternatives

- `cwt` — Computes the CWT using convolutions. `cwt` supports a wider choice of analyzing wavelets than `cwtft`, but may be more computationally expensive. The output of `cwt` is not compatible with the inverse CWT implemented with `icwtft`. To use `icwtft`, obtain the CWT with `cwtft`.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, Philadelphia, PA: Society for Industrial and Applied Mathematics (SIAM), 1992.
- [2] Farge, M. “Wavelet Transforms and Their Application to Turbulence”, *Ann. Rev. Fluid. Mech.*, 1992, 24, 395-457.
- [3] Mallat, S. *A Wavelet Tour of Signal Processing*, San Diego, CA: Academic Press, 1998.
- [4] Sun, W. “Convergence of Morlet's Reconstruction Formula”, *preprint*, 2010.
- [5] Torrence, C. and G.P. Compo. “A Practical Guide to Wavelet Analysis”, *Bull. Am. Meteorol. Soc.*, 79, 61-78, 1998.

See Also

cwt | icwtft | icwtlin

Topics

“Continuous and Discrete Wavelet Transforms”

“Continuous Wavelet Transform and Scale-Based Analysis”

“Inverse Continuous Wavelet Transform”

“Time-Frequency Analysis with the Continuous Wavelet Transform”

Introduced in R2011a

cwtftinfo

Valid analyzing wavelets for FFT-based CWT

Note This function is no longer recommended. Use `cwt` instead.

Syntax

`cwtftinfo`

Description

`cwtftinfo` displays expressions for the Fourier transforms of valid analyzing wavelets for use with `cwtfft`.

Examples

Display a list of Fourier transforms for all valid analyzing wavelets.

`cwtftinfo`

Definitions

Morlet Wavelet

Both non-analytic and analytic Morlet wavelets are supported. The analytic Morlet wavelet, 'morl', is defined in the Fourier domain by:

$$\widehat{\Psi}(s\omega) = \pi^{-1/4} e^{-(s\omega - \omega_0)^2/2} U(s\omega)$$

where $U(\omega)$ is the Heaviside step function.

The non-analytic Morlet wavelet, 'morlex', is defined in the Fourier domain by:

$$\widehat{\psi}(s\omega) = \pi^{-1/4} e^{-(s\omega - \omega_0)^2/2}$$

'morl0' defines a non-analytic Morlet wavelet in the Fourier domain with exact zero mean:

$$\widehat{\psi}(s\omega) = \pi^{-1/4} \left\{ e^{-(s\omega - \omega_0)^2/2} - e^{-\omega_0^2/2} \right\}$$

The default value of ω_0 is 6.

***m*-th Order Derivative of Gaussian Wavelets**

In the Fourier domain, the *m*-th order derivative of Gaussian wavelets, 'dog', is defined by:

$$\widehat{\psi}(s\omega) = -\frac{1}{\sqrt{\Gamma(m + 1/2)}} (js\omega)^m e^{-(s\omega)^2/2}$$

The derivative must be an even order. The default order of the derivative is 2, which is also known as the *Mexican hat* or *Ricker* wavelet.

Because the unit imaginary, *j*, is always raised to an even power, the Fourier transform is real-valued.

Paul Wavelet

The Fourier transform of the Paul wavelet, 'paul', of order *m* is:

$$\widehat{\psi}(s\omega) = \frac{2^m}{\sqrt{m(2m - 1)!}} (s\omega)^m e^{-s\omega} U(s\omega)$$

where *U*(ω) is the Heaviside step function. The Paul wavelet is analytic.

The default order of the Paul wavelet is 4.

Bump wavelet

The Fourier transform of the analytic bump wavelet, 'bump', with parameters μ and σ is

$$\widehat{\psi}(s\omega) = e^{\left(1 - \frac{1}{1 - (s\omega - \mu)^2/\sigma^2}\right)} 1_{[(\mu - \sigma)/s, (\mu + \sigma)/s]}$$

where $1_{[(\mu - \sigma)/s, (\mu + \sigma)/s]}$ is the indicator function for the interval $(\mu - \sigma)/s \leq \omega \leq (\mu + \sigma)/s$.

Valid values for μ are [3,6]. Valid values for σ are [0.1, 1.2]. Smaller values of σ result in a wavelet with superior frequency localization but poorer time localization. Larger values of σ produce a wavelet with better time localization and poorer frequency localization.

The default values for μ and σ are 5 and 0.6 respectively.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, Philadelphia, PA: Society for Industrial and Applied Mathematics (SIAM), 1992.
- [2] Farge, M. *Wavelet Transforms and Their Application to Turbulence*, Ann. Rev. Fluid. Mech., 1992, 24, 395-457.
- [3] Mallat, S. *A Wavelet Tour of Signal Processing*, San Diego, CA: Academic Press, 1998.
- [4] Torrence, C. and G.P. Compo *A Practical Guide to Wavelet Analysis*, Bull. Am. Meteorol. Soc., 79, 61-78, 1998.

See Also

cwtft | icwtft | icwtlin

Topics

“Continuous and Discrete Wavelet Transforms”
“Continuous Wavelet Transform and Scale-Based Analysis”
“Inverse Continuous Wavelet Transform”

Introduced in R2011a

cwtftinfo2

Supported 2-D CWT wavelets and Fourier transforms

Syntax

```
cwtftinfo2  
cwtftinfo2(wname)
```

Description

`cwtftinfo2` lists the supported 2-D continuous wavelet transform (CWT) wavelets and corresponding parameters for use with `cwtft2`.

`cwtftinfo2(wname)` displays the equation for the 2-D Fourier transform of the wavelet, `wname`. The figure with the 2-D Fourier transform of the analyzing wavelet has a drop-down list from which you can select other wavelets.

Examples

Available Wavelets with Parameters

```
cwtftinfo2
```

```
CWTFTINF02 Information on wavelets for CWTFT2  
CWTFTINF02 provides information on the available wavelets  
for 2-D Continuous Wavelet Transform using FFT.  
The wavelets are defined by their Fourier transform.
```

```
The formulae giving the Fourier transform of  
the wavelet which short name (see below) is SNAME  
will be displayed using CWTFTINF02(SNAME).
```

```
The table below gives the short name of each wavelet  
and the associated parameters: first, the name of parameter  
and then the default value.
```

```

WAV_Param_Table = {...
    'morl'
        ,
        defaults: omega0 = 6; sigma = 1; epsilon = 1;
    'mexh'
        ,
        defaults: p = 2; sigmax = 1; sigmay = 1;
    'paul'
        ,
        defaults: p = 4;
    'dog'
        ,
        defaults: alpha = 2;
    'cauchy'
        ,
        defaults: alpha = pi/6; sigma = 1; L = 4; M = 4;
    'escauchy'
        ,
        defaults: alpha = pi/6; sigma = 1; L = 4; M = 4;
    'gaus'
        ,
        defaults: p = 1; sigmax = 1; sigmay = 1;
    'wheel'
        ,
        defaults: sigma = 2;
    'fan'
        ,
        defaults: omega0 = 5.336; sigma = 1; epsilon = 1; J = 6.5;
    'pethat'
        ,
        defaults: No parameters.
    'dogpow'
        ,
        defaults: alpha = 1.25; p = 2;
    'esmorl'
        ,
        defaults: omega0 = 6; sigma = 1; epsilon = 1;
    'esmexh'
        ,
        defaults: sigma = 1; epsilon = 0.5;
    'gaus2'
        ,
        defaults: p = 1; sigmax = 1; sigmay = 1;
    'gaus3'
        ,
        defaults: A = 1; B = 1; p = 1; sigmax = 1; sigmay = 1;
    'isodog'
        ,
        defaults: alpha = 1.25;
    'dog2'
        ,
        defaults = alpha = 1.25;
    'isomorl'
        ,
        defaults: omega0 = 6; sigma = 1;
    'rmorl'
        ,
        defaults: omega0 = 6; sigma = 1; epsilon = 1;
    'endstop1'
        ,
        defaults: omega0 = 6;
    'endstop2'
        ,
        defaults: omega0 = 6; sigma = 1;

```

```
'gabmexh' ,
      defaults: omega0 = 5.336; epsilon = 1;
'sinc' ,
      defaults: Ax = 1; Ay = 1; p = 1; omega0X= 0; Omega0Y = 0;
};
```

The various wavelets may be grouped in families as follow:

```
MORLET: 'morl' , 'esmorl' , 'rmorl' , 'isomorl'
DOG:    'dog' , 'isodog' , 'dog2' , 'dogpow'
GAUSS:  'mexh' , 'gaus' , 'gaus2' , 'gaus3' , 'esmexh'
PAUL:   'paul'
CAUCHY: 'cauchy' , 'escauchy'
WHEEL:  'wheel' , 'pethat'
MISCELLANEOUS : 'endstop1' , 'endstop2' , 'gabmexh' , 'sinc' , 'fan'
```

REFERENCES

Two-Dimensional Wavelets and their Relatives
J.-P. Antoine, R. Murenzi, P. Vandergheynst and S. Twareque Ali
Cambridge University Press - 2004

Two-dimensional wavelet transform profilometry
Fringe Pattern Analysis Using Wavelet
Liverpool John Moores University
<http://www.ljmu.ac.uk>

See also CWTFT2

Display the Expression for the 2-D Fourier Transform

Display the expression for the 2-D Fourier transform of the Cauchy wavelet. After displaying the Fourier transform for any wavelet, you can use the drop-down list in the bottom left to view the Fourier transform for any supported wavelet.

```
cwtftinfo2('cauchy')
```

cauchy

$$\widehat{\psi}(\omega_x, \omega_y) = \left[\sin(\alpha)\omega_x + \cos(\alpha)\omega_y \right]^L \dots$$

$$\left[-\sin(\alpha)\omega_x + \cos(\alpha)\omega_y \right]^M \left| \tan(\alpha)\omega_x > |\omega_y| \right| e^{-\sigma \frac{(\omega_x)^2 + (\omega_y)^2}{2}}$$

$$\sigma \in]0, +\infty[; \alpha \in \left] 0, \frac{\pi}{2} \right[; L, M \in]0, +\infty[$$

Input Arguments

wname — Wavelet name

character vector | string scalar

Wavelet name, specified as a character vector or string scalar. The following table lists the supported wavelets for the 2-D CWT and associated parameters:

Wavelet name	Parameters
'morl'	{'Omega0',6;'Sigma',1;'Epsilon',1}
'mexh'	{'p',2;'sigmax',1;'sigmay',1}
'paul'	{'p',4}
'dog'	{'alpha',1.25}
'cauchy'	{'alpha','pi/6';'sigma',1;'L',4;'M',4}
'escauchy'	{'alpha','pi/6';'sigma',1;'L',4;'M',4}
'gaus'	{'p',1;'sigmax',1;'sigmay',1}
'wheel'	{'sigma',2}
'fan'	{'Omega0X',5.336;'Sigma',1;'Epsilon',1;'J',6.5}
'pethat'	None
'dogpow'	{'alpha',1.25;'p',2}
'esmorl'	{'Omega0',6;'Sigma',1;'Epsilon',1}
'esmexh'	{'Sigma',1;'Epsilon',0.5}
'gaus2'	{'p',1;'sigmax',1;'sigmay',1}
'gaus3'	{'A',1;'B',1;'p',1;'sigmax',1;'sigmay',1}
'isodog'	{'alpha',1.25}
'dog2'	{'alpha',1.25}
'isomorl'	{'Omega0',6;'Sigma',1}
'rmorl'	{'Omega0',6;'Sigma',1;'Epsilon',1}
'endstop1'	{'Omega0',6}
'endstop2'	{'Omega0',6;'Sigma',1}
'gabmexh'	{'Omega0',5.336;'Epsilon',1}

Wavelet name	Parameters
'sinc'	{'Ax',1;'Ay',1;'p',1;'Omega0X',0;'Omega0Y',0}

Example: `cwtftinfo2('paul')`

Data Types: char

Introduced in R2013b

cwtft2

2-D continuous wavelet transform

Syntax

```
cwtstruct = cwtft2(x)
cwtstruct = cwtft2(x, 'plot')
cwtstruct = cwtft2(x, Name, Value)
```

Description

`cwtstruct = cwtft2(x)` returns the 2-D continuous wavelet transform (CWT) of the 2-D matrix, `x`. `cwtft2` uses a Fourier transform-based algorithm in which the 2-D Fourier transforms of the input data and analyzing wavelet are multiplied together and inverted.

`cwtstruct = cwtft2(x, 'plot')` plots the data and the 2-D CWT.

`cwtstruct = cwtft2(x, Name, Value)` uses additional options specified by one or more `Name, Value` pair arguments.

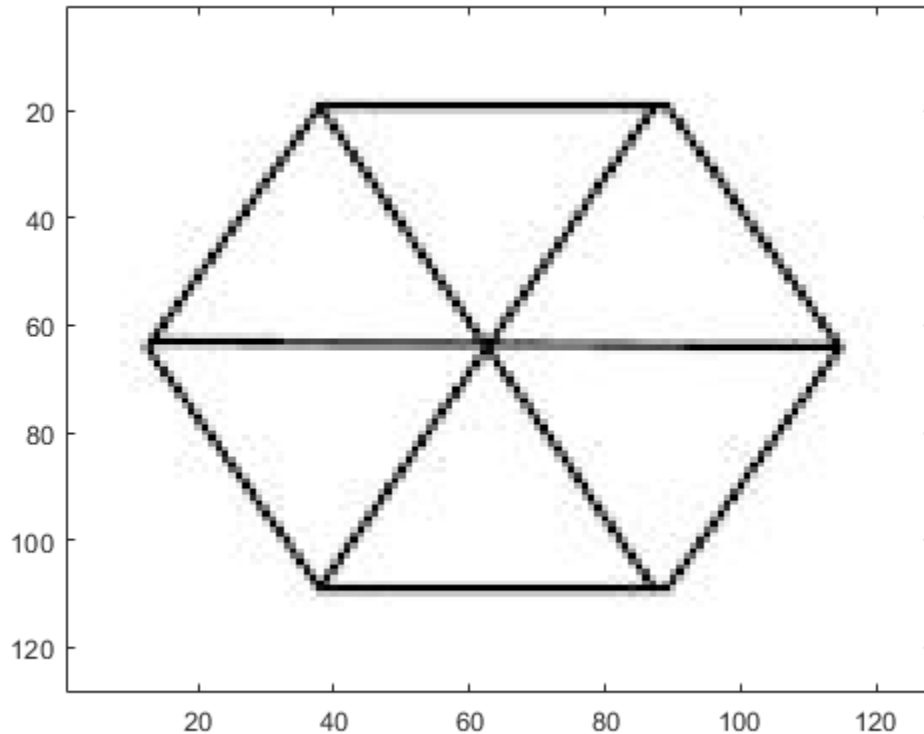
Examples

Compare Isotropic and Anisotropic Wavelets

Shows how an isotropic wavelet does not discern the orientation of features while an anisotropic wavelet does. The example uses the Mexican hat isotropic wavelet and the directional (anisotropic) Cauchy wavelet.

Load and view the hexagon image.

```
Im = imread('hexagon.jpg');
imagesc(Im); colormap(jet);
```

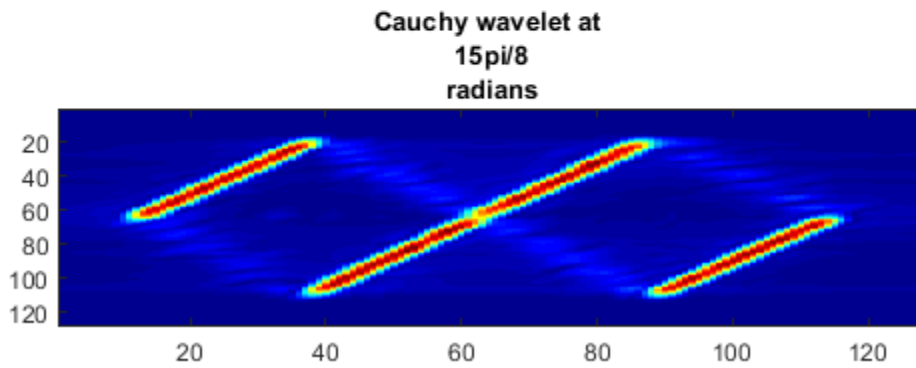
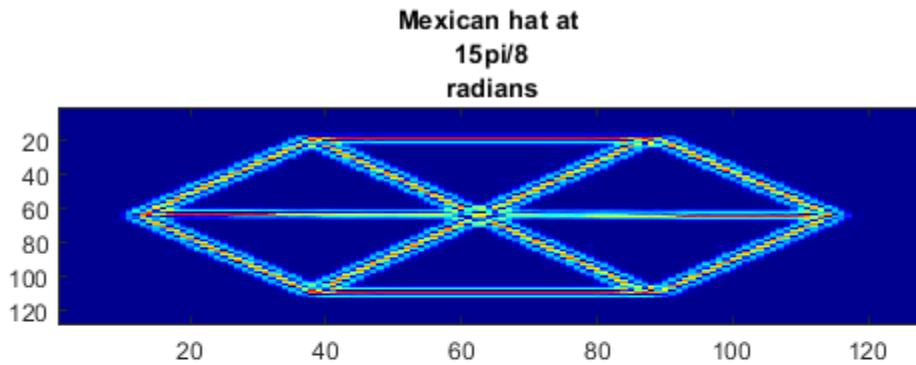
Obtain the scale-one 2-D CWT with both the Mexican hat and Cauchy wavelets. Specify a vector of angles going from 0 to $15\pi/8$ in $\pi/8$ increments.

```
cwtcauchy = cwtft2(Im,'wavelet','cauchy','scales',1,...
    'angles',0:pi/8:2*pi-pi/8);
cwtmexh = cwtft2(Im,'wavelet','mexh','scales',1,...
    'angles',0:pi/8:2*pi-pi/8);
```

Visualize the scale-one 2-D CWT coefficient magnitudes at each angle.

```
angz = {'0', 'pi/8', 'pi/4', '3pi/8', 'pi/2', '5pi/8', '3pi/4', ...
    '7pi/8', 'pi', '9pi/8', '5pi/4', '11pi/8', '3pi/2', ...
    '13pi/8', '7pi/4', '15pi/8'};
for angn = 1:length(angz)
```

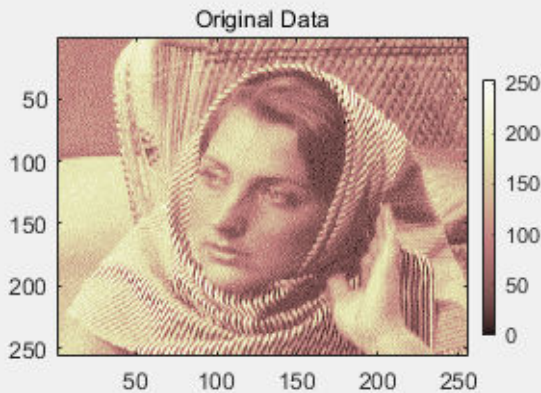
```
subplot(211)
imagesc(abs(cwtmexh.cfs(:,:,1,1,angn)));
title(['Mexican hat at ' angz(angn) 'radians']);
subplot(212)
imagesc(abs(cwtcauchy.cfs(:,:,1,1,angn)));
title(['Cauchy wavelet at ' angz(angn) 'radians']);
pause(1);
end
```



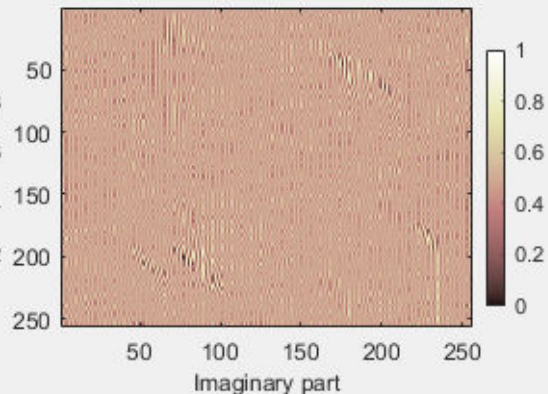
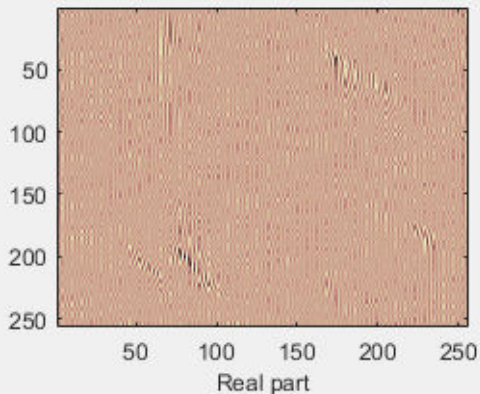
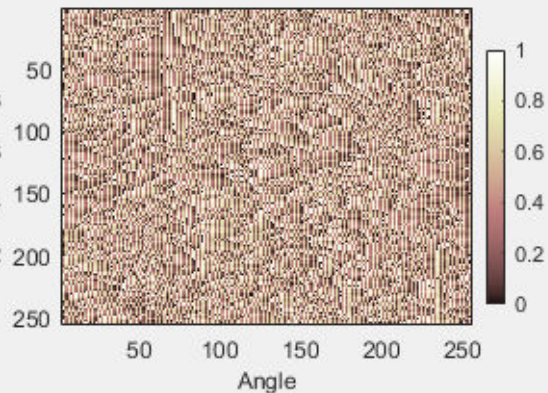
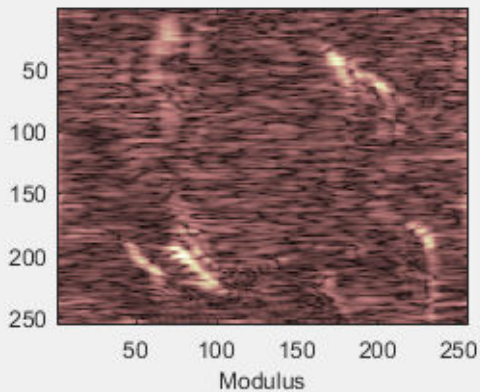
Plot 2-D CWT

Load an image of a woman, obtain the 2-D CWT using the Morlet wavelet, and plot the CWT coefficients.

```
load woman;  
cwtmorl = cwtft2(X, 'scales', 1:4, 'angles', 0:pi/2:3*pi/2, 'plot');
```



Scale: Index 1 - Value 1.000 -- Angle: Index 1 - Value 0 pi [rad] = 0.00 [dgr]



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Idx. Scale1 -- Value: 1.000

Idx. Angle1 -- Value: 0 pi

pink

2-D CWT with Morlet Wavelet

Obtain the 2-D CWT of the star image using the default Morlet wavelet, scales $2^{(0:5)}$, and an angle of 0.

```
Im = imread('star.jpg');  
cwtout = cwtft2(Im);
```

Input Arguments

x — Input data

array

Input data, specified as a 2-D matrix or 3-D array. If the input data is a 3-D array, the input matrix is a truecolor image.

```
Example: X = imread('stars.jpg');
```

Data Types: double | uint8

Name-Value Pair Arguments

Specify optional comma-separated pairs of **Name**, **Value** arguments. **Name** is the argument name and **Value** is the corresponding value. **Name** must appear inside quotes. You can specify several name and value pair arguments in any order as **Name1**, **Value1**, ..., **NameN**, **ValueN**.

Example: 'wavelet', 'paul', 'scales', $2^{(0:5)}$ specifies to use the Paul wavelet and a vector of scales.

angles — Angles

0 (default) | scalar | vector

Angles in radians, specified as a comma-separated pair consisting of 'angles' and either a scalar or a vector.

```
Example: 'angles', [0 pi/2 pi]
```

norm — Normalization`'L2'` (default) | `'L1'` | `'L0'`

Normalization used in the 2-D CWT, specified as a comma-separated pair consisting of `'norm'` and one of these character vectors:

- `'L2'` — The Fourier transform of the analyzing wavelet at a given scale is multiplied by the corresponding scale. `'L2'` is the default normalization.
- `'L1'` — The Fourier transform of the analyzing wavelet is multiplied by 1 at all scales.
- `'L0'` — The Fourier transform of the analyzing wavelet at a given scale is multiplied by the square of the corresponding scale.

Example: `'norm','L1'`

scales — Scales`2^(0:5)` (default) | scalar | vector

Scales, specified as a comma-separated pair consisting of `'scales'` and either a positive real-valued scalar or a vector of positive real numbers.

Example: `'scales',2^(1:6)`

wavelet — Analyzing wavelet`'morl'` (default) | character vector | string scalar | structure | cell array

Analyzing wavelet, specified as a comma-separated pair consisting of `'wavelet'` and a character vector, a string scalar, a structure, or a cell array. `cwtftinfo2` provides a comprehensive list of supported wavelets and associated parameters.

If you specify `'wavelet'` as a structure, the structure must contain two fields:

- `name` — the character vector or string scalar corresponding to a supported wavelet.
- `param` — a cell array with the parameters of the wavelet.

If you specify `'wavelet'` as a cell array, `wav`, the cell array must contain two elements:

- `wav{1}` — the character vector or string scalar corresponding to a supported wavelet.
- `wav{2}` — a cell array with the parameters of the wavelet.

Example: `'wavelet',{'morl',{6,1,1}}`

Example: `'wavelet',struct('name','paul','param',{'p',2})`

Output Arguments

cwtstruct — 2-D CWT

structure

The 2-D CWT, returned as a structure with the following fields:

wav — Analyzing wavelet and parameters

structure

Analyzing wavelet and parameters, returned as a structure with the following fields:

- **wname** — name
- **param** — parameters

wav_norm — Normalization constants

matrix

Normalization constants, returned as a M -by- N matrix where M is the number of scales and N is the number of angles.

cfs — CWT coefficients

array

CWT coefficients, returned as an N-D array. The row and column dimensions of the array equal the row and column dimensions of the input data. The third page of the array is equal to 1 or 3 depending on whether the input data is a grayscale or truecolor image. The fourth page of the array is equal to the number of scales and the fifth page of the array is equal to the number of angles.

scales — Scales

vector

Scales for the 2-D CWT, returned as a row vector.

angles — Angles

vector

Angles for the 2-D CWT, returned as a row vector.

meanSIG — Mean

scalar

Mean of the input data, returned as a scalar

See Also

`cwtftinfo2`

Topics

“Two-Dimensional CWT of Noisy Pattern”

“2-D Continuous Wavelet Transform App”

Introduced in R2013b

dbaux

Daubechies wavelet filter computation

The dbaux function generates the scaling filter coefficients for the "extremal phase" Daubechies wavelets.

Syntax

$W = \text{dbaux}(N)$
 $W = \text{dbaux}(N, \text{SUMW})$

Description

$W = \text{dbaux}(N)$ is the order N Daubechies scaling filter such that $\text{sum}(W) = 1$.

Note

- Instability may occur when N is too large. Starting with values of N in the 30s range, function output will no longer accurately represent scaling filter coefficients.
-

$W = \text{dbaux}(N, \text{SUMW})$ is the order N Daubechies scaling filter such that $\text{sum}(W) = \text{SUMW}$.

$W = \text{dbaux}(N, 0)$ is equivalent to $W = \text{dbaux}(N, 1)$.

Examples

Daubechies Extremal Phase Scaling Filter with Specified Sum

This example shows how to determine the Daubechies extremal phase scaling filter with a specified sum. The two most common values for the sum are $\sqrt{2}$ and 1.

Construct two versions of the db4 scaling filter. One scaling filter sums to $\sqrt{2}$ and the other version sums to 1.

```
NumVanishingMoments = 4;  
h = dbaux(NumVanishingMoments,sqrt(2));  
m0 = dbaux(NumVanishingMoments,1);
```

The filter with sum equal to $\sqrt{2}$ is the synthesis (reconstruction) filter returned by `wfilters` and used in the discrete wavelet transform.

```
[LoD,HiD,LoR,HiR] = wfilters('db4');  
max(abs(LoR-h))
```

```
ans = 4.2590e-13
```

For orthogonal wavelets, the analysis (decomposition) filter is the time-reverse of the synthesis filter.

```
max(abs(LoD-fliplr(h)))
```

```
ans = 4.2590e-13
```

Extremal Phase

This example demonstrates that for a given support, the cumulative sum of the squared coefficients of a scaling filter increase more rapidly for an extremal phase wavelet than other wavelets.

First, set the order to 15 and generate the scaling filter coefficients for the Daubechies wavelet and Symlet. Both wavelets have support of length 29.

```
n = 15;  
[~,~,LoR_db,~] = wfilters('db15');  
[~,~,LoR_sym,~] = wfilters('sym15');
```

Next, generate the scaling filter coefficients for the order 5 Coiflet. This wavelet also has support of length 29.

```
[~,~,LoR_coif,~] = wfilters('coif5');
```

Confirm the sum of the coefficients for all three wavelets equals $\sqrt{2}$.

```
sqrt(2)-sum(LoR_db)
```

```
ans = 2.2204e-16
```

```
sqrt(2)-sum(LoR_sym)
```

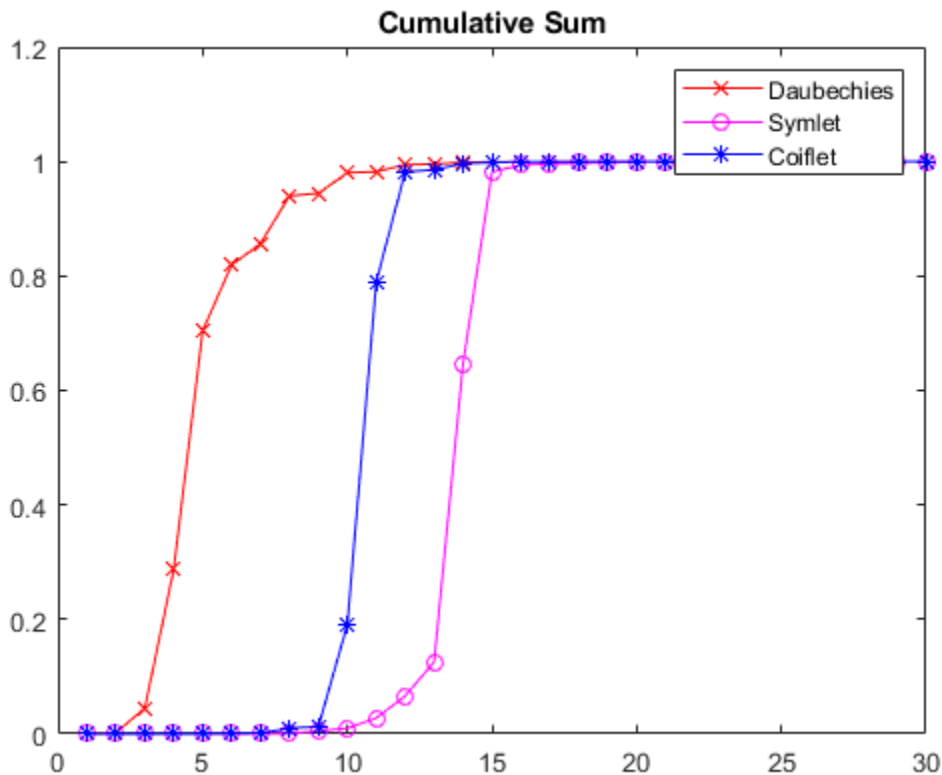
```
ans = 0
```

```
sqrt(2)-sum(LoR_coif)
```

```
ans = 2.2204e-16
```

Plot the cumulative sums of the squared coefficients. Note how rapidly the Daubechies sum increases. This is because its energy is concentrated at small abscissas. Since the Daubechies wavelet has extremal phase, the cumulative sum of its squared coefficients increases more rapidly than the other two wavelets.

```
plot(cumsum(LoR_db.^2), 'rx-')
hold on
plot(cumsum(LoR_sym.^2), 'mo-')
plot(cumsum(LoR_coif.^2), 'b*-')
legend('Daubechies', 'Symlet', 'Coiflet')
title('Cumulative Sum')
```



Input Arguments

N — Order of Daubechies scaling filter

positive integer

Order of Daubechies scaling filter, specified as a positive integer.

Data Types: `single` | `double`

SUMW — Sum of coefficients

1 (default) | positive scalar

Sum of coefficients, specified as a positive scalar. Set to `sqrt(2)` to generate vector of coefficients whose norm is 1.

Data Types: `single` | `double`

Output Arguments

W — Scaling filter coefficients

vector

Scaling filter coefficients returned as a vector.

The scaling filter coefficients satisfy a number of properties. As the example “Daubechies Extremal Phase Scaling Filter with Specified Sum” on page 1-243 demonstrates, you can construct scaling filter coefficients with a specific sum. If $\{h_k\}$ denotes the set of order N

Daubechies scaling filter coefficients, where $n = 1, \dots, 2N$, then $\sum_{n=1}^{2N} h_n^2 = 1$. The

coefficients also satisfy the relation $\sum_n h(n)h(n-2k) = \delta(k)$. You can use these properties to check your results.

Limitations

- The computation of the dbN Daubechies scaling filter requires the extraction of the roots of a polynomial of order 4N. Instability may occur beginning with values of N in the 30s.

Definitions

Extremal Phase

Constructing a compactly supported orthogonal wavelet basis involves choosing roots of a particular polynomial equation. Different choices of roots will result in wavelets whose phases are different. Choosing roots that lie within the unit circle in the complex plane results in a filter with highly nonlinear phase. Such a wavelet is said to have extremal phase, and has energy concentrated at small abscissas. Let $\{h_k\}$ denote the set of scaling coefficients associated with an extremal phase wavelet, where $k = 1, \dots, N$. Then for any

other set of scaling coefficients $\{g_k\}$ resulting from a different choice of roots, the following inequality will hold for all $J = 1, \dots, N$:

$$\sum_{k=1}^J g_k^2 \leq \sum_{k=1}^J h_k^2$$

The inequality is illustrated in the example “Extremal Phase” on page 1-244. The $\{h_k\}$ are sometimes called a *minimal delay filter* [2].

Algorithms

The algorithm used is based on a result obtained by Shensa [3], showing a correspondence between the “Lagrange à trous” filters and the convolutional squares of the Daubechies wavelet filters.

The computation of the order N Daubechies scaling filter w proceeds in two steps: compute a “Lagrange à trous” filter P , and extract a square root. More precisely:

- P the associated “Lagrange à trous” filter is a symmetric filter of length $4N-1$. P is defined by

$$P = [a(N) \ 0 \ a(N-1) \ 0 \ \dots \ 0 \ a(1) \ 1 \ a(1) \ 0 \ a(2) \ 0 \ \dots \ 0 \ a(N)]$$

- where

$$\alpha(k) = \frac{\prod_{\substack{i=-N+1 \\ i \neq k}}^N \left(\frac{1}{2} - i\right)}{\prod_{\substack{i=-N+1 \\ i \neq k}}^N (k-i)} \quad \text{for } k = 1, \dots, N$$

- Then, if w denotes db N Daubechies scaling filter of sum $\sqrt{2}$, w is a square root of P :

$$P = \text{conv}(w \text{rev}(w), w) \quad \text{where } w \text{ is a filter of length } 2N.$$

The corresponding polynomial has N zeros located at -1 and $N-1$ zeros less than 1 in modulus.

Note that other methods can be used; see various solutions of the spectral factorization problem in Strang-Nguyen [4] (p. 157).

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, CBMS-NSF conference series in applied mathematics, SIAM Ed., 1992.
- [2] Oppenheim, Alan V., and Ronald W. Schafer. *Discrete-Time Signal Processing*. Englewood Cliffs, NJ: Prentice Hall, 1989.
- [3] Shensa, M.J. (1992), "The discrete wavelet transform: wedding the a trous and Mallat Algorithms," *IEEE Trans. on Signal Processing*, vol. 40, 10, pp. 2464-2482.
- [4] Strang, G., and T. Nguyen. *Wavelets and Filter Banks*. Wellesley, MA: Wellesley-Cambridge Press, 1996.

See Also

dbwavf | symaux | wfilters

Introduced before R2006a

dbwavf

Daubechies wavelet filter

Syntax

```
f = dbwavf(wname)
```

Description

`f = dbwavf(wname)` returns the scaling filter associated with the Daubechies wavelet specified by `wname`. `f` is a real-valued vector.

Examples

Scaling Filter Associated With Daubechies Wavelet

Specify the order 4 Daubechies wavelet.

```
wname = 'db4';
```

Compute the corresponding scaling filter.

```
f = dbwavf(wname);  
f'
```

```
ans = 8×1
```

```
0.1629  
0.5055  
0.4461  
-0.0198  
-0.1323  
0.0218  
0.0233  
-0.0075
```


Input Arguments

wname — Daubechies wavelet

'dbN'

Daubechies wavelet with N vanishing moments, where N is a positive integer in the closed interval $[1, 45]$.

See Also

dbaux | waveinfo | wfilters

Introduced before R2006a

ddencmp

Default values for denoising or compression

ddencmp returns default values for denoising or compression for the critically sampled discrete wavelet or wavelet packet transform.

Syntax

```
[thr,sorh,keepapp] = ddencmp(in1,in2,x)
[ ___,crit] = ddencmp(in1,'wp',x)
```

Description

[thr,sorh,keepapp] = ddencmp(in1,in2,x) returns default values for denoising or compression, using wavelets or wavelet packets, of the input data x . x is a real-valued vector or 2-D matrix. thr is the threshold, and sorh indicates soft or hard thresholding. keepapp can be used as a flag to set whether or not the approximation coefficients are thresholded.

- Set in1 to 'den' for denoising or 'cmp' for compression.
- Set in2 to 'wv' to use wavelets or 'wp' to use wavelet packets.

[___,crit] = ddencmp(in1,'wp',x) also returns the entropy type, crit. See wentropy for more information.

Examples

Default Global Threshold for Wavelet Denoising

Determine the default global denoising threshold for an $N(0,1)$ white noise input. Create an $N(0,1)$ white noise input. Change the DWT extension mode to periodic. Set the random number generator to the default initial settings for reproducible results.

```
dwtmode('per','nodisp');
rng default;
x = randn(512,1);
```

Use `ddencmp` to obtain the default global threshold for wavelet denoising. Demonstrate that the threshold is equal to the universal threshold of Donoho and Johnstone scaled by a robust estimate of the variance.

```
[thr,sorh,keepapp] = ddencmp('den','wv',x);
[A,D] = dwt(x,'db1');
noiselev = median(abs(D))/0.6745;
thresh = sqrt(2*log(length(x)))*noiselev;
```

Compare the value of the variable `thr` to the value of `thresh`.

```
thr
thr = 3.3639

thresh
thresh = 3.3639
```

Default Global Threshold for Wavelet Packet Compression

Determine the default global compression threshold for an $N(0,1)$ white noise input.

Create an $N(0,1)$ white noise input. Set the DWT extension mode to periodic. Set the random number generator to the default initial settings for reproducible results.

```
dwtmode('per','nodisp')
rng default
x = randn(512,1);
```

Use `ddencmp` with the `'cmp'` and `'wp'` input arguments to return the default global compression threshold for a wavelet packet transform.

```
[thr,sorh,keepapp,crit] = ddencmp('cmp','wp',x)

thr = 0.6424

sorh =
'h'
```

```
keepapp = 1
```

```
crit =  
'threshold'
```

Compare with the default values returned for denoising.

```
[thr,sorh,keepapp,crit] = ddencomp('den','wp',x)
```

```
thr = 4.1074
```

```
sorh =  
'h'
```

```
keepapp = 1
```

```
crit =  
'sure'
```

Input Arguments

in1 — Purpose

'den' | 'cmp'

Purpose of `ddencomp` output, specified as:

- 'den' — Denoising
- 'cmp' — Compression

in2 — Transform type

'wv' | 'wp'

Transform type to be used for denoising or compression, specified as:

- 'wv' — Critically sampled discrete wavelet transform. This output can be used with `wdencomp`.
- 'wp' — Critically sampled wavelet packet transform. This output can be used with `wpdencmp`.

x — Input data

real-valued vector or matrix

Input data to be denoised or compressed, specified as a real-valued vector or 2-D matrix.

Data Types: `double`

Output Arguments

thr — Threshold

real number

Threshold for denoising or compression, returned as a real number. Use this output with `wdencomp` or `wpdencmp`.

sorh — Thresholding type

character

Thresholding type for denoising or compression, returned as a character.

- 's' — Soft thresholding
- 'h' — Hard thresholding

Use this output with `wdencomp` or `wpdencmp`.

keepapp — Threshold approximation setting

1 (default)

Threshold approximation setting, returned as 1. Use this output with `wdencomp` or `wpdencmp`. If `keepapp = 1`, the approximation coefficients are not thresholded.

crit — Entropy type

character vector

Entropy type when denoising or compressing with wavelet packets, returned as a character vector. Use this output only with `wpdencmp`. See `wentropy` for more information.

References

- [1] Donoho, D. L. "De-noising by Soft-Thresholding." *IEEE Transactions on Information Theory*, Vol. 42, Number 3, pp. 613–627, 1995.

[2] Donoho, D. L., and Johnstone, I. M. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*, Vol. 81, pp. 425-455, 1994.

[3] Donoho, D. L., and I. M. Johnstone. "Ideal denoising in an orthonormal basis chosen from a library of bases." *Comptes Rendus Acad. Sci. Paris, Ser. I*, Vol. 319, pp. 1317-1322, 1994.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

`wdecmp` | `wdenoise` | `wenergy` | `wpdencmp`

Introduced before R2006a

dddtree

Dual-tree and double-density 1-D wavelet transform

Syntax

```
wt = dddtree(tytree,x,level,fdf,df)
wt = dddtree(tytree,x,level,fname)
wt = dddtree(tytree,x,level,fname1,fname2)
```

Description

`wt = dddtree(tytree,x,level,fdf,df)` returns the `tytree` discrete wavelet transform (DWT) of the 1-D input signal, `x`, down to level, `level`. The wavelet transform uses the decomposition (analysis) filters, `fdf`, for the first level and the analysis filters, `df`, for subsequent levels. Supported wavelet transforms are the critically sampled DWT, double-density, dual-tree complex, and dual-tree double-density complex wavelet transform. The critically sampled DWT is a filter bank decomposition in an orthogonal or biorthogonal basis (nonredundant). The other wavelet transforms are oversampled filter banks.

`wt = dddtree(tytree,x,level,fname)` uses the filters specified by `fname` to obtain the wavelet transform. Valid filter specifications depend on the type of wavelet transform. See `dtfilters` for details.

`wt = dddtree(tytree,x,level,fname1,fname2)` uses the filters specified in `fname1` for the first stage of the dual-tree wavelet transform and the filters specified in `fname2` for subsequent stages of the dual-tree wavelet transform. Specifying different filters for stage 1 is valid and necessary only when `tytree` is `'cplxdt'` or `'cplxddd'`.

Examples

Complex Dual-Tree Wavelet Transform

Obtain the complex dual-tree wavelet transform of the noisy Doppler signal. The FIR filters in the first and subsequent stages result in an approximately analytic wavelet as required.

Create the first-stage analysis filters for the two trees.

```
Faf{1} = [0          0
          -0.0884   -0.0112
           0.0884    0.0112
           0.6959    0.0884
           0.6959    0.0884
           0.0884   -0.6959
          -0.0884    0.6959
           0.0112   -0.0884
           0.0112   -0.0884
           0          0];
Faf{2} = [ 0.0112  0
          0.0112  0
          -0.0884 -0.0884
           0.0884 -0.0884
           0.6959  0.6959
           0.6959 -0.6959
           0.0884  0.0884
          -0.0884  0.0884
           0       0.0112
           0      -0.0112];
```

Create the analysis filters for subsequent stages of the multiresolution analysis.

```
af{1} = [ 0.0352          0
          0          0
          -0.0883   -0.1143
           0.2339          0
           0.7603    0.5875
           0.5875   -0.7603
           0         0.2339
          -0.1143    0.0883
           0          0
           0        -0.0352];
af{2} = [0   -0.0352
          0    0
          -0.1143  0.0883
```



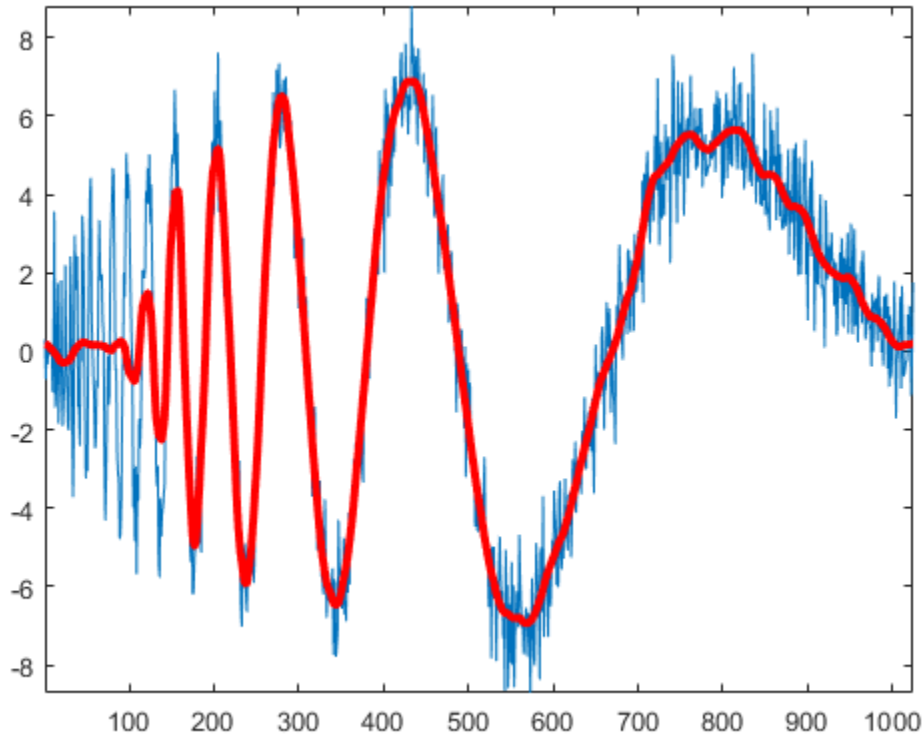
```
      0      0.2339
0.5875 -0.7603
0.7603  0.5875
0.2339      0
-0.0883 -0.1143
      0      0
0.0352      0];
```

Load the noisy Doppler signal and obtain the complex dual-tree wavelet transform down to level 4.

```
load noisdopp;
wt = dddtree('cplxdt',noisdopp,4,Faf,af);
```

Plot an approximation based on the level-four approximation coefficients.

```
xapp = dddtreecfs('r',wt,'scale',{5});
plot(noisdopp); hold on;
plot(cell2mat(xapp),'r','linewidth',3);
axis tight;
```



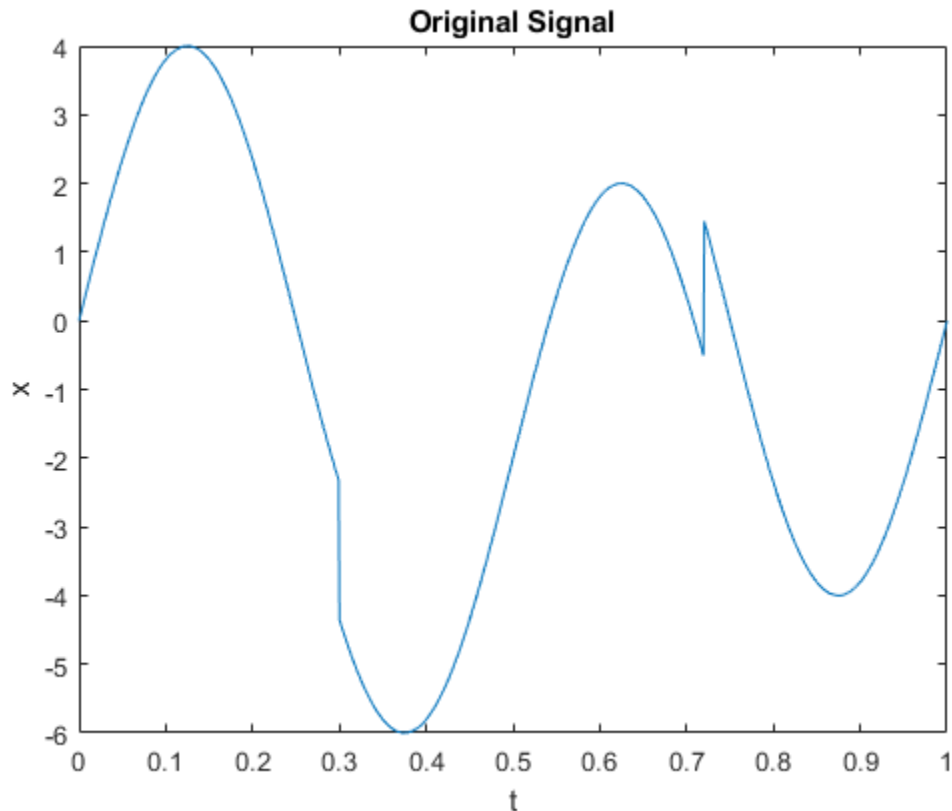
Double-Density Wavelet Transform

Obtain the double-density wavelet transform of a signal with two discontinuities. Use the level-one detail coefficients to localize the discontinuities.

Create a signal consisting of a 2-Hz sine wave with a duration of 1 second. The sine wave has discontinuities at 0.3 and 0.72 seconds.

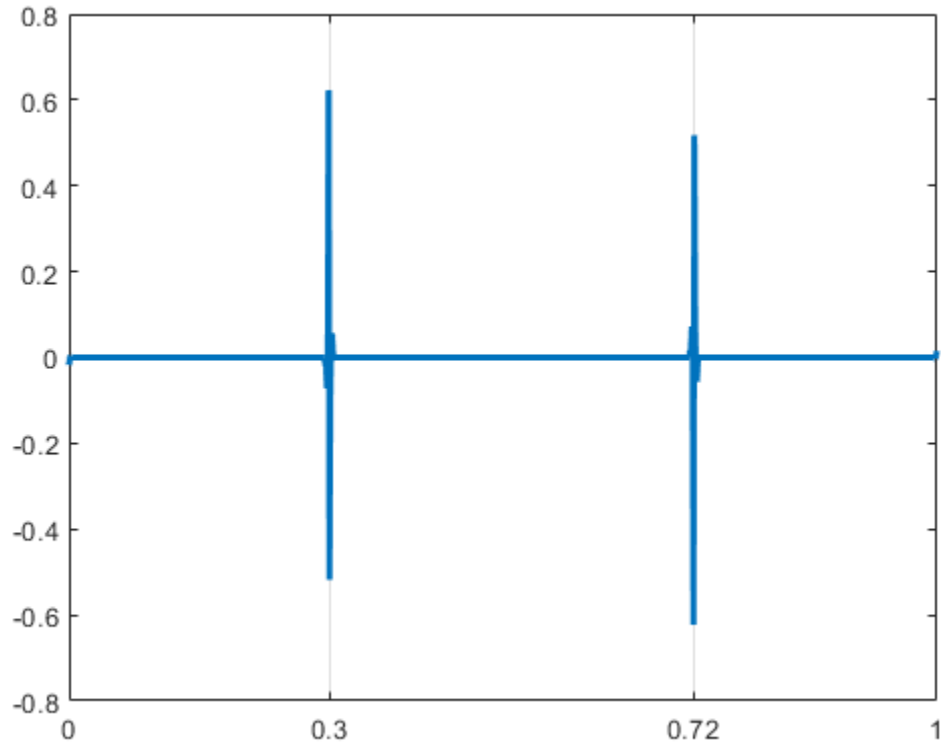
```
N = 1024;  
t = linspace(0,1,1024);  
x = 4*sin(4*pi*t);
```

```
x = x - sign(t - .3) - sign(.72 - t);
plot(t,x); xlabel('t'); ylabel('x');
title('Original Signal');
```



Obtain the double-density wavelet transform of the signal, reconstruct an approximation based on the level-one detail coefficients, and plot the result.

```
wt = dddtree('ddt',x,1,'filters1');
wt.cfs{2} = zeros(1,512);
xrec = idddtree(wt);
plot(t,xrec,'linewidth',2)
set(gca,'xtick',[0 0.3 0.72 1]); set(gca,'xgrid','on');
```



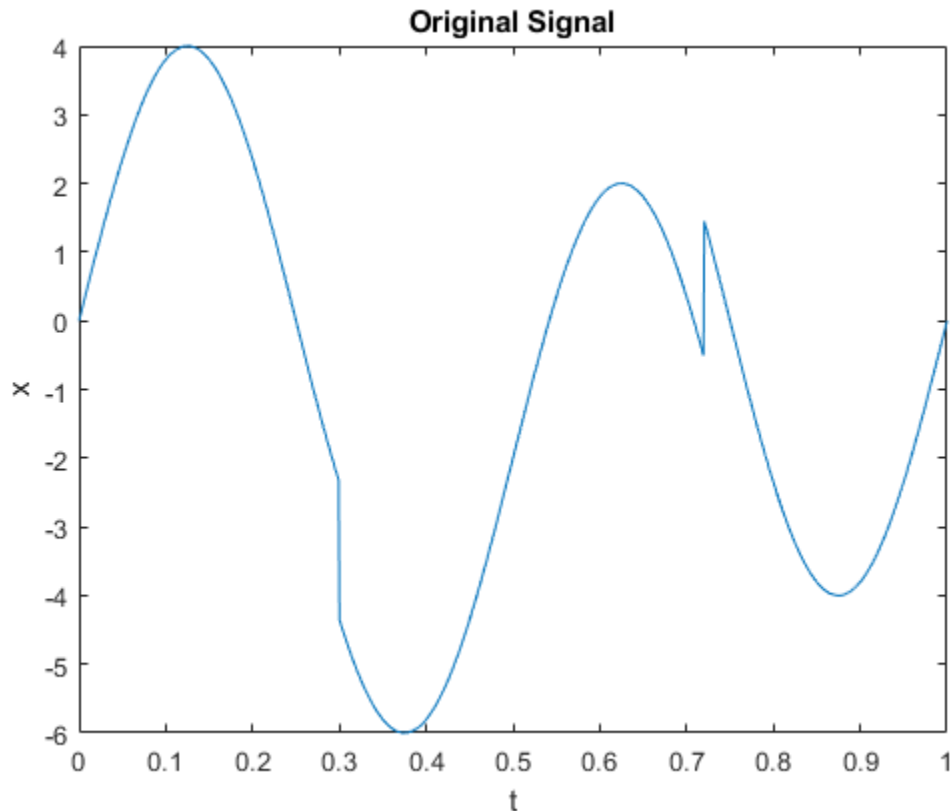
First-Level Detail Coefficients Approximation — Complex Dual-Tree

Obtain the complex dual-tree wavelet transform of a signal with two discontinuities. Use the first-level detail coefficients to localize the discontinuities.

Create a signal consisting of a 2-Hz sine wave with a duration of 1 second. The sine wave has discontinuities at 0.3 and 0.72 seconds.

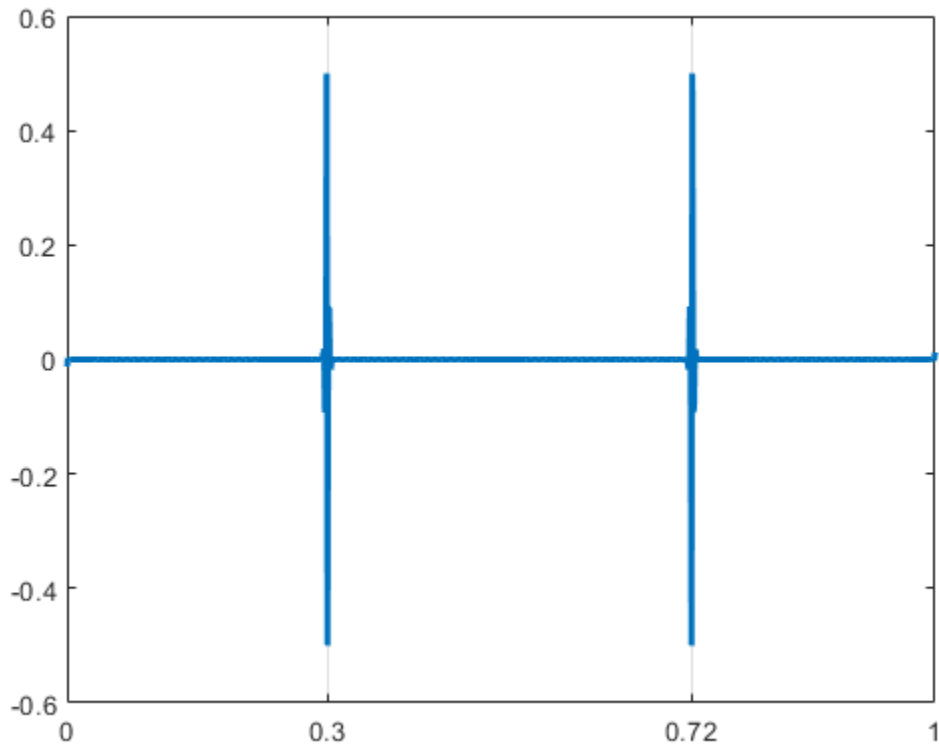
```
N = 1024;  
t = linspace(0,1,1024);  
x = 4*sin(4*pi*t);
```

```
x = x - sign(t - .3) - sign(.72 - t);
plot(t,x); xlabel('t'); ylabel('x');
title('Original Signal');
```



Obtain the dual-tree wavelet transform of the signal, reconstruct an approximation based on the level-one detail coefficients, and plot the result.

```
wt = dddtree('cplxdt',x,1,'FSfarras','qshift06');
wt.cfs{2} = zeros(1,512,2);
xrec = idddtree(wt);
plot(t,xrec,'linewidth',2)
set(gca,'xtick',[0 0.3 0.72 1]); set(gca,'xgrid','on');
```



Input Arguments

typetree — Type of wavelet decomposition

'dwt' | 'ddt' | 'cplxdt' | 'cplxdddtt'

Type of wavelet decomposition, specified as one of 'dwt', 'ddt', 'cplxdt', or 'cplxdddtt'. The type, 'dwt', gives a critically sampled (nonredundant) discrete wavelet transform. The other decomposition types produce oversampled wavelet transforms. 'ddt' produces a double-density wavelet transform. 'cplxdt' produces a dual-tree complex wavelet transform. 'cplxdddtt' produces a double-density dual-tree complex wavelet transform.

x — Input signal

vector

Input signal, specified as an even-length row or column vector. If L is the value of the `level` of the wavelet decomposition, 2^L must divide the length of x . Additionally, the length of the signal must be greater than or equal to the product of the maximum length of the decomposition (analysis) filters and $2^{(L-1)}$.

Data Types: double

level — Level of wavelet decomposition

positive integer

Level of the wavelet decomposition, specified as an integer. If L is the value of `level`, 2^L must divide the length of x . Additionally, the length of the signal must be greater than or equal to the product of the maximum length of the decomposition (analysis) filters and $2^{(L-1)}$.

Data Types: double

fdf — Level-one analysis filters

matrix | cell array

The level-one analysis filters, specified as a matrix or cell array of matrices. Specify `fdf` as a matrix when `typetree` is `'dwt'` or `'ddt'`. The size and structure of the matrix depend on the `typetree` input as follows:

- `'dwt'` — This is the critically sampled discrete wavelet transform. In this case, `fdf` is a two-column matrix with the lowpass (scaling) filter in the first column and the highpass (wavelet) filter in the second column.
- `'ddt'` — This is the double-density wavelet transform. The double-density DWT is a three-channel perfect reconstruction filter bank. `fdf` is a three-column matrix with the lowpass (scaling) filter in the first column and the two highpass (wavelet) filters in the second and third columns. In the double-density wavelet transform, the single lowpass and two highpass filters constitute a three-channel perfect reconstruction filter bank. This is equivalent to the three filters forming a tight frame. You cannot arbitrarily choose the two wavelet filters in the double-density DWT. The three filters together must form a tight frame.

Specify `fdf` as a 1-by-2 cell array of matrices when `typetree` is a dual-tree transform, `'cplxdt'` or `'cplxddt'`. The size and structure of the matrix elements depend on the `typetree` input as follows:

- For the dual-tree complex wavelet transform, 'cplxdt', `fdf{1}` is a two-column matrix containing the lowpass (scaling) filter and highpass (wavelet) filters for the first tree. The scaling filter is the first column and the wavelet filter is the second column. `fdf{2}` is a two-column matrix containing the lowpass (scaling) and highpass (wavelet) filters for the second tree. The scaling filter is the first column and the wavelet filter is the second column.
- For the double-density dual-tree complex wavelet transform, 'cplxddd', `fdf{1}` is a three-column matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the first tree and `fdf{2}` is a three-column matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the second tree.

Data Types: `double`

df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1, specified as a matrix or cell array of matrices. Specify `df` as a matrix when `typetree` is 'dwt' or 'ddt'. The size and structure of the matrix depend on the `typetree` input as follows:

- 'dwt' — This is the critically sampled discrete wavelet transform. In this case, `df` is a two-column matrix with the lowpass (scaling) filter in the first column and the highpass (wavelet) filter in the second column. For the critically sampled orthogonal or biorthogonal DWT, the filters in `df` and `fdf` must be identical.
- 'ddt' — This is the double-density wavelet transform. The double-density DWT is a three-channel perfect reconstruction filter bank. `df` is a three-column matrix with the lowpass (scaling) filter in the first column and the two highpass (wavelet) filters in the second and third columns. In the double-density wavelet transform, the single lowpass and two highpass filters must constitute a three-channel perfect reconstruction filter bank. This is equivalent to the three filters forming a tight frame. For the double-density DWT, the filters in `df` and `fdf` must be identical.

Specify `df` as a 1-by-2 cell array of matrices when `typetree` is a dual-tree transform, 'cplxdt' or 'cplxddd'. For dual-tree transforms, the filters in `fdf` and `df` must be different. The size and structure of the matrix elements in the cell array depend on the `typetree` input as follows:

- For the dual-tree complex wavelet transform, 'cplxdt', `df{1}` is a two-column matrix containing the lowpass (scaling) and highpass (wavelet) filters for the first tree. The scaling filter is the first column and the wavelet filter is the second column. `df{2}` is a two-column matrix containing the lowpass (scaling) and highpass (wavelet) filters

for the second tree. The scaling filter is the first column and the wavelet filter is the second column.

- For the double-density dual-tree complex wavelet transform, 'cplxdddtt', $df\{1\}$ is a three-column matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the first tree and $df\{2\}$ is a three-column matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the second tree.

Data Types: double

fname — Filter name

character vector | string scalar

Filter name, specified as a character vector or string scalar. For the critically sampled DWT, specify any valid orthogonal or biorthogonal wavelet filter. See `wfilters` for details. For the double-density wavelet transform, 'ddt', valid choices are 'filters1' and 'filters2'. For the complex dual-tree wavelet transform, valid choices are 'dtfP' with $P = 1, 2, 3, 4$. For the double-density dual-tree wavelet transform, the only valid choice is 'dddttf1'. See `dtfilters` for more details on valid filter names for the oversampled wavelet filter banks.

Data Types: char

fname1 — First-stage filter name

character vector | string scalar

First-stage filter name, specified as a character vector or string scalar. Specifying a different filter for the first stage is valid and necessary only in the dual-tree transforms, 'cplxdt' and 'cplxdddtt'. In the complex dual-tree wavelet transform, you can use any valid wavelet filter for the first stage. In the double-density dual-tree wavelet transform, the first-stage filters must form a three-channel perfect reconstruction filter bank.

Data Types: char

fname2 — Filter name for stages > 1

character vector | string scalar

Filter name for stages > 1, specified as a character vector or string scalar. You must specify a first-level filter that is different from the wavelet and scaling filters in subsequent levels when using the dual-tree wavelet transforms, 'cplxdt' or 'cplxdddtt'. See `dtfilters` for valid choices.

Data Types: char

Output Arguments

wt — Wavelet transform

structure

Wavelet transform, returned as a structure with these fields:

type — Type of wavelet decomposition (filter bank)

'dwt' | 'ddt' | 'cplxdt' | 'cplxdddtt'

Type of wavelet decomposition (filter bank) used in the analysis, returned as one of 'dwt', 'ddt', 'cplxdt', or 'cplxdddtt'. The type, 'dwt', gives a critically sampled discrete wavelet transform. The other types correspond to oversampled wavelet transforms. 'ddt' is a double-density wavelet transform, 'cplxdt' is a dual-tree complex wavelet transform, and 'cplxdddtt' is a double-density dual-tree complex wavelet transform.

level — Level of the wavelet decomposition

positive integer

Level of wavelet decomposition, returned as a positive integer.

filters — Decomposition (analysis) and reconstruction (synthesis) filters

structure

Decomposition (analysis) and reconstruction (synthesis) filters, returned as a structure with these fields:

Fdf — First-stage analysis filters

matrix | cell array

First-stage analysis filters, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

Df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1 , returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the analysis filters for the corresponding tree.

FrF — First-level reconstruction filters

matrix | cell array

First-level reconstruction filters, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

Rf — Reconstruction filters for levels > 1

matrix | cell array

Reconstruction filters for levels > 1 , returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the synthesis filters for the corresponding tree.

cfs — Wavelet transform coefficients

cell array of matrices

Wavelet transform coefficients, returned as a 1 -by- $(\text{level}+1)$ cell array of matrices. The size and structure of the matrix elements of the cell array depend on the type of wavelet transform, `typetree`, as follows:

- 'dwt' — `cfs{j}`

- $j = 1, 2, \dots$ level is the level.
- $cfs\{\text{level}+1\}$ are the lowpass, or scaling, coefficients.
- 'ddt' — $cfs\{j\}(:, :, k)$
 - $j = 1, 2, \dots$ level is the level.
 - $k = 1, 2$ is the wavelet filter.
 - $cfs\{\text{level}+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdt' — $cfs\{j\}(:, :, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'cplxddd' — $cfs\{j\}(:, :, k, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $k = 1, 2$ is the wavelet filter.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.

See Also

dddtree2 | dddtreecfs | dtfilters | idddtree

Topics

“Analytic Wavelets Using the Dual-Tree Wavelet Transform”
“Critically Sampled and Oversampled Wavelet Filter Banks”

Introduced in R2013b

dddtreecfs

Extract dual-tree/double-density wavelet coefficients or projections

Syntax

```
out = dddtreecfs(outputtype,wt,outputspec,outputindices)
out = dddtreecfs(outputtype,wt,outputspec,outputindices,'plot')
```

Description

`out = dddtreecfs(outputtype,wt,outputspec,outputindices)` extracts the coefficients or subspace projections from the 1-D or 2-D wavelet decomposition, `wt`. If `outputtype` equals 'e', `out` contains wavelet or scaling coefficients. If `outputtype` equals 'r', `out` contains wavelet or scaling subspace projections (reconstructions).

`out = dddtreecfs(outputtype,wt,outputspec,outputindices,'plot')` plots the signal or image reconstruction or specified analysis coefficients. You can include the 'plot' option anywhere after the `wt` input.

Examples

Reconstruction from 1-D Complex Dual-Tree Wavelet Transform

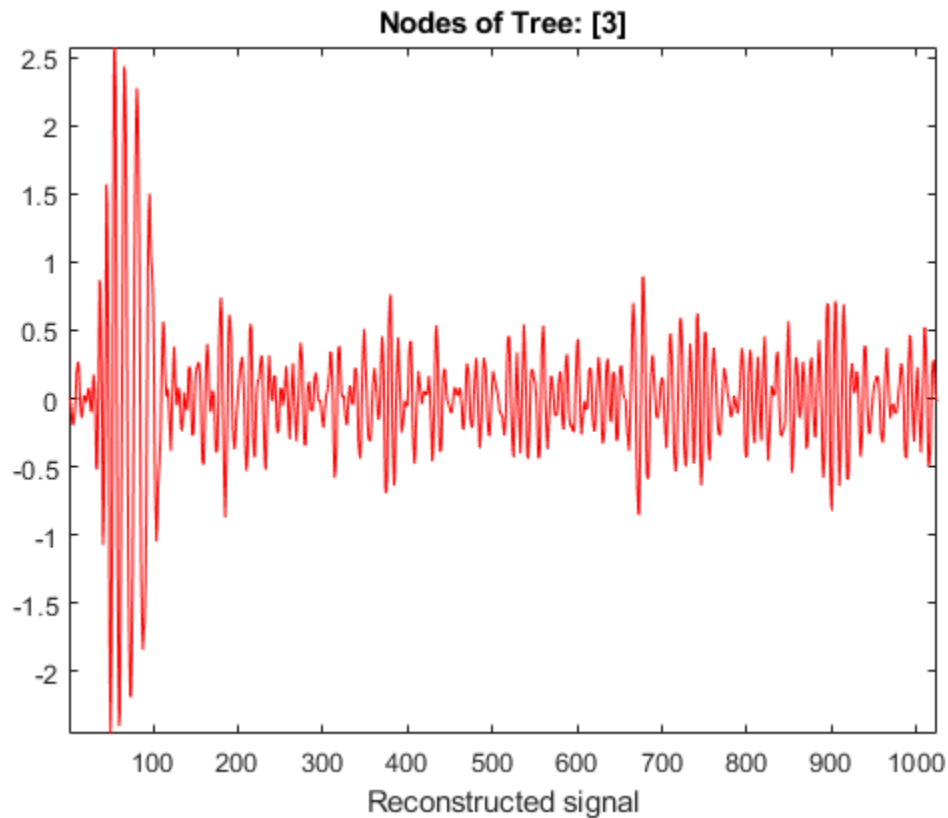
Obtain the complex dual-tree wavelet transform of the 1-D noisy Doppler signal. Reconstruct an approximation based on the level-three detail coefficients

Load the noisy Doppler signal. Obtain the complex dual-tree transform down to level 3.

```
load noisdopp;
wt = dddtree('cplxdt',noisdopp,3,'dtf1');
```

Plot a reconstruction of the original signal based on the level-three detail coefficients

```
xr = dddtreecfs('r',wt,'scale',{3},'plot');
```



Coefficients from 1-D Complex Dual-Tree Wavelet Transform

Load the noisy Doppler signal. Obtain the complex dual-tree transform down to level 3.

```
load noisdopp;  
wt = dddtree('cplxdt',noisdopp,3,'dtf1');
```

Create a cell array of vectors to obtain the second- and third-level detail coefficients from each of the wavelet filter bank trees.

```
outputindices = {[2 1]; [2 2]; [3 1]; [3 2]};
```

The first element of each vector in the cell array denotes the level, or stage. The second element denotes the tree.

Extract the detail coefficients.

```
out = dddtrecfs('e',wt,'ind',outputindices);
```

out is a 1-by-4 cell array. The cell array elements contain the wavelet coefficients corresponding to the elements in outputindices. For example, out{1} contains the level-two detail coefficients from the first tree.

1-D Complex Dual-Tree Wavelet Transform Structure

Load the noisy Doppler signal. Obtain the complex dual-tree transform down to level 3.

```
load noisdopp;
wt = dddtree('cplxdt',noisdopp,3,'dtf1');
```

Create a cell array of vectors to obtain the second- and third-level detail coefficients from each of the wavelet filter bank trees.

```
outputindices = {[2 1]; [2 2]; [3 1];[3 2]};
```

The first element of each vector in the cell array denotes the level, or stage. The second element denotes the tree.

Create a structure array identical to the wt output of dddtree with all the coefficients equal to zero except the first- and second-level detail coefficients.

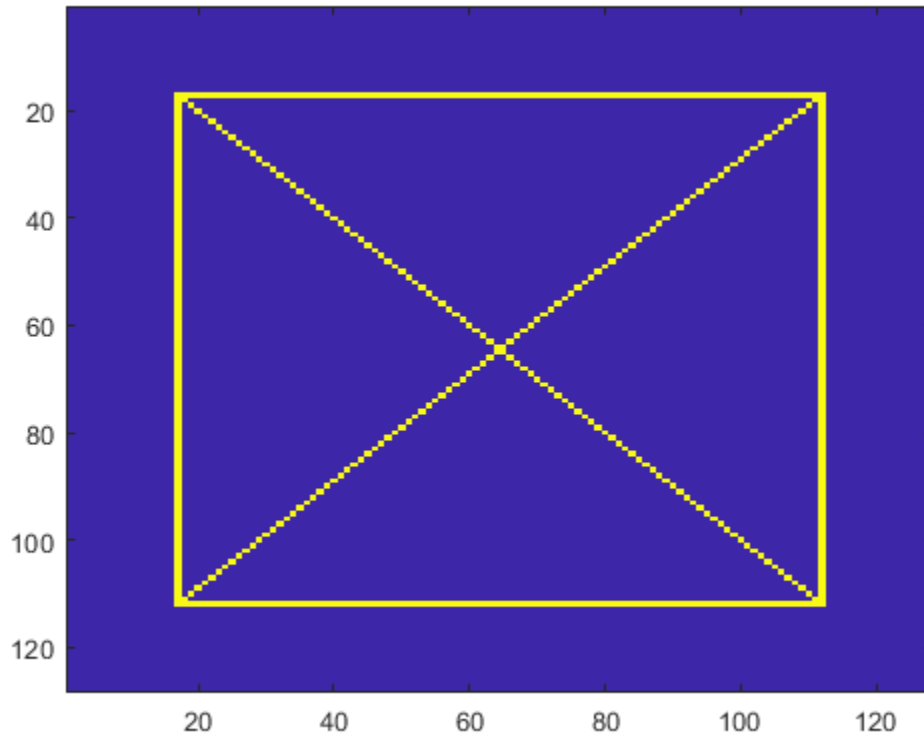
```
out = dddtrecfs('e',wt,'cumind',outputindices);
```

Extract Diagonal Features from Image

Use the complex dual-tree wavelet transform to isolate diagonal features in an image at +45 and -45 degrees.

Load and display the xbox image.

```
load xbox;  
imagesc(xbox)
```

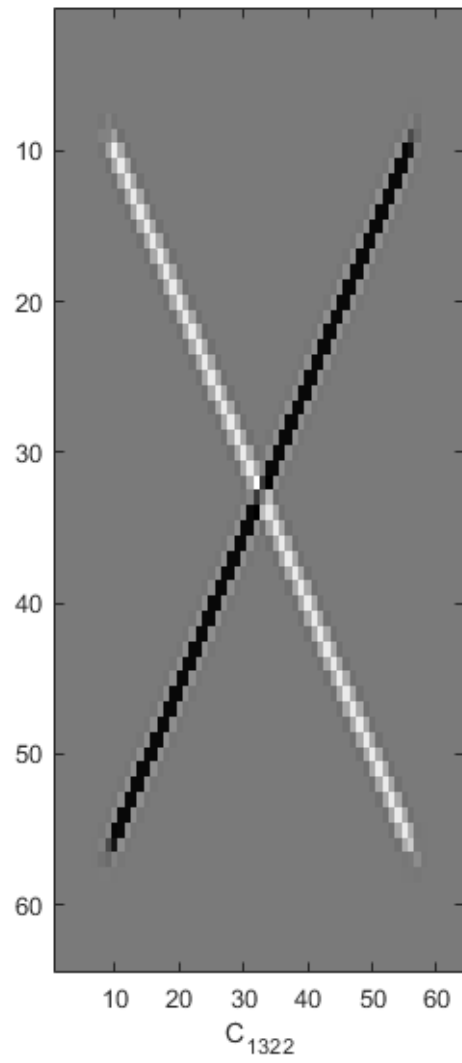
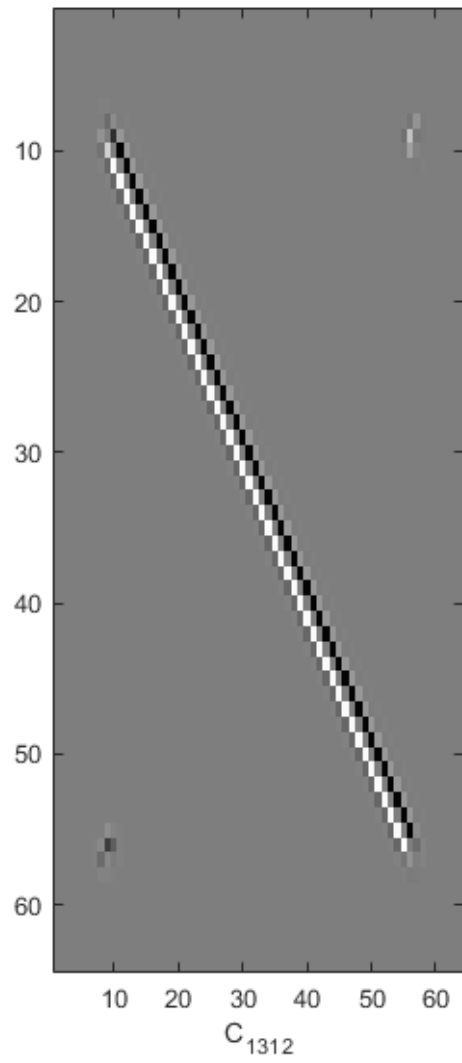


Obtain the complex dual-tree wavelet transform down to level 3.

```
fdf = dtfilters('FSfarras');  
df = dtfilters('qshift10');  
wt = dddtree2('cplxdt',xbox,3,fdf,df);
```

Isolate the +45 and -45 diagonal image features in the level-one wavelet coefficients. Plot the result.

```
out = dddtreecfs('e',wt,'ind',{[1 3 1 2]; [1 3 2 2]},'plot');
```

Input Arguments

outputtype — Output type

'e' | 'r'

Output type, specified as 'e' or 'r'. Use 'e' to obtain the scaling or wavelet coefficients. Use 'r' to obtain a projection, or reconstruction, onto the appropriate scaling or wavelet subspace.

wt — Wavelet transform

structure

Wavelet transform, specified as a structure. The structure array is the output of `dddtree` or `dddtree2`.

outputspec — Output specification

'lowpass' | 'scale' | 'ind' | 'cumind'

Output specification, specified as one of 'lowpass', 'scale', 'ind', or 'cumind'. The output specifications are defined as follows:

- 'lowpass' — Outputs the lowpass, or scaling, coefficients or a signal/image approximation based on the scaling coefficients. If you set the output specification to 'lowpass', do not specify `outputindices`. If the `outputtype` is 'e', `out` is a structure array with fields identical to the input structure array `wt` except that all wavelet (detail) coefficients are equal to zero. If the `outputtype` is 'r', `out` is a signal or image approximation based on the scaling coefficients. The signal or image approximation is equal in size to the original input to `dddtree` or `dddtree2`.
- 'scale' — Outputs the coefficients or a signal/image approximation based on the scales specified in `outputindices`. If the `outputtype` is 'e', `out` is a cell array of structure arrays. The fields of the structure arrays in `out` are identical to the fields of the input structure array `wt`. The coefficients in the `cfs` field are all equal to zero except the coefficients corresponding to the scales in `outputindices`. If the `outputtype` is 'r', `out` is a signal or image approximation based on the scales in `outputindices`. The signal or image approximation is equal in size to the original input to `dddtree` or `dddtree2`.
- 'ind' — Outputs the coefficients or a signal/image approximation based on the tree-position indices specified in `outputindices`. If the `outputtype` is 'e', `out` is a cell array of vectors or matrices containing the coefficients specified by the tree-position indices in `outputindices`. If the `outputtype` is 'r', `out` is a cell array of vectors or

matrices containing signal or image approximations based on the corresponding tree-position indices in `outputindices`.

- `'cumind'` — Outputs the coefficients or a signal/image approximation based on the tree-position indices specified in `outputindices`. If the `outputtype` is `'e'`, `out` is a structure array. The fields of the structure array are identical to the fields of the input structure array `wt`. The coefficients in the `cfs` field are all equal to zero except the coefficients corresponding to the tree positions in `outputindices`. If the `outputtype` is `'r'`, `out` is a signal or image approximation based on the coefficients corresponding to the tree-position indices in `outputindices`.

Example: `'ind', {[1 1]; [1 2]}`

outputindices — Output indices

cell array

Output indices, specified as a cell array with scalar or vector elements. If `outputspec` equals `'scale'`, a scalar element selects the corresponding element in the `cfs` field of `wt`. If `outputspec` equals `'ind'` or `'cumind'`, the elements of `outputspec` are row vectors. The first element of the row vector corresponds to the element in the `cfs` field of `wt`. Subsequent elements in the row vector correspond to the indices of the array contained in the cell array element.

Example: `'scale', {1;2;3}`

Output Arguments

out — Signal or image reconstruction or coefficients

cell array | structure | vector | matrix

Signal or image reconstruction or coefficients, returned as a vector, matrix, structure array, cell array of vectors or matrices, or cell array of structure arrays. The form of `out` depends on the value of `outputspec` and `outputindices`.

See Also

`dddtree` | `dddtree2` | `plotdt`

Introduced in R2013b

dddtree2

Dual-tree and double-density 2-D wavelet transform

Syntax

```
wt = dddtree2(typetree,x,level,fdf,df)
wt = dddtree2(typetree,x,level,fname)
wt = dddtree2(typetree,x,level,fname1,fname2)
```

Description

`wt = dddtree2(typetree,x,level,fdf,df)` returns the `typetree` discrete wavelet transform of the 2-D input image, `x`, down to `level`, `level`. The wavelet transform uses the decomposition (analysis) filters, `fdf`, for the first level and the analysis filters, `df`, for subsequent levels. Supported wavelet transforms are the critically sampled DWT, double-density, real oriented dual-tree, complex oriented dual-tree, real oriented dual-tree double-density, and complex oriented dual-tree double-density wavelet transform. The critically sampled DWT is a filter bank decomposition in an orthogonal or biorthogonal basis (nonredundant). The other wavelet transforms are oversampled filter banks with differing degrees of directional selectivity.

`wt = dddtree2(typetree,x,level,fname)` uses the filters specified by `fname` to obtain the wavelet transform. Valid filter specifications depend on the type of wavelet transform. See `dtfilters` for details.

`wt = dddtree2(typetree,x,level,fname1,fname2)` uses the filters specified in `fname1` for the first stage of the dual-tree wavelet transform and the filters specified in `fname2` for subsequent stages of the dual-tree wavelet transform. Specifying different filters for stage 1 is valid and necessary only when `typetree` is `'realdt'`, `'cplxdt'`, `'realddt'`, or `'cplxddt'`.

Examples

Real Oriented Dual-Tree Wavelets

Visualize the six directional wavelets of the real oriented dual-tree wavelet transform.

Create the first-stage analysis filters for the two trees.

```
Faf{1} = [0      0
          -0.0884 -0.0112
           0.0884  0.0112
           0.6959  0.0884
           0.6959  0.0884
           0.0884 -0.6959
          -0.0884  0.6959
           0.0112 -0.0884
           0.0112 -0.0884
           0      0];
Faf{2} = [ 0.0112  0
          0.0112  0
          -0.0884 -0.0884
           0.0884 -0.0884
           0.6959  0.6959
           0.6959 -0.6959
           0.0884  0.0884
          -0.0884  0.0884
           0      0.0112
           0     -0.0112];
```

Create the analysis filters for subsequent stages of the multiresolution analysis.

```
af{1} = [ 0.0352  0
          0      0
          -0.0883 -0.1143
           0.2339  0
           0.7603  0.5875
           0.5875 -0.7603
           0      0.2339
          -0.1143  0.0883
           0      0
           0     -0.0352];
af{2} = [0     -0.0352
          0      0
          -0.1143  0.0883
           0      0.2339
           0.5875 -0.7603]
```

```
    0.7603    0.5875
    0.2339     0
   -0.0883   -0.1143
         0     0
    0.0352     0];
```

Obtain the real dual-tree wavelet transform of an image of zeros down to level 4.

```
J = 4;
L = 3*2^(J+1);
N = L/2^J;
x = zeros(2*L,3*L);
wt = dddtree2('realdt',x,J,Faf,af);
```

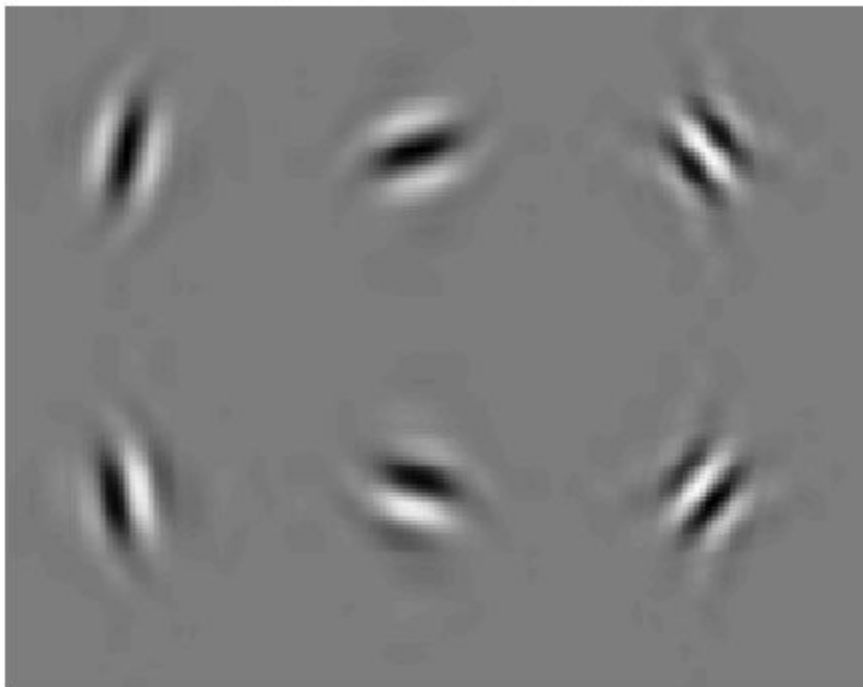
Insert a 1 in one position of the six subbands and invert the wavelet transform.

```
wt.cfs{4}(N/2,N/2+0*N,1,1) = 1;
wt.cfs{4}(N/2,N/2+1*N,2,1) = 1;
wt.cfs{4}(N/2,N/2+2*N,3,1) = 1;
wt.cfs{4}(N/2+N,N/2+0*N,1,2) = 1;
wt.cfs{4}(N/2+N,N/2+1*N,2,2) = 1;
wt.cfs{4}(N/2+N,N/2+2*N,3,2) = 1;
xrec = idddtree2(wt);
```

Visualize the six directional wavelets.

```
imagesc(xrec);
colormap gray; axis off;
title('Real Oriented Dual-Tree Wavelets')
```

Real Oriented Dual-Tree Wavelets

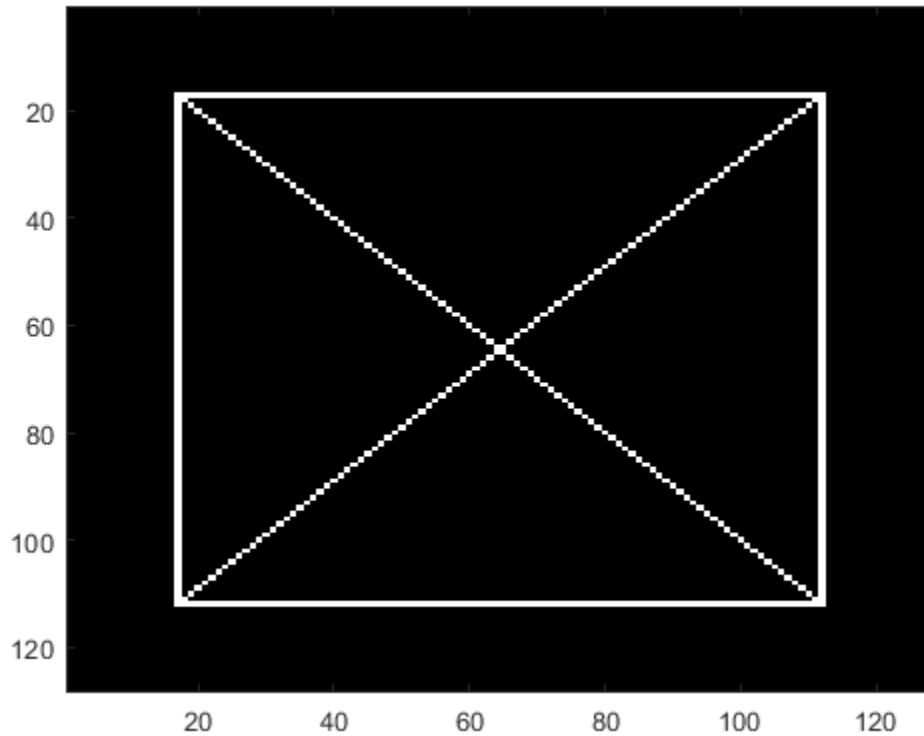


Double-Density Wavelet Transform

Obtain the double-density wavelet transform of an image.

Load the image and obtain the double-density wavelet transform.

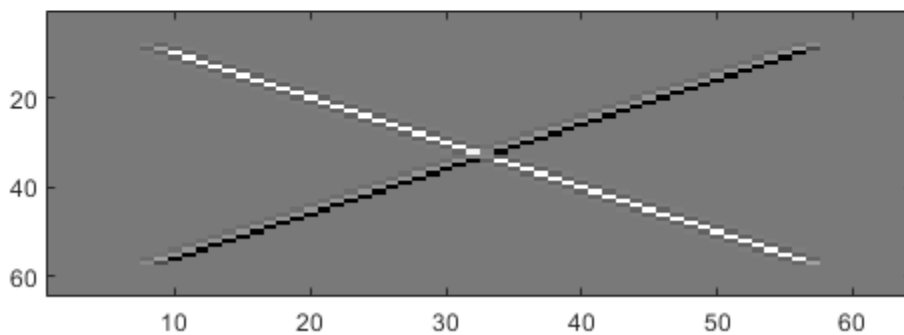
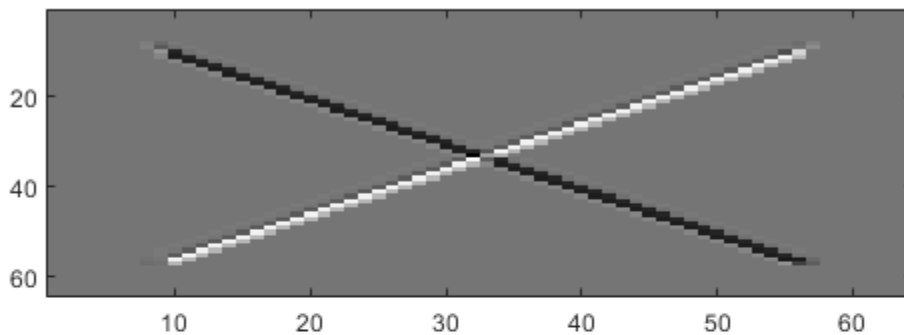
```
load xbox;  
imagesc(xbox); colormap gray;
```



```
wt = dddtree2('ddt',xbox,1,'filters1');
```

Visualize the diagonal details in the two wavelet HH subbands.

```
HH1 = wt.cfs{1}(:, :, 5);  
HH2 = wt.cfs{1}(:, :, 8);  
subplot(211)  
imagesc(HH1);  
colormap gray;  
subplot(212);  
imagesc(HH2);
```

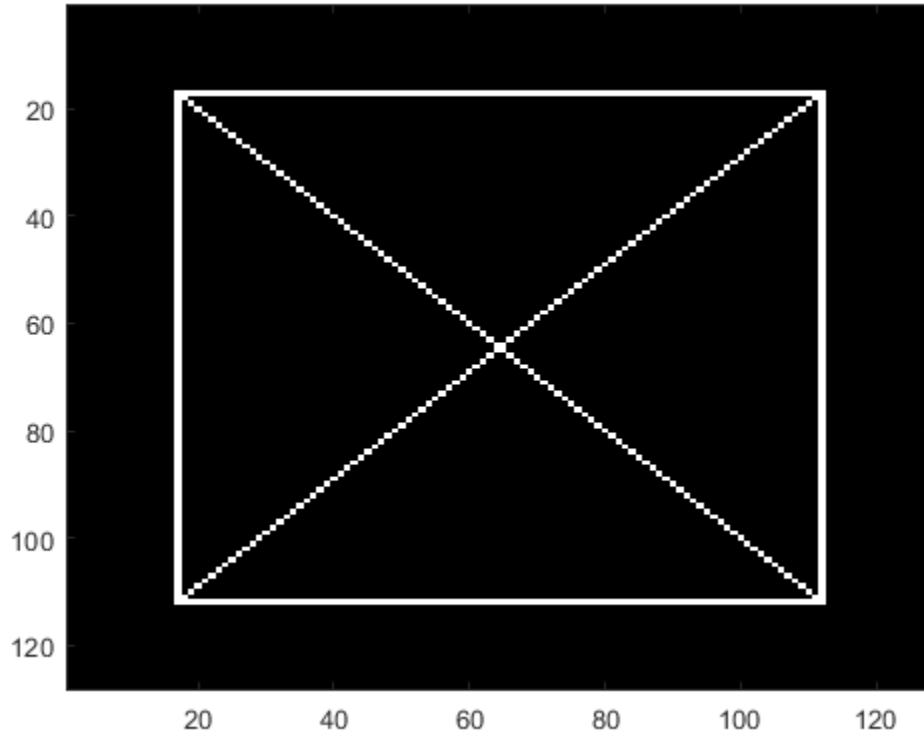



Complex Dual-Tree Wavelet Transform

Obtain the complex dual-tree wavelet transform of an image. Show that the complex dual-tree wavelet transform can detect the two different diagonal directions.

Load the image and obtain the complex dual-tree wavelet transform.

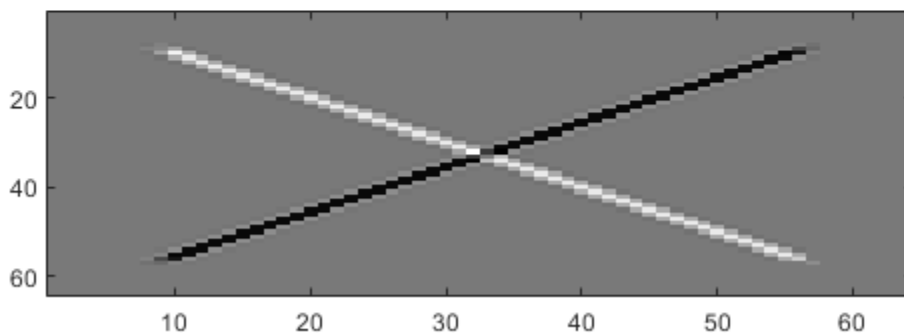
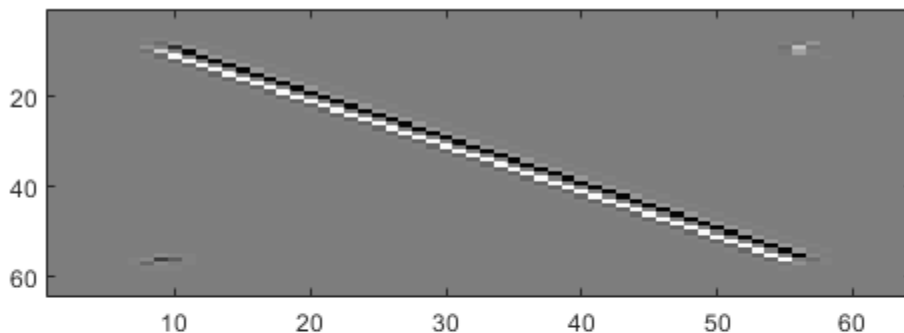
```
load xbox;  
imagesc(xbox); colormap gray;
```



```
wt = dddtree2('cplxdt',xbox,1,'FSfarras','qshift10');
```

Obtain and display the imaginary parts of the 2 trees.

```
waveletcfs = wt.cfs{1};  
subplot(211)  
imagesc(waveletcfs(:,:,3,1,2));  
colormap gray;  
subplot(212)  
imagesc(waveletcfs(:,:,3,2,2));
```



Input Arguments

typetree — Type of wavelet decomposition

'dwt' | 'ddt' | 'realdt' | 'cplxdt' | 'realdddt' | 'cplxdddtt'

Type of wavelet decomposition, specified as one of 'dwt', 'ddt', 'realdt', 'cplxdt', 'realdddt', or 'cplxdddtt'. The type, 'dwt', produces a critically sampled (nonredundant) discrete wavelet transform. The other decomposition types produce oversampled wavelet transforms. 'ddt' produces a double-density wavelet transform with one scaling and two wavelet filters for both row and column filtering. The double-density wavelet transform uses the same filters at all stages. 'realdt' and 'cplxdt'

produce oriented dual-tree wavelet transforms consisting of two and four separable wavelet transforms. 'realdddt' and 'cplxdddt' produce double-density dual-tree wavelet transforms. The dual-tree wavelet transforms use different filters for the first stage (level).

x — Input image

matrix

Input image, specified as a matrix with even-length row and column dimensions. Both the row and column dimensions must be divisible by 2^L , where L is the level of the wavelet transform. Additionally, the minimum of the row and column dimensions of the image must be greater than or equal to the product of the maximum length of the decomposition (analysis) filters and $2^{(L-1)}$.

Data Types: double

level — Level of wavelet decomposition

integer

Level of the wavelet decomposition, specified as a positive integer. If L is the value of level, 2^L must divide both the row and column dimensions of `x`. Additionally, the minimum of the row and column dimensions of the image must be greater than or equal to the product of the maximum length of the decomposition (analysis) filters and $2^{(L-1)}$.

fdf — Level-one analysis filters

matrix | cell array

The level-one analysis filters, specified as a matrix or cell array of matrices. Specify `fdf` as a matrix when `typetree` is 'dwt' or 'ddt'. The size and structure of the matrix depend on the `typetree` input as follows:

- 'dwt' — This is the critically sampled discrete wavelet transform. In this case, `fdf` is a two-column matrix with the lowpass (scaling) filter in the first column and the highpass (wavelet) filter in the second column.
- 'ddt' — This is the double-density wavelet transform. The double-density DWT is a three-channel perfect reconstruction filter bank. `fdf` is a three-column matrix with the lowpass (scaling) filter in the first column and the two highpass (wavelet) filters in the second and third columns. In the double-density wavelet transform, the single lowpass and two highpass filters constitute a three-channel perfect reconstruction filter bank. This is equivalent to the three filters forming a tight frame. You cannot arbitrarily choose the two wavelet filters in the double-density DWT. The three filters together must form a tight frame.

Specify `fdf` as a 1-by-2 cell array of matrices when `typetree` is a dual-tree transform, `'realdt'`, `'cplxdt'`, `'realdtdt'`, or `'cplxdddtdt'`. The size and structure of the matrix elements in the cell array depend on the `typetree` input as follows:

- For the dual-tree complex wavelet transforms, `'realdt'` and `'cplxdt'`, `fdf{1}` is an N -by-2 matrix containing the lowpass (scaling) and highpass (wavelet) filters for the first tree and `fdf{2}` is an N -by-2 matrix containing the lowpass (scaling) and highpass (wavelet) filters for the second tree.
- For the double-density dual-tree complex wavelet transforms, `'realdtdt'` and `'cplxdddtdt'`, `fdf{1}` is an N -by-3 matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the first tree and `fdf{2}` is an N -by-3 matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the second tree.

df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1, specified as a matrix or cell array of matrices. Specify `df` as a matrix when `typetree` is `'dwt'` or `'ddt'`. The size and structure of the matrix depend on the `typetree` input as follows:

- `'dwt'` — This is the critically sampled discrete wavelet transform. In this case, `df` is a two-column matrix with the lowpass (scaling) filter in the first column and the highpass (wavelet) filter in the second column. For the critically sampled orthogonal or biorthogonal DWT, the filters in `df` and `fdf` must be identical.
- `'ddt'` — This is the double-density wavelet transform. The double-density DWT is a three-channel perfect reconstruction filter bank. `df` is a three-column matrix with the lowpass (scaling) filter in the first column and the two highpass (wavelet) filters in the second and third columns. In the double-density wavelet transform, the single lowpass and two highpass filters constitute a three-channel perfect reconstruction filter bank. This is equivalent to the three filters forming a tight frame. For the double-density DWT, the filters in `df` and `fdf` must be identical.

Specify `df` as a 1-by-2 cell array of matrices when `typetree` is a dual-tree transform, `'realdt'`, `'cplxdt'`, `'realdtdt'`, or `'cplxdddtdt'`. For dual-tree transforms, the filters in `fdf` and `df` must be different. The size and structure of the matrix elements in the cell array depend on the `typetree` input as follows:

- For the dual-tree wavelet transforms, `'realdt'` and `'cplxdt'`, `df{1}` is an N -by-2 matrix containing the lowpass (scaling) and highpass (wavelet) filters for the first tree and `df{2}` is an N -by-2 matrix containing the lowpass (scaling) and highpass (wavelet) filters for the second tree.

- For the double-density dual-tree complex wavelet transforms, 'realdddtt' and 'cplxdddtt', `df{1}` is an N -by-3 matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the first tree and `df{2}` is an N -by-3 matrix containing the lowpass (scaling) and two highpass (wavelet) filters for the second tree.

fname — Filter name

character vector | string scalar

Filter name, specified as a character vector or string scalar. For the critically sampled DWT, specify any valid orthogonal or biorthogonal wavelet filter. See `wfilters` for details. For the redundant wavelet transforms, see `dtfilters` for valid filter names.

fname1 — First-stage filter name

character vector | string scalar

First-stage filter name, specified as a character vector or string scalar. Specifying a first-level filter that is different from the wavelet and scaling filters in subsequent levels is valid and necessary only with the dual-tree wavelet transforms, 'realddt', 'cplxdt', 'realdddtt', and 'cplxdddtt'.

fname2 — Filter name for stages > 1

character vector | string scalar

Filter name for stages > 1, specified as a character vector or string scalar. Specifying a different filter for stages > 1 is valid and necessary only with the dual-tree wavelet transforms, 'realddt', 'cplxdt', 'realdddtt', and 'cplxdddtt'.

Output Arguments

wt — Wavelet transform

structure

Wavelet transform, returned as a structure with these fields:

type — Type of wavelet decomposition (filter bank)

'dwt' | 'ddt' | 'realddt' | 'cplxdt' | 'realdddtt' | 'cplxdddtt'

Type of wavelet decomposition used in the analysis returned as one of 'dwt', 'ddt', 'realddt', 'cplxdt', 'realdddtt', or 'cplxdddtt'. 'dwt' is the critically sampled DWT. 'ddt' produces a double-density wavelet transform with one scaling and two

wavelet filters for both row and column filtering. 'realdt' and 'cplxdt' produce oriented dual-tree wavelet transforms consisting of 2 and 4 separable wavelet transforms. 'realdddt' and 'cplxdddt' produce double-density dual-tree wavelet transforms consisting of two and four separable wavelet transforms.

Level — Level of wavelet decomposition

positive integer

Level of wavelet decomposition, returned as a positive integer.

filters — Decomposition (analysis) and reconstruction (synthesis) filters

structure

Decomposition (analysis) and reconstruction (synthesis) filters, returned as a structure with these fields:

Fdf — First-stage analysis filters

matrix | cell array

First-stage analysis filters, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

Df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the analysis filters for the corresponding tree.

Frf — First-level reconstruction filters

matrix | cell array

First-level reconstruction filters, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

Rf — Reconstruction filters for levels > 1

matrix | cell array

Reconstruction filters for levels > 1, returned as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

cfs — Wavelet transform coefficients

cell array of matrices

Wavelet transform coefficients, specified as a 1-by-(level+1) cell array of matrices. The size and structure of the matrix elements of the cell array depend on the type of wavelet transform, `typetree` as follows:

- 'dwt' — `cfs{j}(:, :, d)`
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - `cfs{level+1}(:, :, :)` are the lowpass, or scaling, coefficients.
- 'ddt' — `cfs{j}(:, :, d)`
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3, 4, 5, 6, 7, 8$ is the orientation.
 - `cfs{level+1}(:, :, :)` are the lowpass, or scaling, coefficients.
- 'realddt' — `cfs{j}(:, :, d, k)`

- $j = 1, 2, \dots$ level is the level.
- $d = 1, 2, 3$ is the orientation.
- $k = 1, 2$ is the wavelet transform tree.
- $cfs\{level+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{level+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'realdddt' — $cfs\{j\}(:, :, d, k)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $cfs\{level+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdddt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{level+1\}(:, :)$ are the lowpass, or scaling, coefficients.

See Also

dddtree | dddtreecfs | dtfilters | idddtree2

Topics

“Analytic Wavelets Using the Dual-Tree Wavelet Transform”

“Critically Sampled and Oversampled Wavelet Filter Banks”

Introduced in R2013b

depo2ind

Node depth-position to node index

Syntax

$N = \text{depo2ind}(\text{ORD}, [D \ P])$

Description

depo2ind is a tree-management utility.

For a tree of order ORD , $N = \text{depo2ind}(\text{ORD}, [D \ P])$ computes the indices N of the nodes whose depths and positions are encoded within $[D, P]$.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

D and P are column vectors. The values of depths D and positions P must be such that $D \geq 0$ and $0 \leq P \leq \text{ORD}^{D-1}$.

Output indices N are such that $0 \leq N < (\text{ORD}^{\max(D)-1}) / \text{ORD} - 1$.

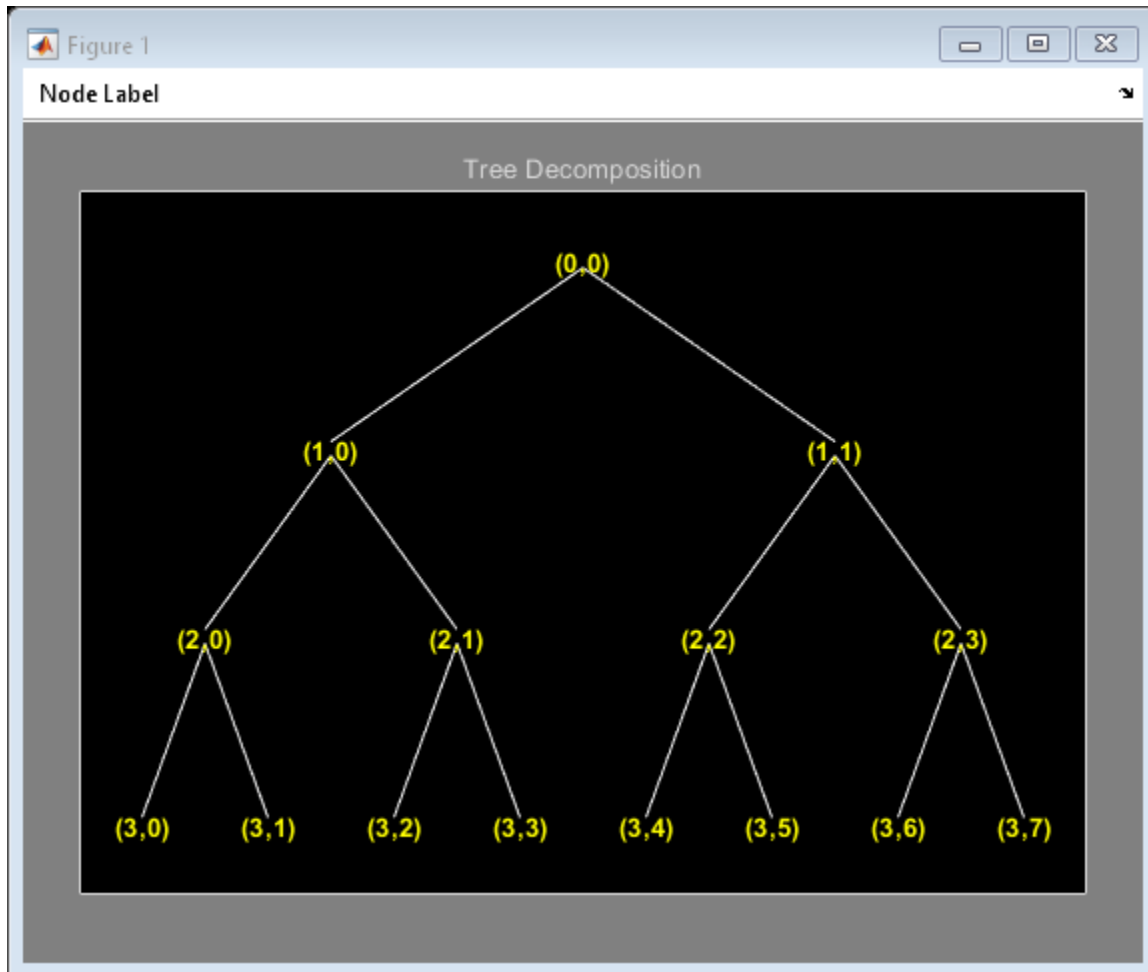
Note that for a column vector X , we have $\text{depo2ind}(0, X) = X$.

Examples

Convert Depth-Position to Node Index

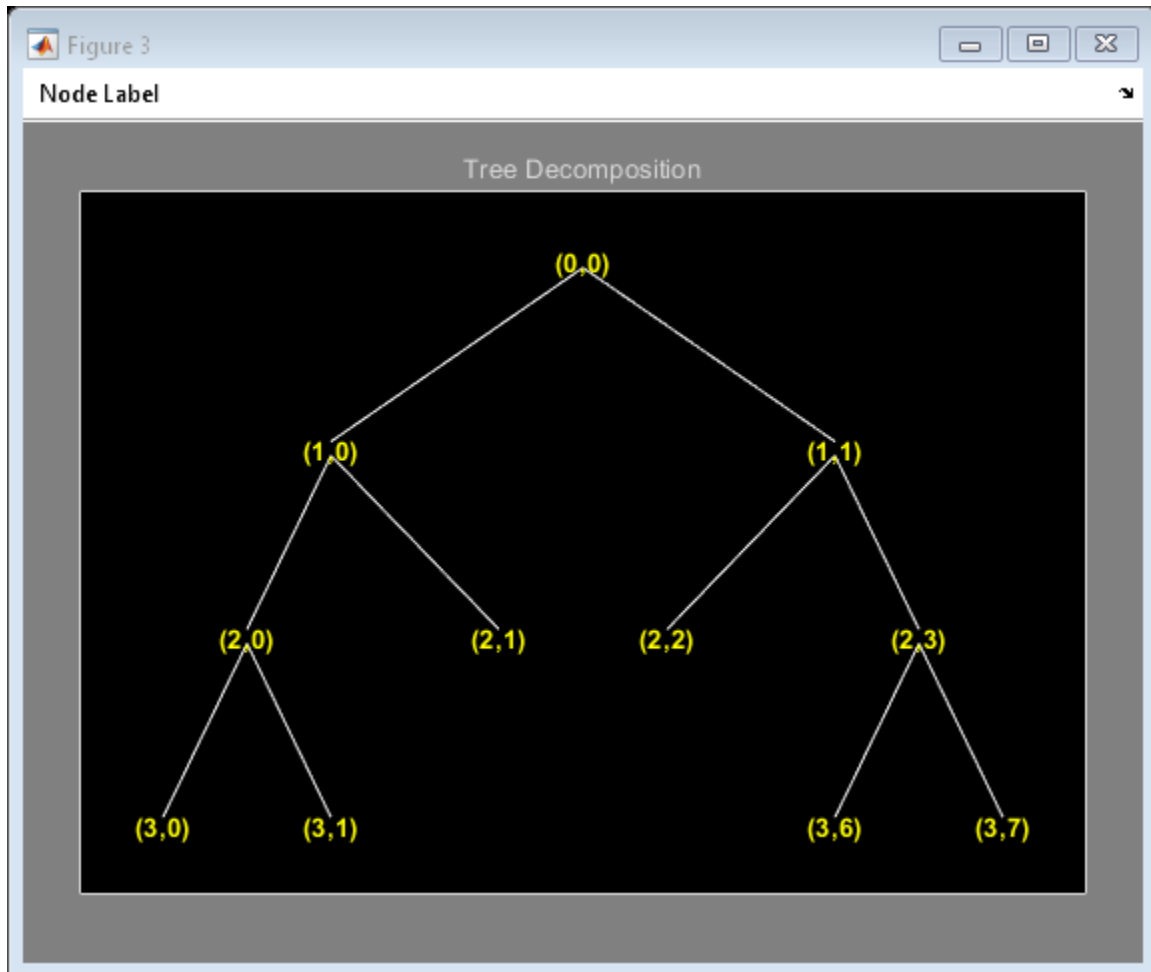
Create a binary tree of depth 3. Plot the tree.

```
ord = 2;  
t = ntree(ord,3);  
plot(t)
```



Merge the nodes of indices 4 and 5. Plot the new tree.

```
t = nodejoin(t,5);  
t = nodejoin(t,4);  
figure  
plot(t)
```



List the depth-position of the tree nodes.

```
aln_depo = allnodes(t, 'depos')
```

```
aln_depo = 11x2
```

```
0    0
1    0
1    1
2    0
```

```
2     1
2     2
2     3
3     0
3     1
3     6
:
```

Convert the depth-position to index.

```
aln_ind = depo2ind(ord,aln_depo)
```

```
aln_ind = 11x1
```

```
0
1
2
3
4
5
6
7
8
13
:
```

See Also

ind2depo

Introduced before R2006a

detcoef

1-D detail coefficients

Syntax

`D = detcoef(C,L,N)`
`D = detcoef(C,L)`

Description

`detcoef` is a one-dimensional wavelet analysis function.

`D = detcoef(C,L,N)` extracts the detail coefficients at level N from the wavelet decomposition structure $[C,L]$. See `wavedec` for more information on C and L .

Level N must be an integer such that $1 \leq N \leq NMAX$ where $NMAX = \text{length}(L) - 2$.

`D = detcoef(C,L)` extracts the detail coefficients at last level $NMAX$.

If N is a vector of integers such that $1 \leq N(j) \leq NMAX$:

- `DCELL = detcoef(C,L,N,'cells')` returns a cell array where `DCELL{j}` contains the coefficients of detail $N(j)$.
- If $\text{length}(N) > 1$, `DCELL = detcoef(C,L,N)` is equivalent to `DCELL = detcoef(C,L,N,'cells')`.
- `DCELL = detcoef(C,L,'cells')` is equivalent to `DCELL = detcoef(C,L,[1:NMAX])`.
- `[D1, ..., Dp] = detcoef(C,L,[N(1), ..., N(p)])` extracts the details coefficients at levels $[N(1), \dots, N(p)]$.

Examples

Detail Coefficients for 1-D Signal

This example shows how to obtain and plot the detail coefficients for an electrical current signal. This example uses zero-padding (see `dwtmode`).

Load the signal and select the first 3920 samples.

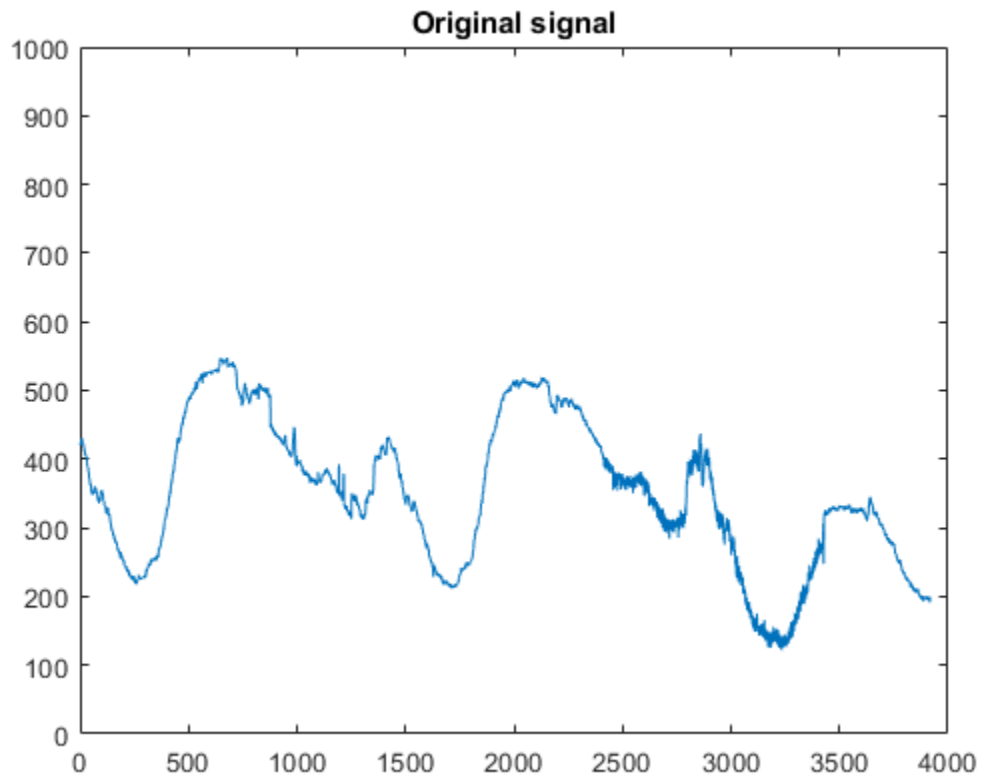
```
load leleccum;  
s = leleccum(1:3920);
```

Perform the decomposition at level 3 using `db1`. Extract the detail coefficients at levels 1, 2, and 3 from the decomposition structure.

```
[c,l] = wavedec(s,3,'db1');  
[cd1,cd2,cd3] = detcoef(c,l,[1 2 3]);
```

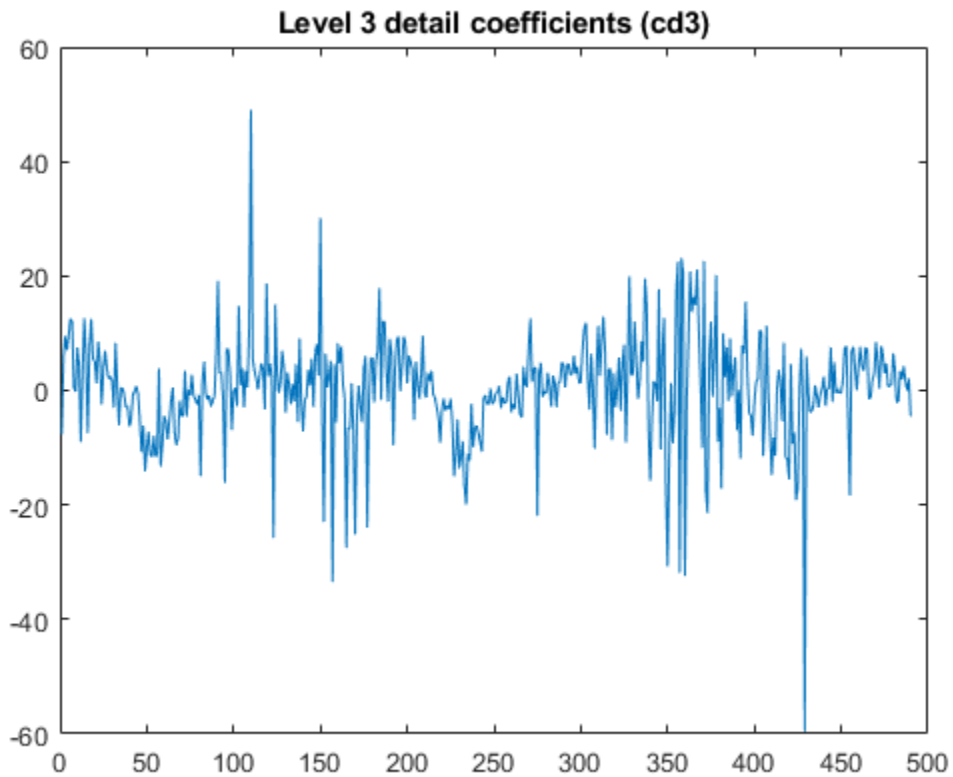
Plot the original signal.

```
plot(s)  
title('Original signal')  
ylim([0 1000])
```

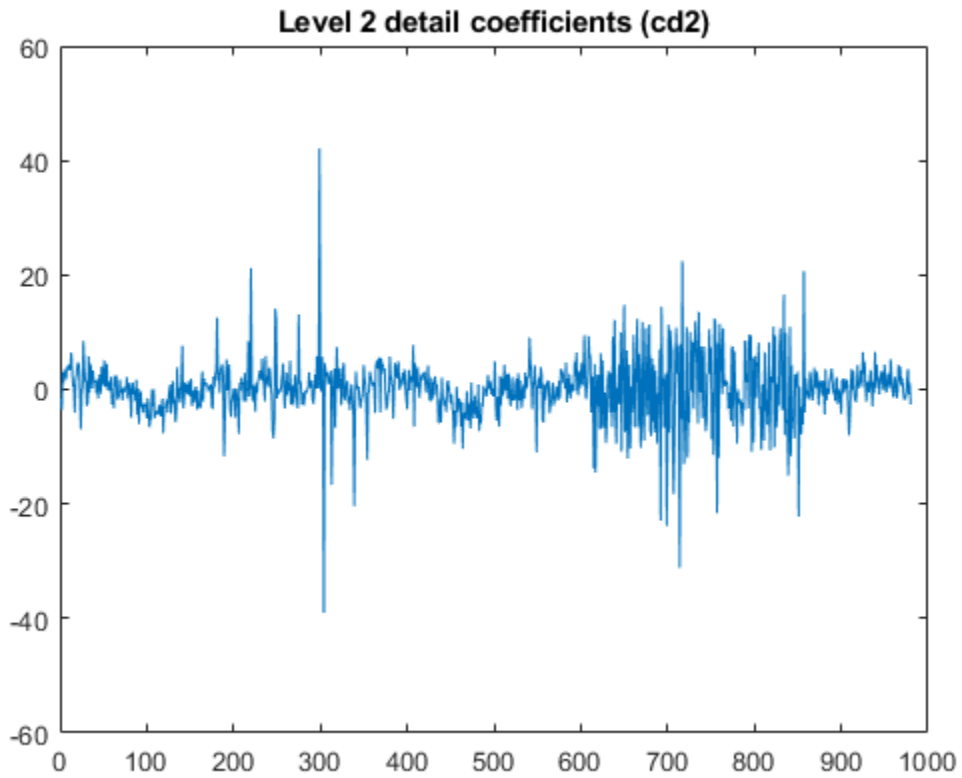
Plot the level 3 detail coefficients.

```
plot(cd3)
title('Level 3 detail coefficients (cd3)')
ylim([-60 60])
```



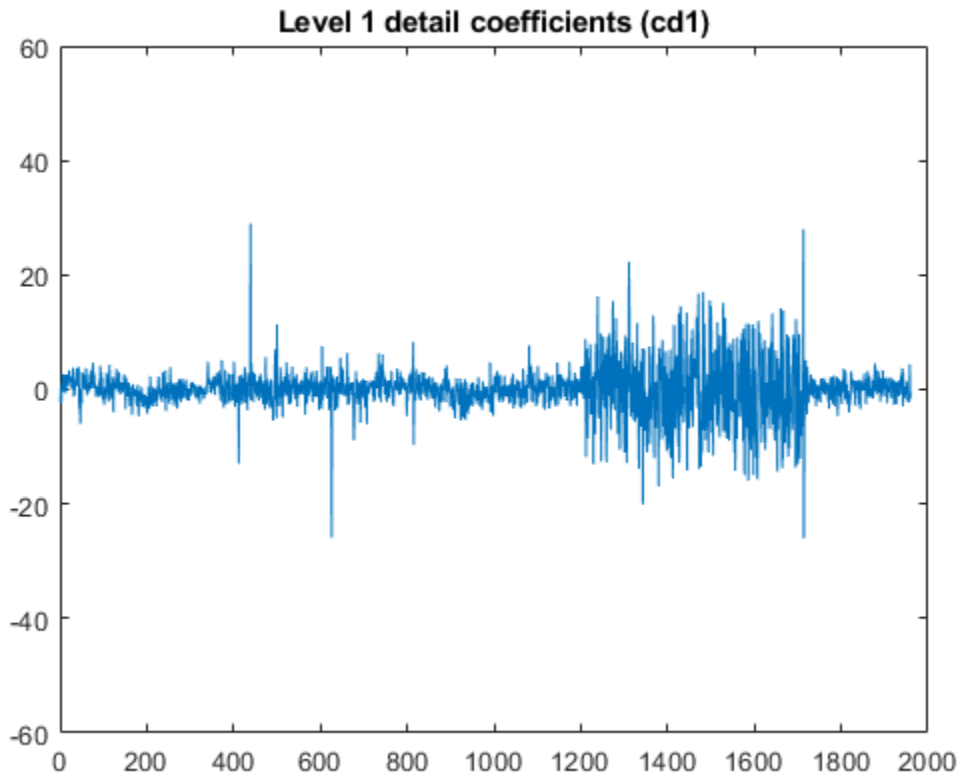
Plot the level 2 detail coefficients.

```
plot (cd2)  
title('Level 2 detail coefficients (cd2)')  
ylim([-60 60])
```



Plot the level 1 detail coefficients.

```
plot (cd1)
title('Level 1 detail coefficients (cd1)')
ylim([-60 60])
```



Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

[appcoef](#) | [wavedec](#)

Introduced before R2006a

detcoef2

2-D detail coefficients

Syntax

```
D = detcoef2(0,C,S,N)
```

Description

detcoef2 is a two-dimensional wavelet analysis function.

`D = detcoef2(0,C,S,N)` extracts from the wavelet decomposition structure `[C,S]` the horizontal, vertical, or diagonal detail coefficients for `0 = 'h'` (or `'v'` or `'d'`, respectively), at level `N`, where `N` must be an integer such that $1 \leq N \leq \text{size}(S,1) - 2$. See `wavedec2` for more information on `C` and `S`.

`[H,V,D] = detcoef2('all',C,S,N)` returns the horizontal `H`, vertical `V`, and diagonal `D` detail coefficients at level `N`.

`D = detcoef2('compact',C,S,N)` returns the detail coefficients at level `N`, stored row-wise.

`detcoef2('a',C,S,N)` is equivalent to `detcoef2('all',C,S,N)`.

`detcoef2('c',C,S,N)` is equivalent to `detcoef2('compact',C,S,N)`.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load original image.  
load woman;
```

```
% X contains the loaded image.
```

```
% Perform decomposition at level 2
% of X using db1.
[c,s] = wavedec2(X,2,'db1');
sizeX = size(X)
sizeX =
    256    256

sizec = size(c)
sizec =
     1   65536

val_s = s
val_s =
    64    64
    64    64
   128   128
   256   256

% Extract details coefficients at level 2
% in each orientation, from wavelet decomposition
% structure [c,s].
[chd2,cvd2,cdd2] = detcoef2('all',c,s,2);
sizecd2 = size(chd2)
sizecd2 =
    64    64

% Extract details coefficients at level 1
% in each orientation, from wavelet decomposition
% structure [c,s].
[chd1,cvd1,cdd1] = detcoef2('all',c,s,1);
sizecd1 = size(chd1)
sizecd1 =
   128   128
```

Tips

If C and S are obtained from an indexed image analysis or a truecolor image analysis, D is an m-by-n matrix or an m-by-n-by-3 array, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

appcoef2 | wavedec2

Introduced before R2006a

disp

WPTREE information

Syntax

`disp(T)`

Description

`disp(T)` displays the content of the WPTREE object *T*.

Examples

```
% Compute a wavelet packets tree
x = rand(1,1000);
t = wpdec(x,2,'db2');
disp(t)
```

Wavelet Packet Object Structure

```
=====
Size of initial data      : [1 1000]
Order                    : 2
Depth                    : 2
Terminal nodes           : [3 4 5 6]
```

```
-----
Wavelet Name              : db2
Low Decomposition filter  : [-0.1294  0.2241  0.8365  0.483]
High Decomposition filter : [ -0.483  0.8365 -0.2241 -0.1294]
Low Reconstruction filter : [  0.483  0.8365  0.2241 -0.1294]
High Reconstruction filter: [-0.1294 -0.2241  0.8365 -0.483]
```

```
-----
Entropy Name              : shannon
Entropy Parameter         : 0
-----
```

See Also

get | read | set | write

Introduced before R2006a

displs

Display lifting scheme

Syntax

`S = displs(LS,FRM)`

Description

`S = displs(LS,FRM)` returns the character array describing the lifting scheme LS. The formatting operator FRM (see `sprintf`) builds S.

`displs(LS)` is equivalent to `displs(LS, '%12.8f')`.

Examples

Display Lifting Scheme

Start with the Haar wavelet and get the corresponding lifting scheme.

```
lshaar = liftwave('haar');
```

Visualized the lifting scheme.

```
displs(lshaar);
```

```
lshaar = {...
'd'          [ -1.00000000] [0]
'p'          [  0.50000000] [0]
[  1.41421356] [  0.70710678] []
};
```

Add a primal elementary lifting step to the lifting scheme. Display the resulting scheme.

```
els = {'p',[-0.125 0.125],0};  
lsnew = addlift(lshaar,els);  
displs(lsnew);  
  
lsnew = {...  
'd'          [ -1.00000000]          [0]  
'p'          [  0.50000000]          [0]  
'p'          [ -0.12500000  0.12500000] [0]  
[  1.41421356] [  0.70710678]          []  
};
```

Input Arguments

LS — Lifting scheme

lifting scheme

Lifting scheme associated with a wavelet. See `liftwave`.

Example: `LS = liftwave('db4')` returns the lifting scheme associated with the Daubechies wavelet `db4`.

FRM — Formatting operator

formatting operator

Formatting operator used to build LS. See `sprintf`.

Example: `'%12.3f'`

See Also

`liftwave` | `lsinfo`

Introduced before R2006a

drawtree

Draw wavelet packet decomposition tree (GUI)

Syntax

```
drawtree(T)  
F = drawtree(T)  
drawtree(T,F)
```

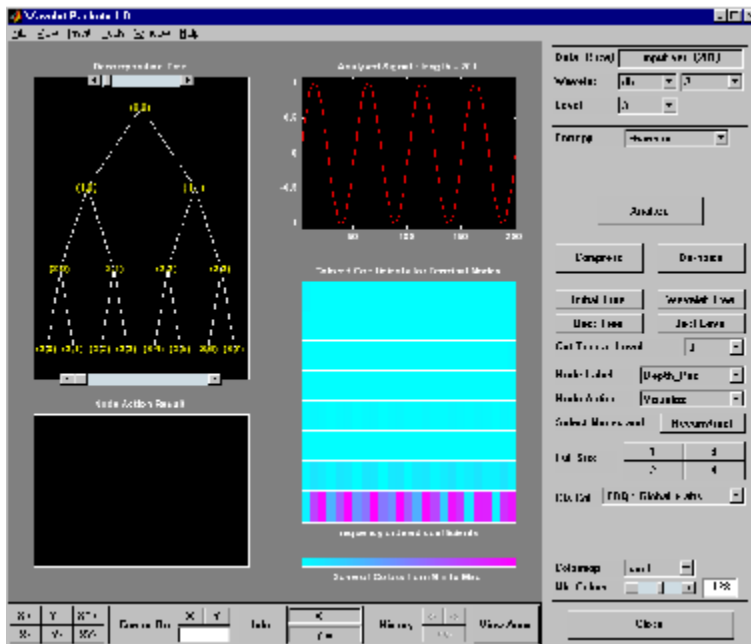
Description

`drawtree(T)` draws the wavelet packet tree *T*, and `F = drawtree(T)` also returns the figure's handle.

For an existing figure *F* produced by a previous call to the `drawtree` function, `drawtree(T,F)` draws the wavelet packet tree *T* in the figure whose handle is *F*.

Examples

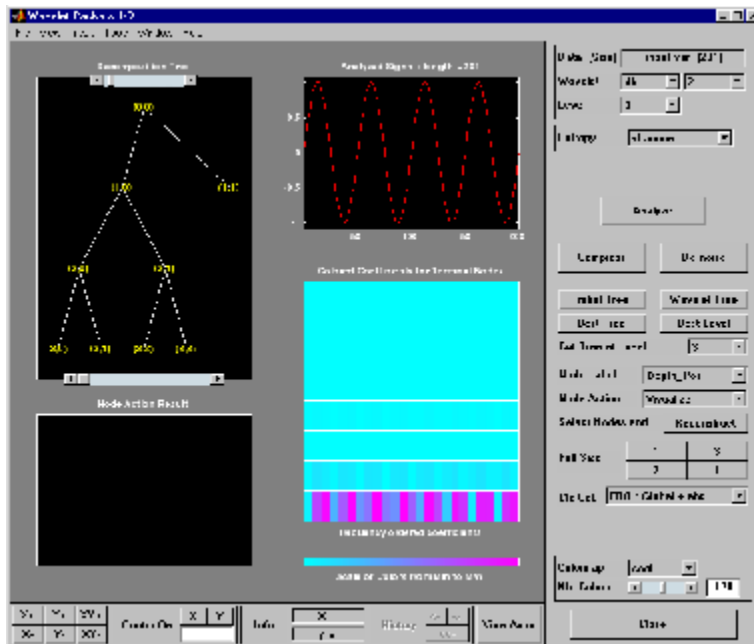
```
x = sin(8*pi*[0:0.005:1]);  
t = wpdec(x,3,'db2');  
fig = drawtree(t);
```



```

%-----
% Use command line function to modify t.
%-----
t = wpjoin(t,2);
drawtree(t,fig);

```



See Also

readtree

Introduced before R2006a

dtfilters

Analysis and synthesis filters for oversampled wavelet filter banks

Syntax

```
df = dtfilters(name)
[df,rf] = dtfilters(name)
```

Description

`df = dtfilters(name)` returns the decomposition (analysis) filters corresponding to `name`. These filters are used most often as input arguments to `dddtree` and `dddtree2`.

`[df,rf] = dtfilters(name)` returns the reconstruction (synthesis) filters corresponding to `name`.

Examples

Filters for Complex Dual-Tree Wavelet Transform

Obtain valid filters for the complex dual-tree wavelet transform. The transform uses Farras nearly symmetric filters for the first stage and Kingsbury Q-shift filters with 10 taps for subsequent stages.

Load the noisy Doppler signal. Obtain the filters for the first and subsequent stages of the complex dual-tree wavelet transform. Demonstrate perfect reconstruction using the complex dual-tree wavelet transform.

```
load noisdopp;
df = dtfilters('dtf2');
dt = dddtree('cplxdt',noisdopp,5,df{1},df{2});
xrec = idddtree(dt);
max(abs(noisdopp-xrec))

ans = 1.3678e-13
```


Filters for Double-Density Wavelet Transform

Obtain valid filters for the double-density wavelet transform.

Load the noisy Doppler signal. Obtain the filters for the double-density wavelet transform. The double-density wavelet transform uses the same filters at all stages. Demonstrate perfect reconstruction using the double-density wavelet transform.

```
df = dtfilters('filters1');
load noisdopp;
dt = dddtree('ddt',noisdopp,5,df,df);
xrec = idddtree(dt);
max(abs(noisdopp-xrec))
```

```
ans = 2.3803e-13
```

Input Arguments

name — Filter name

'dtf1' | 'dddtf1' | 'self1' | 'self2' | ...

Filter name, specified as a character vector or string scalar. Valid entries for name are:

- Any valid orthogonal or biorthogonal wavelet name. See `wfilters` for details. An orthogonal or biorthogonal wavelet is only valid when the filter bank type is `'dwt'`, or when you use the filter as the first stage in a complex dual-tree transform, `'realdt'` or `'cplxdt'`. An orthogonal or biorthogonal wavelet filter is not a valid filter if you have a double-density, `'ddt'` or dual-tree double-density, `'realdddt'` or `'cplxdddt'`, filter bank. An orthogonal or biorthogonal wavelet filter is not a valid filter for complex dual-tree filter banks for stages greater than 1.
- `'dtfP'` — With P equal to 1, 2, 3, 4, or 5 returns the first-stage Farras filters (`'FSfarras'`) and Kingsbury Q-shift filters (`'qshiftN'`) for subsequent stages. This input is only valid for a dual-tree transform, `'realdt'` or `'cplxdt'`. Setting P = 1, 2, 3, 4, or 5 specifies the Kingsbury Q-shift filters with N = 6, 10, 14, 16, or 18 taps, respectively.

- `'dddtf1'` — Returns the filters for the first and subsequent stages of the double-density dual-tree transform. This input is only valid for the double-density dual-tree transforms, `'realdddt'` and `'cplxdddt'`.
- `'self1'` — Returns 10-tap filters for the double-density wavelet transform. This option is only valid for double-density wavelet transforms, `'ddt'`, `'realdddt'`, and `'cplxdddt'`.
- `'self2'` — Returns 16-tap filters for the double-density wavelet transform. This option is only valid for double-density wavelet transforms, `'ddt'`, `'realdddt'`, and `'cplxdddt'`.
- `'filters1'` — Returns 6-tap filters for the double-density wavelet transform, `'ddt'`.
- `'filters2'` — Returns 12-tap filters for the double-density wavelet transform, `'ddt'`.
- `'farras'` — Farras nearly symmetric filters for a two-channel perfect reconstruction filter bank. This option is meant to be used for one-tree transforms and is valid only for an orthogonal critically sampled wavelet transform, `'dwt'`. The output of `dtfilters` is a two-column matrix. The first column of the matrix is a scaling (lowpass) filter, and the second column is a wavelet (highpass) filter.
- `'FSfarras'` — Farras nearly symmetric first-stage filters intended for a dual-tree wavelet transform. With this option, the output of `dtfilters` is a cell array with two elements, one for each tree. Each element is a two-column matrix. The first column of the matrix is a scaling (lowpass) filter, and the second column is a wavelet (highpass) filter.
- `'qshiftN'` — Kingsbury Q-shift N-tap filters with $N = 6, 10, 14, 16, \text{ or } 18$. The Kingsbury Q-shift filters are used most commonly in dual-tree wavelet transforms for stages greater than 1.
- `'doubledualfilt'` — Filters for one stage of the double-density dual-tree wavelet transforms, `'realdddt'` or `'cplxdddt'`.

Output Arguments

df — Decomposition (analysis) filters

matrix | cell array

Decomposition (analysis) filters, returned as a matrix or cell array of matrices.

rf — Reconstruction (synthesis) filters

matrix | cell array

Reconstruction (synthesis) filters, returned as a matrix or cell array of matrices.

See Also

dddtree | dddtree2

Introduced in R2013b

dtree

DTREE constructor

Syntax

```
T = dtree(ORD,D,X)
T = dtree(ORD,D,X,U)
[T,NB] = dtree(...)
[T,NB] = dtree('PropName1',PropValue1,'PropName2',PropValue2,...)
```

Description

`T = dtree(ORD,D,X)` returns a complete data tree (DTREE) object of order *ORD* and depth *D*. The data associated with the tree *T* is *X*.

With `T = dtree(ORD,D,X,U)` you can set a user data field.

`[T,NB] = dtree(...)` returns also the number of terminal nodes (leaves) of *T*.

`[T,NB] = dtree('PropName1',PropValue1,'PropName2',PropValue2,...)` is the most general syntax to construct a DTREE object.

The valid choices for '*PropName*' are

'order'	Order of the tree
'depth'	Depth of the tree
'data'	Data associated to the tree
'spsch'	Split scheme for nodes
'ud'	User data field

The split scheme field is an order *ORD* by 1 logical array. The root of the tree can be split and it has *ORD* children. If `spsch(j) = 1`, you can split the *j*-th child. Each node that you can split has the same property as the root node.

For more information on object fields, type `help dtree/get`.

Class DTREE (Parent class: NTREE)

Fields

dtree	Parent object
allNI	All nodes information
terNI	Terminal nodes information

Examples

```
% Create a data tree.  
x = [1:10];  
t = dtree(3,2,x);  
t = nodejoin(t,2);
```

See Also

ntree | wtbo

Introduced before R2006a

dualtree3

3-D dual-tree complex wavelet transform

Syntax

```
[a,d] = dualtree3(x)
[a,d] = dualtree3(x,level)

[a,d] = dualtree3( ____,Name,Value)

[a,d] = dualtree3( ____, 'excludeL1')
```

Description

`[a,d] = dualtree3(x)` returns the 3-D dual-tree complex wavelet transform of `x` at the maximum level, `floor(log2(min(size(x))))`.

`[a,d] = dualtree3(x,level)` returns the 3-D dual-tree wavelet transform down to `level`.

`[a,d] = dualtree3(____,Name,Value)` specifies options using name-value pair arguments in addition to any of the input arguments in previous syntaxes.

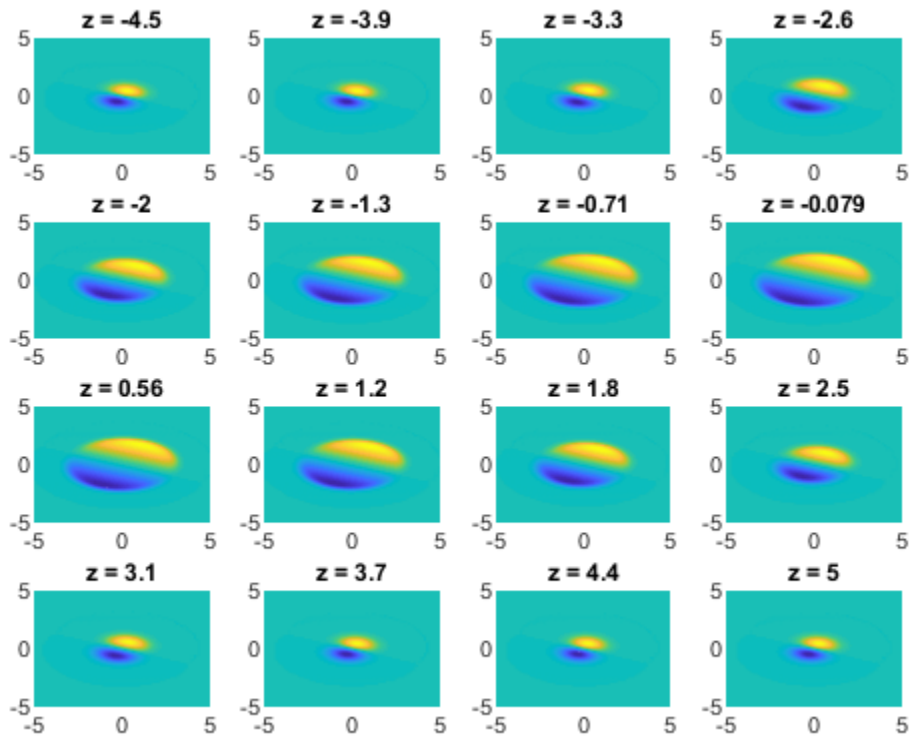
`[a,d] = dualtree3(____, 'excludeL1')` excludes the first-level wavelet (detail) coefficients. Excluding the first-level wavelet coefficients can speed up the algorithm and saves memory. The first level does not exhibit the directional selectivity of levels 2 and higher. The perfect reconstruction property of the dual-tree wavelet transform holds only if the first-level wavelet coefficients are included. If you do not specify this option, or append `'includeL1'`, then the function includes the first-level coefficients.

Examples

Three-Dimensional Dual-Tree Transform of Volumetric Data

Generate a volumetric data set. Plot several cross-sections of the data seen from above. The data are not symmetric about the x-axis or the y-axis.

```
x1 = 64;
xx = linspace(-5,5,x1);
[x,y,z] = meshgrid(xx);
G = (x+3*y)./(1+exp((x.^2+2*y.^2+z.^2)-10));
for k = 1:16
    subplot(4,4,k)
    surf(xx,xx,G(:,:,4*k))
    view(2)
    shading interp
    title(['z = ' num2str(xx(4*k),2)])
end
```



Compute the 3-D dual-tree transform of the data down to level 4. Specify a Hilbert Q-shift filter-pair length of 14.

```
[a,d] = dualtree3(G,4,'FilterLength',14);
```

Plot the real and imaginary parts of the first-level wavelet coefficients for selected subbands. The coefficients have the same directionality as the data. The imaginary parts are shifted versions of the real parts.

```
m = 1;
```

```
for k = 1:8
    subplot(4,4,2*k-1)
    surf(real(d{m}(:,:,3*k)))
```



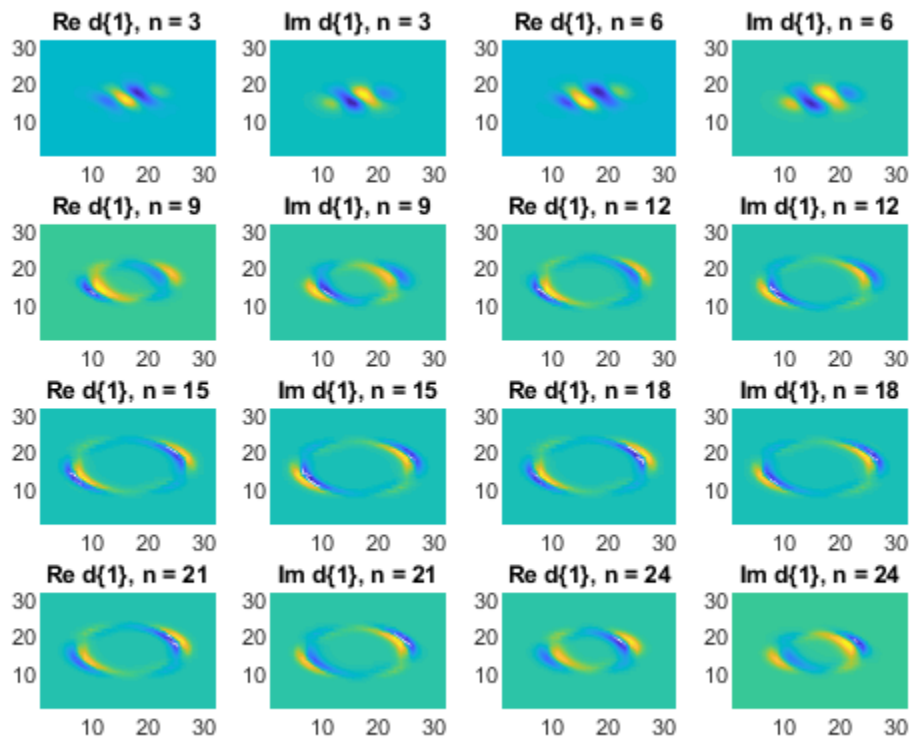
```

view(2)
shading interp
axis tight
title(['Re d\{1\}, n = ' int2str(3*k)])

subplot(4,4,2*k)
surf(imag(d{m}(:, :, 3*k)))

view(2)
shading interp
axis tight
title(['Im d\{1\}, n = ' int2str(3*k)])
end

```



Repeat the procedure for the second-level coefficients. When the level number increases by one, the array of wavelet coefficients decreases by half along the first two dimensions.

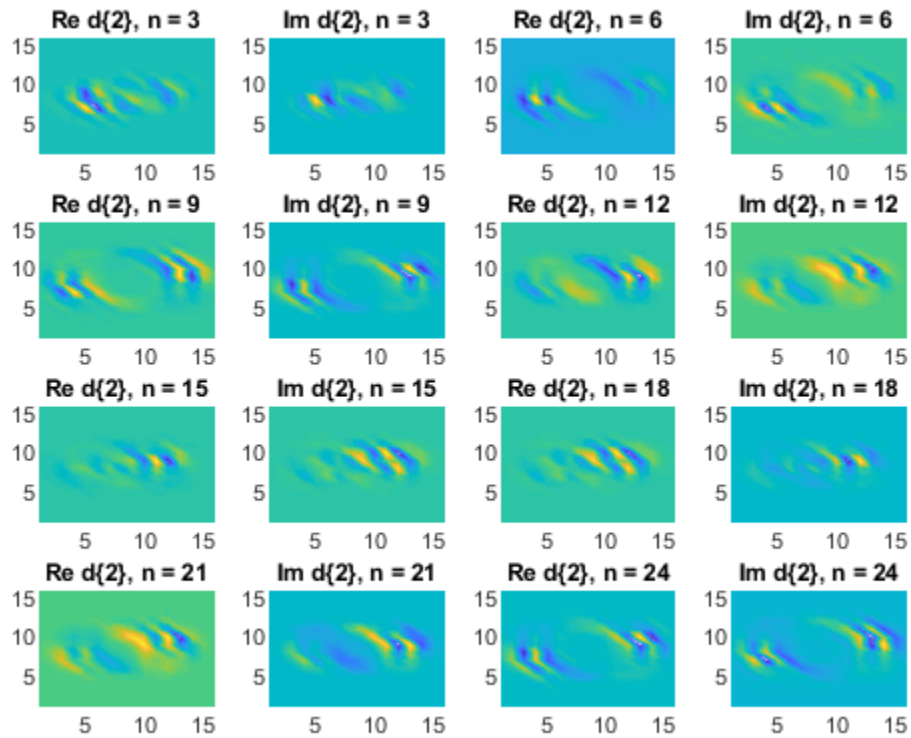
```
m = 2;

for k = 1:8
    subplot(4,4,2*k-1)
    surf(real(d{m}(:,:,3*k)))

    view(2)
    shading interp
    axis tight
    title(['Re d\{2\}, n = ' int2str(3*k)])

    subplot(4,4,2*k)
    surf(imag(d{m}(:,:,3*k)))

    view(2)
    shading interp
    axis tight
    title(['Im d\{2\}, n = ' int2str(3*k)])
end
```



Invert the transform, specifying the same filter-pair length. Check for perfect reconstruction.

```
xrec = idualtree3(a,d,'FilterLength',14);
max(abs(xrec(:)-G(:)))
```

```
ans = 7.1054e-15
```

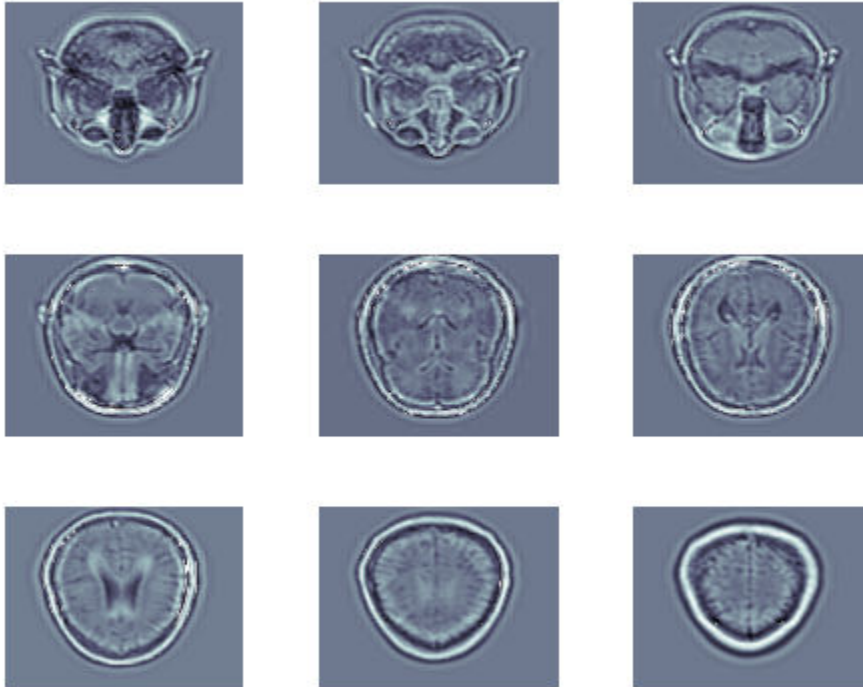
3-D Dual-Tree Transform of MRI Data

Load a set of MRI measurements of a human head. Truncate the data so that it is even along the third dimension. Compute the 3-D dual-tree transform, excluding the first-level wavelet coefficients.

```
load wmri  
  
[A,D] = dualtree3(X(:,:,1:26),2,'excludel1');
```

Reconstruct the data by inverting the transform. Set the final-level scaling coefficients explicitly to 0. Display an evenly spaced selection of reconstructed images.

```
imrec = idualtree3(A*0,D);  
  
colormap bone  
for kj = 1:9  
    subplot(3,3,kj)  
    surf(imrec(:,:,3*kj-2))  
    shading interp  
    view(2)  
    axis tight off  
end
```



Input Arguments

x — Input data

real 3-D array

Input data, specified as a real 3-D array. All three dimensions of **x** must be even and greater than or equal to 4.

Data Types: `double` | `single`

level — Transform level

$\text{floor}(\log_2(\min(\text{size}(x))))$ (default) | positive integer

Transform level, specified as a positive integer greater than or equal to 2 and less than or equal to $\text{floor}(\log_2(\min(\text{size}(x))))$.

Data Types: `double` | `single`

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'LevelOneFilter', 'legall', 'FilterLength', 6` computes a transform using LeGall analysis filters with scaling length 5 and wavelet length 3 at level 1, and length-6 Q-shift filters at levels 2 and greater.

FilterLength — Hilbert Q-shift filter-pair length

10 (default) | 6 | 14 | 16 | 18

Hilbert Q-shift filter-pair length, specified as the comma-separated pair consisting of `'FilterLength'` and one of 6, 10, 14, 16, or 18. `dualtree3` uses the orthogonal Hilbert Q-shift filter pair of length `'FilterLength'` for levels 2 and greater.

Data Types: `double` | `single`

LevelOneFilter — First-level biorthogonal analysis filter

`'nearsym5_7'` (default) | `'nearsym13_19'` | `'antonini'` | `'legall'`

First-level biorthogonal analysis filter, specified as the comma-separated pair consisting of `'LevelOneFilter'` and a character vector or string. By default, `dualtree3` uses for level 1 the near-symmetric biorthogonal wavelet filter with lengths 5 (scaling filter) and 7 (wavelet filter).

Data Types: `char` | `string`

Output Arguments

a — Final-level scaling coefficients

real-valued matrix

Final-level scaling (lowpass) coefficients, returned as a real-valued matrix.

Data Types: double

d — Wavelet coefficients

1-by-level cell array

Wavelet coefficients, returned as a 1-by-level cell array. There are 28 wavelet subbands in the 3-D dual-tree transform at each level.

Data Types: double

References

- [1] Chen, H., and N. G. Kingsbury. "Efficient Registration of Nonrigid 3-D Bodies." *IEEE® Transactions on Image Processing*. Vol 21, January 2012, pp. 262-272.
- [2] Kingsbury, N. G. "Complex Wavelets for Shift Invariant Analysis and Filtering of Signals." *Journal of Applied and Computational Harmonic Analysis*. Vol. 10, May 2001, pp. 234-253.

See Also

dddtree2 | idualtree3 | wavedec3 | waverec3

Topics

Dual-Tree Wavelet Transforms

Introduced in R2017a

dwt

Single-level 1-D discrete wavelet transform

Syntax

```
[cA, cD] = dwt(x, wname)  
[cA, cD] = dwt(x, LoD, HiD)  
[cA, cD] = dwt( ___, 'mode', extmode)
```

Description

`[cA, cD] = dwt(x, wname)` returns the single-level discrete wavelet transform (DWT) of the vector `x` using the wavelet specified by `wname`. The wavelet must be recognized by `wavemngr`. `dwt` returns the approximation coefficients vector `cA` and detail coefficients vector `cD` of the DWT.

Note If your application requires a multilevel wavelet decomposition, consider using `wavedec`.

`[cA, cD] = dwt(x, LoD, HiD)` returns the single-level DWT using the scaling (lowpass) filter `LoD` and wavelet (highpass) filter `HiD`. The filters must have the same even length.

`[cA, cD] = dwt(___, 'mode', extmode)` returns the single-level DWT with the specified extension mode `extmode`. For more information, see `dwtmode`.

Examples

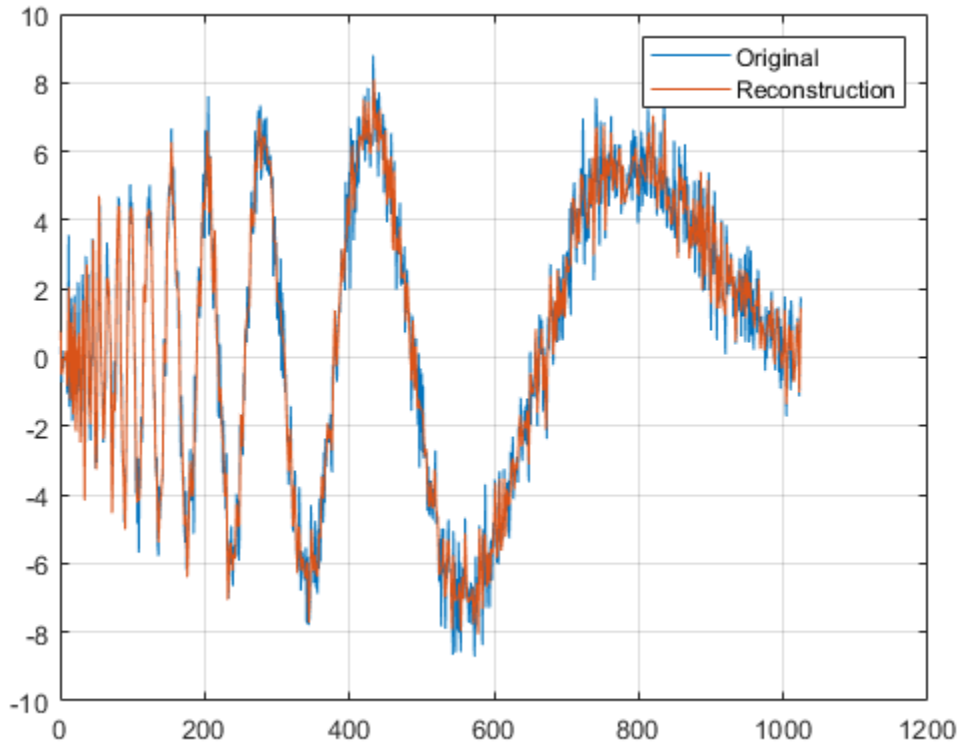
DWT Using Wavelet Name

Obtain the single-level DWT of the noisy Doppler signal using a wavelet name.


```
load noisdopp;  
[cA,cD] = dwt(noisdopp,'sym4');
```

Reconstruct a smoothed version of the signal using the approximation coefficients. Plot and compare with the original signal.

```
xrec = idwt(cA,zeros(size(cA)),'sym4');  
plot(noisdopp)  
hold on  
grid on  
plot(xrec)  
legend('Original','Reconstruction')
```



DWT Using Wavelet and Scaling Filters

Obtain the single-level DWT of a noisy Doppler signal using the wavelet (highpass) and scaling (lowpass) filters.

```
load noisdopp;  
[LoD,HiD] = wfilters('bior3.5','d');  
[cA,cD] = dwt(noisdopp,LoD,HiD);
```

Create a DWT filter bank that can be applied to the noisy Doppler signal using the same wavelet. Obtain the highpass and lowpass filters from the filter bank.

```
len = length(noisdopp);  
fb = dwtfilterbank('SignalLength',len,'Wavelet','bior3.5');  
[lo,hi] = filters(fb);
```

For the `bior3.5` wavelet, `lo` and `hi` are 12-by-2 matrices. `lo` are the lowpass filters, and `hi` are the highpass filters. The first columns of `lo` and `hi` are used for analysis and the second columns are used for synthesis. Compare the first column of `lo` and `hi` with `LoD` and `HiD` respectively. Confirm they are equal.

```
disp('Lowpass Analysis Filters')
```

```
Lowpass Analysis Filters
```

```
[lo(:,1) LoD']
```

```
ans = 12×2
```

```
-0.0138    -0.0138  
 0.0414     0.0414  
 0.0525     0.0525  
-0.2679    -0.2679  
-0.0718    -0.0718  
 0.9667     0.9667  
 0.9667     0.9667  
-0.0718    -0.0718  
-0.2679    -0.2679  
 0.0525     0.0525  
  :
```

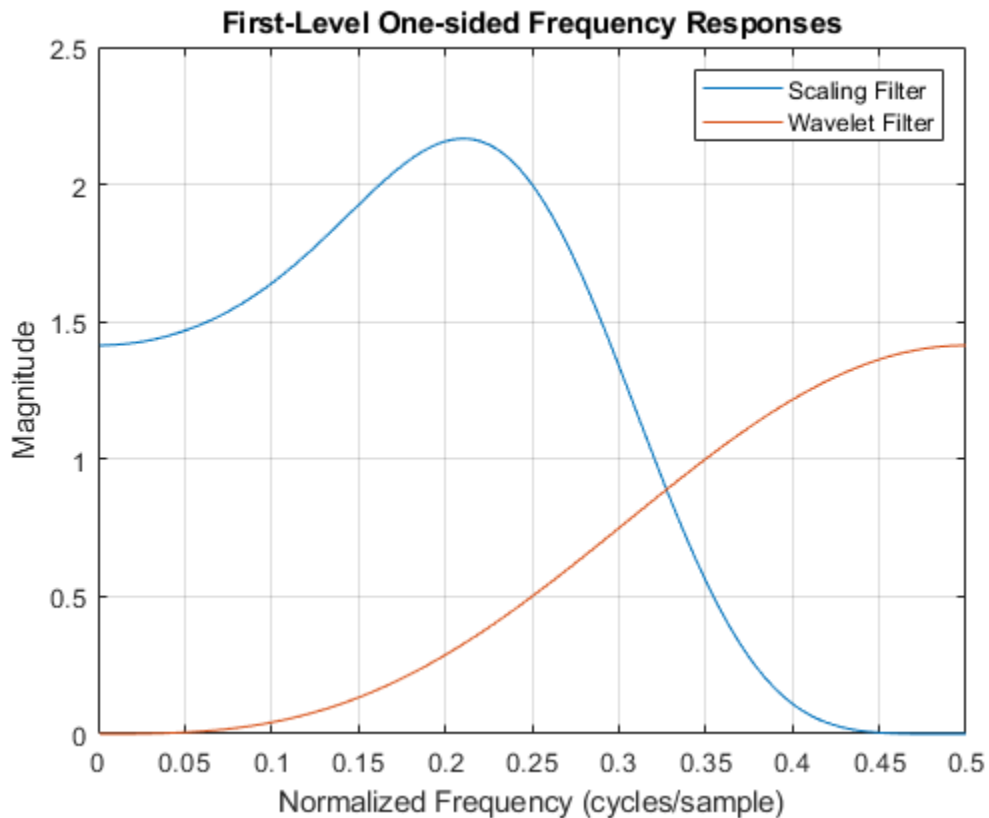
```
disp('Highpass Analysis Filters')
```

```
Highpass Analysis Filters
```

```
[hi(:,1) HiD']
ans = 12x2
    0         0
    0         0
    0         0
    0         0
   -0.1768   -0.1768
    0.5303    0.5303
   -0.5303   -0.5303
    0.1768    0.1768
    0         0
    0         0
    :
```

Plot the one-sided magnitude frequency responses of the first-level wavelet and scaling filters.

```
[psidft,f,phidft] = freqz(fb);
level = 1;
plot(f(len/2+1:end),abs(phidft(level,len/2+1:end)))
hold on
plot(f(len/2+1:end),abs(psidft(level,len/2+1:end)))
grid on
legend('Scaling Filter','Wavelet Filter')
title('First-Level One-sided Frequency Responses')
xlabel('Normalized Frequency (cycles/sample)')
ylabel('Magnitude')
```



Input Arguments

x — Input data

real-valued vector

Input data, specified as a real-valued vector.

Data Types: double

wname — Analyzing wavelet

character vector | string scalar

Analyzing wavelet used to compute the single-level DWT, specified as a character vector or string scalar. The wavelet must be recognized by `wavemngr`. The analyzing wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wfilters` for the wavelets available in each family.

Example: `'db4'`

LoD — Scaling filter

even-length real-valued vector

Scaling (lowpass) filter, specified as an even-length real-valued vector. LoD must be the same length as HiD. See `wfilters` for details.

Data Types: `double`

HiD — Wavelet filter

even-length real-valued vector

Wavelet (highpass) filter, specified as an even-length real-valued vector. HiD must be the same length as LoD. See `wfilters` for details.

Data Types: `double`

extmode — Extension mode

`'zpd'` | `'sp0'` | `'spd'` | ...

Extension mode used when performing the DWT, specified as one of the following:

mode	DWT Extension Mode
<code>'zpd'</code>	Zero extension
<code>'sp0'</code>	Smooth extension of order 0
<code>'spd'</code> (or <code>'sp1'</code>)	Smooth extension of order 1
<code>'sym'</code> or <code>'symh'</code>	Symmetric extension (half point): boundary value symmetric replication
<code>'symw'</code>	Symmetric extension (whole point): boundary value symmetric replication
<code>'asym'</code> or <code>'asymh'</code>	Antisymmetric extension (half point): boundary value antisymmetric replication

mode	DWT Extension Mode
'asymw'	Antisymmetric extension (whole point): boundary value antisymmetric replication
'ppd'	Periodized extension (1)
'per'	Periodized extension (2) If the signal length is odd, <code>wextend</code> adds to the right an extra sample that is equal to the last value, and performs the extension using the 'ppd' mode. Otherwise, 'per' reduces to 'ppd'. This rule also applies to images.

The global variable managed by `dwtmode` specifies the default extension mode. See `dwtmode` for extension mode descriptions.

Example: `[cA,cD] = dwt(x,'db4','mode','symw')` returns the single-level DWT of `x` using the order 4 Daubechies extremal phase wavelet and whole point symmetric extension.

Output Arguments

cA — Approximation coefficients

real-valued vector

Approximation coefficients obtained from the wavelet decomposition, returned as a vector. Convolving the input signal `x` with the scaling filter `LoD`, followed by dyadic decimation, yields the approximation coefficients. Let `sx = size(x)` and `lf` = the length of the decomposition filters.

- If the DWT extension mode is set to periodization, `cA` is a vector of length `ceil(sx/2)`.
- For the other extension modes, `cA` is a vector of length `floor((sx+lf-1)/2)`.

Data Types: `double`

cD — Detail coefficients

real-valued vector

Detail coefficients obtained from the wavelet decomposition, returned as a vector. Convolving the input signal `x` with the wavelet filter `HiD`, followed by dyadic decimation,

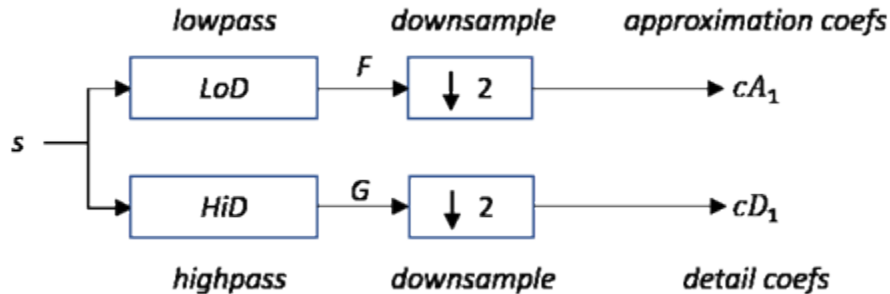
yields the detail coefficients. Let $sx = \text{size}(x)$ and $lf =$ the length of the decomposition filters.

- If the DWT extension mode is set to periodization, cD is a vector of length $\text{ceil}(sx/2)$.
- For the other extension modes, cD is a vector of length $\text{floor}((sx+lf-1)/2)$.

Data Types: `double`

Algorithms

Starting from a signal s of length N , two sets of coefficients are computed: approximation coefficients cA_1 , and detail coefficients cD_1 . Convolution of s with the scaling filter LoD , followed by dyadic decimation, yields the approximation coefficients. Similarly, convolution of s with the wavelet filter HiD , followed by dyadic decimation, yields the detail coefficients.



where

- \boxed{X} — Convolve with filter X
- $\boxed{\downarrow 2}$ — Downsample (keep the even-indexed elements)

The length of each filter is equal to $2n$. If $N = \text{length}(s)$, the signals F and G are of length $N + 2n - 1$ and the coefficients cA_1 and cD_1 are of length $\text{floor}\left(\frac{N-1}{2}\right) + n$.

To deal with signal-end effects resulting from a convolution-based algorithm, a global variable managed by `dwtmode` defines the kind of signal extension mode used. The possible options include zero-padding and symmetric extension, which is the default mode.

Note For the same input, this `dwt` function and the DWT block in the DSP System Toolbox™ do not produce the same results. The DWT block is designed for real-time implementation while Wavelet Toolbox software is designed for analysis, so the products handle boundary conditions and filter states differently.

To make the `dwt` function output match the DWT block output, set the function boundary condition to zero-padding by typing `dwtmode('zpd')` at the MATLAB command prompt. To match the latency of the DWT block, which is implemented using FIR filters, add zeros to the input of the `dwt` function. The number of zeros you add must be equal to half the filter length.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*. CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: Society for Industrial and Applied Mathematics, 1992.
- [2] Mallat, S. G. "A Theory for Multiresolution Signal Decomposition: The Wavelet Representation." *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 11, Issue 7, July 1989, pp. 674-693.
- [3] Meyer, Y. *Wavelets and Operators*. Translated by D. H. Salinger. Cambridge, UK: Cambridge University Press, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`dwtfilterbank` | `dwtmode` | `idwt` | `wavedec` | `waveinfo`

Introduced before R2006a

dwt2

Single-level discrete 2-D wavelet transform

`dwt2` computes the single-level 2-D wavelet decomposition. Compare `dwt2` with `wavedec2` which may be more useful for your application. The decomposition is done with respect to either a particular wavelet (see `wfilters` for more information) or particular wavelet decomposition filters.

Syntax

```
[cA, cH, cV, cD] = dwt2(X, wname)
[cA, cH, cV, cD] = dwt2(X, LoD, HiD)
[cA, cH, cV, cD] = dwt2( ____, 'mode', extmode)
```

Description

`[cA, cH, cV, cD] = dwt2(X, wname)` computes the single-level 2-D discrete wavelet transform (DWT) of the input data `X` using the `wname` wavelet. `dwt2` returns the approximation coefficients matrix `cA` and detail coefficients matrices `cH`, `cV`, and `cD` (horizontal, vertical, and diagonal, respectively).

`[cA, cH, cV, cD] = dwt2(X, LoD, HiD)` computes the single-level 2-D DWT using the wavelet decomposition lowpass filter `LoD` and highpass filter `HiD`. The decomposition filters must have the same length.

`[cA, cH, cV, cD] = dwt2(____, 'mode', extmode)` computes the single-level 2-D DWT with the extension mode `extmode`. Include this argument after all other arguments.

Examples

Single-Level 2-D Discrete Wavelet Transform of an Image

Load and display an image.

```
load woman
imagesc(X)
colormap(map)
```

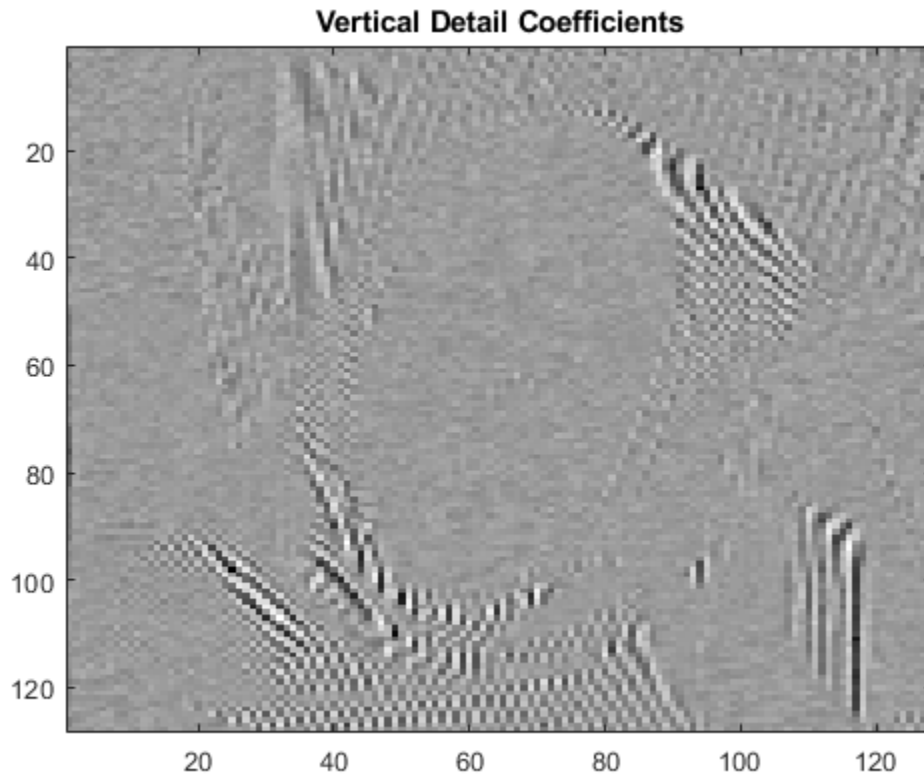


Obtain the single-level 2-D discrete wavelet transform of the image using the order 4 symlet and periodic extension.

```
[cA,cH,cV,cD] = dwt2(X, 'sym4', 'mode', 'per');
```

Display the vertical detail coefficients and the approximation coefficients.

```
imagesc(cV)
title('Vertical Detail Coefficients')
```



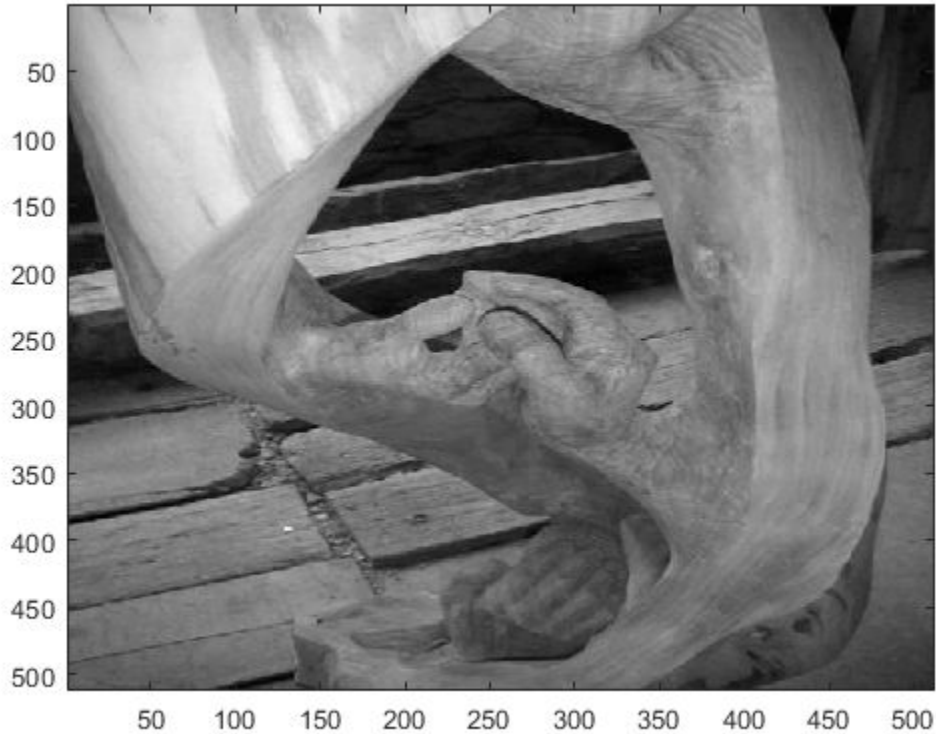
```
imagesc(cA)  
title('Approximation Coefficients')
```



Single-Level 2-D Discrete Wavelet Transform Using Filters

Load and display an image.

```
load sculpture
imagesc(X)
colormap gray
```



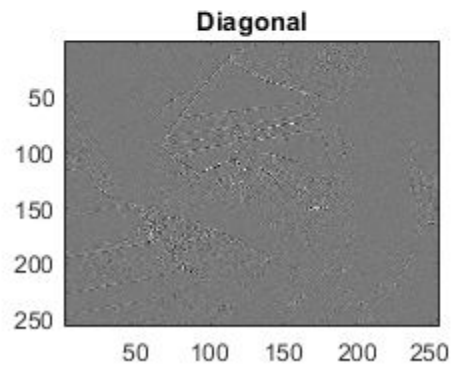
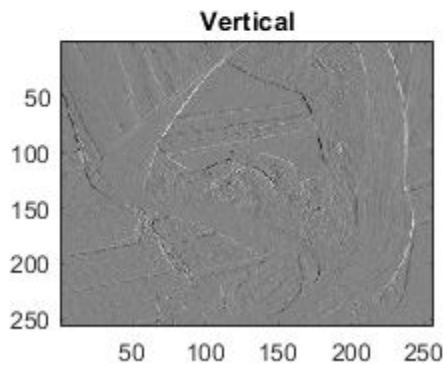
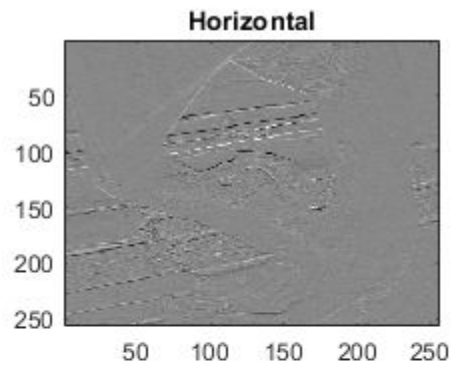
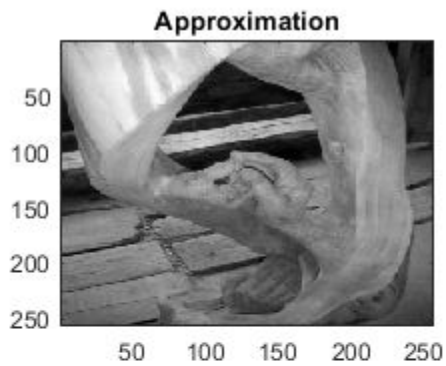
Generate the lowpass and highpass decomposition filters for the Haar wavelet.

```
[LoD,HiD] = wfilters('haar','d');
```

Use the filters to perform a single-level 2-D wavelet decomposition. Use half-point symmetric extension. Display the approximation and detail coefficients.

```
[cA,cH,cV,cD] = dwt2(X,LoD,HiD,'mode','symh');  
subplot(2,2,1)  
imagesc(cA)  
colormap gray  
title('Approximation')  
subplot(2,2,2)  
imagesc(cH)
```

```
colormap gray
title('Horizontal')
subplot(2,2,3)
imagesc(cV)
colormap gray
title('Vertical')
subplot(2,2,4)
imagesc(cD)
colormap gray
title('Diagonal')
```



Input Arguments

X — Input data

matrix | 3-D array of RGB triplets

Input data, specified as an *m*-by-*n* matrix representing an indexed image or an *m*-by-*n*-by-3 array representing a truecolor image. For more information on truecolor images, see “RGB (Truecolor) Images” (MATLAB).

Data Types: double

wname — Analyzing wavelet

character vector | string scalar

Analyzing wavelet used to compute the 2-D DWT, specified as a character vector or string scalar. The analyzing wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wfilters` for the wavelets available in each family.

LoD — Wavelet decomposition lowpass filter

even-length real-valued vector

Wavelet decomposition lowpass filter, specified as an even-length real-valued vector. `LoD` must be of the same length as `HiD`.

HiD — Wavelet decomposition highpass filter

even-length real-valued vector

Wavelet decomposition highpass filter, specified as an even-length real-valued vector. `HiD` must be of the same length as `LoD`.

extmode — Extension mode

'zpd' | 'sp0' | 'spd' | ...

Extension mode used when performing the DWT, specified as one of the following:

mode	DWT Extension Mode
'zpd'	Zero extension
'sp0'	Smooth extension of order 0
'spd' (or 'sp1')	Smooth extension of order 1

mode	DWT Extension Mode
'sym' or 'symh'	Symmetric extension (half point): boundary value symmetric replication
'symw'	Symmetric extension (whole point): boundary value symmetric replication
'asym' or 'asymh'	Antisymmetric extension (half point): boundary value antisymmetric replication
'asymw'	Antisymmetric extension (whole point): boundary value antisymmetric replication
'ppd'	Periodized extension (1)
'per'	Periodized extension (2) If the signal length is odd, <code>wextend</code> adds to the right an extra sample that is equal to the last value, and performs the extension using the 'ppd' mode. Otherwise, 'per' reduces to 'ppd'. This rule also applies to images.

The global variable managed by `dwtmode` specifies the default extension mode.

Example: `[cA,cH,cV,cD] = dwt2(x,'db4','mode','symw');`

Output Arguments

cA — Approximation coefficients

array

Approximation coefficients, returned as an array whose size depends on `X`. Let `sx = size(X)` and `lf =` the length of the decomposition filters.

- If the DWT extension mode is set to periodization, then this output is of size `ceil(sx/2)`.
- For the other extension modes, this output is of size `floor((sx+lf-1)/2)`.

Data Types: `double`

cH — Horizontal detail coefficients

array

Horizontal detail coefficients, returned as an array whose size depends on X . Let $sx = \text{size}(X)$ and $lf =$ the length of the decomposition filters.

- If the DWT extension mode is set to periodization, then this output is of size $\text{ceil}(sx/2)$.
- For the other extension modes, this output is of size $\text{floor}((sx+lf-1)/2)$.

Data Types: `double`

cV — Vertical detail coefficients

array

Vertical detail coefficients, returned as an array whose size depends on X . Let $sx = \text{size}(X)$ and $lf =$ the length of the decomposition filters.

- If the DWT extension mode is set to periodization, then this output is of size $\text{ceil}(sx/2)$.
- For the other extension modes, this output is of size $\text{floor}((sx+lf-1)/2)$.

Data Types: `double`

cD — Diagonal detail coefficients

array

Diagonal detail coefficients, returned as an array whose size depends on X . Let $sx = \text{size}(X)$ and $lf =$ the length of the decomposition filters.

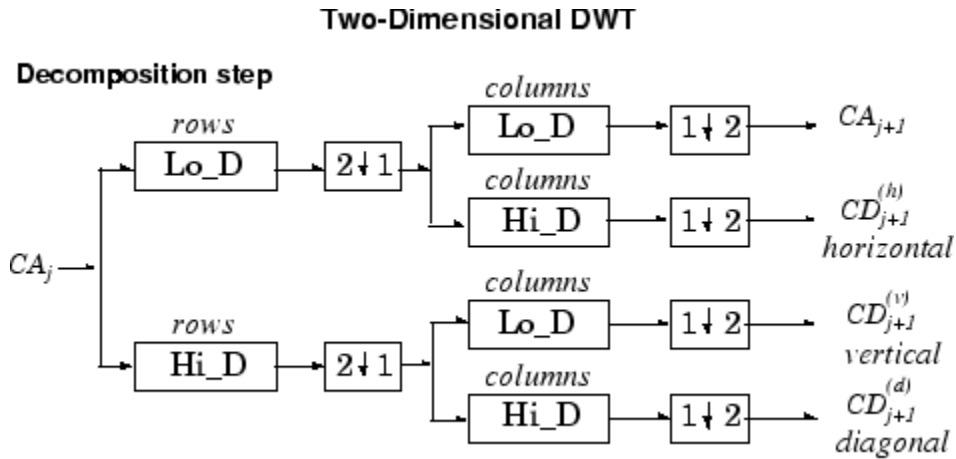
- If the DWT extension mode is set to periodization, then this output is of size $\text{ceil}(sx/2)$.
- For the other extension modes, this output is of size $\text{floor}((sx+lf-1)/2)$.

Data Types: `double`

Algorithms

The 2-D wavelet decomposition algorithm for images is similar to the one-dimensional case. The two-dimensional wavelet and scaling functions are obtained by taking the tensor products of the one-dimensional wavelet and scaling functions. This kind of two-dimensional DWT leads to a decomposition of approximation coefficients at level j in four components: the approximation at level $j + 1$, and the details in three orientations

(horizontal, vertical, and diagonal). The following chart describes the basic decomposition steps for images.



- $\boxed{2 \downarrow 1}$ — Downsample columns: keep the even indexed columns
- $\boxed{1 \downarrow 2}$ — Downsample rows: keep the even indexed rows
- $\begin{matrix} \text{rows} \\ \boxed{X} \end{matrix}$ — Convolve with filter X the rows of the entry
- $\begin{matrix} \text{columns} \\ \boxed{X} \end{matrix}$ — Convolve with filter X the columns of the entry

The decomposition is initialized by setting the approximation coefficients equal to the image s : $CA_0 = s$.

Note To deal with signal-end effects introduced by a convolution-based algorithm, the 1-D and 2-D DWT use a global variable managed by `dwtmode`. This variable defines the kind of signal extension mode used. The possible options include zero-padding and symmetric extension, which is the default mode.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: SIAM Ed, 1992.
- [2] Mallat, S. G. "A Theory for Multiresolution Signal Decomposition: The Wavelet Representation," *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 11, Issue 7, July 1989, pp. 674-693.
- [3] Meyer, Y. *Wavelets and Operators*. Translated by D. H. Salinger. Cambridge, UK: Cambridge University Press, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`dwtmode` | `haart2` | `idwt2` | `ihaart2` | `wavedec2` | `waveinfo` | `waverec2` | `wfilters`

Introduced before R2006a

dwt3

Single-level discrete 3-D wavelet transform

Syntax

```
wt = dwt3(x,wname)
wt = dwt3(x,wname,'mode',extM)
wt = dwt3(x,w, ___)
wt = dwt3(x,wf, ___)
```

Description

`wt = dwt3(x,wname)` returns the single-level three-dimensional wavelet decomposition `wt` of the input data `x` using the `wname` wavelet. The default extension mode of the 3-D discrete wavelet transform (DWT) is 'sym' (see `dwtmode`).

`wt = dwt3(x,wname,'mode',extM)` uses the extension mode `extM` (see `dwtmode`).

`wt = dwt3(x,w, ___)` specifies three wavelets, one for each direction. `w` is a cell array, string array, or structure, and can be followed by 'mode', `extM`.

`wt = dwt3(x,wf, ___)` specifies four filters, two for decomposition and two for reconstruction, or 3×4 filters (one quadruplet by direction). `wf` is a cell array or structure, and can be followed by 'mode', `extM`.

Examples

Single-Level Three-Dimensional Wavelet Decomposition

Define the original 3-D data.

```
X = reshape(1:64,4,4,4)
X =
X(:,:,1) =
```

```
1    5    9   13
2    6   10   14
3    7   11   15
4    8   12   16
```

```
X(:, :, 2) =
```

```
17   21   25   29
18   22   26   30
19   23   27   31
20   24   28   32
```

```
X(:, :, 3) =
```

```
33   37   41   45
34   38   42   46
35   39   43   47
36   40   44   48
```

```
X(:, :, 4) =
```

```
49   53   57   61
50   54   58   62
51   55   59   63
52   56   60   64
```

Perform single-level decomposition of X using 'db1'.

```
wt = dwt3(X, 'db1')
```

```
wt = struct with fields:
  sizeINI: [4 4 4]
  filters: [1x1 struct]
  mode: 'sym'
  dec: {2x2x2 cell}
```

Decompose X using 'db2'.

```
[LoD,HiD,LoR,HiR] = wfilters('db2');
wt = dwt3(X,{LoD,HiD,LoR,HiR})
```

```
wt = struct with fields:
  sizeINI: [4 4 4]
  filters: [1x1 struct]
  mode: 'sym'
  dec: {2x2x2 cell}
```

Decompose X using different wavelets, one for each orientation: 'db1', 'db2', and again 'db1'.

```
WS = struct('w1','db1','w2','db2','w3','db1');
wt = dwt3(X,WS,'mode','per')
```

```
wt = struct with fields:
  sizeINI: [4 4 4]
  filters: [1x1 struct]
  mode: 'per'
  dec: {2x2x2 cell}
```

Decompose X using the filters given by WF and set the extension mode to symmetric.

```
WF = wt.filters;
wtBIS = dwt3(X,WF,'mode','sym')
```

```
wtBIS = struct with fields:
  sizeINI: [4 4 4]
  filters: [1x1 struct]
  mode: 'sym'
  dec: {2x2x2 cell}
```

Input Arguments

x — Input data

3-D array

Input data, specified as a 3-D array.

Data Types: double

wname — Analyzing wavelet

character vector | string scalar

Analyzing wavelet used to compute the 2-D DWT, specified as a character vector or string scalar. The analyzing wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wfilters` for the wavelets available in each family.

w — Analyzing wavelets

cell array of character vectors | string array | structure

Analyzing wavelets to use in the 3-D wavelet decomposition, one for each direction, specified as a cell array of character vectors, a string array, or a structure. `w = {'wname1', 'wname2', 'wname3'}`, or `w = ["wname1", "wname2", "wname3"]`, or `w` is a structure with 3 fields 'w1', 'w2', 'w3' containing character vectors or string scalars that are the names of wavelets.

Example: `wt = dwt3(x, ["db2", "db4", "db6"]);`

wf — Wavelet filters

cell array | structure

Wavelet filters to use in the 3-D wavelet decomposition, specified as either a cell array or structure. `wf` specifies four filters, two for decomposition and two for reconstruction, or 3 × 4 filters (one quadruplet by direction). `wf` is either a cell array (1 × 4) or (3 × 4) : {LoD, HiD, LoR, HiR} or a structure with the four fields 'LoD', 'HiD', 'LoR', 'HiR'.

extM — Extension mode

'zpd' | 'sp0' | 'spd' | ...

Extension mode used when performing the 3-D DWT, specified as one of the following:

mode	DWT Extension Mode
'zpd'	Zero extension
'sp0'	Smooth extension of order 0
'spd' (or 'sp1')	Smooth extension of order 1
'sym' or 'symh'	Symmetric extension (half point): boundary value symmetric replication

mode	DWT Extension Mode
'symw'	Symmetric extension (whole point): boundary value symmetric replication
'asym' or 'asymh'	Antisymmetric extension (half point): boundary value antisymmetric replication
'asymw'	Antisymmetric extension (whole point): boundary value antisymmetric replication
'ppd'	Periodized extension (1)
'per'	Periodized extension (2) If the signal length is odd, <code>wextend</code> adds to the right an extra sample that is equal to the last value, and performs the extension using the 'ppd' mode. Otherwise, 'per' reduces to 'ppd'. This rule also applies to images.

The global variable managed by `dwtmode` specifies the default extension mode.

Output Arguments

wt — Single-level 3-D wavelet decomposition

structure

Single-level 3-D wavelet decomposition, returned as a structure with the following fields:

<code>sizeINI</code>	Size of the three-dimensional array <code>X</code> .
<code>mode</code>	Name of the wavelet transform extension mode.
<code>filters</code>	Structure with four fields: <code>LoD</code> , <code>HiD</code> , <code>LoR</code> , <code>HiR</code> , which are the filters used for DWT.

dec	<p>$2 \times 2 \times 2$ cell array containing the coefficients of the decomposition.</p> <p><code>dec{i,j,k}</code>, $i,j,k = 1$ or 2 contains the coefficients obtained by lowpass filtering (for i or j or $k = 1$) or highpass filtering (for i or j or $k = 2$).</p> <p>The i element filters along the rows of X, the j element filters along the columns, and the k element filters along the third dimension. For example, <code>dec{1,2,1}</code> is obtained by filtering X along the rows with the lowpass (scaling) filter, along the columns with the highpass (wavelet) filter, and along the third dimension with the lowpass (scaling) filter.</p>
-----	--

See Also

`dwtmode` | `idwt3` | `wavedec3` | `waveinfo` | `waverec3` | `wfilters`

Introduced in R2010a

dwtfilterbank

Discrete wavelet transform filter bank

Description

Use `dwtfilterbank` to create a discrete wavelet transform (DWT) filter bank. With the filter bank, you can visualize wavelets and scaling functions in time and frequency. You can measure the 3-dB bandwidths of the wavelet and scaling functions. You can measure the energy concentration of the wavelet and scaling functions in the theoretic DWT passbands. You can determine if the filter bank is orthogonal or biorthogonal. You can also determine the frame bounds of the filter bank. You can create a filter bank using your own custom filters.

Creation

Syntax

```
fb = dwtfilterbank
fb = dwtfilterbank(Name,Value)
```

Description

`fb = dwtfilterbank` create a discrete wavelet transform (DWT) filter bank. The default filter bank is designed for a signal with 1024 samples. The default filter bank uses the analysis (decomposition) `sym4` wavelet and scaling filter with seven resolution levels.

`fb = dwtfilterbank(Name,Value)` creates a DWT filter bank `fb` with properties specified by one or more `Name,Value` pair arguments. Properties can be specified in any order as `Name1,Value1,...,NameN,ValueN`. Enclose each property name in quotes.

Note You cannot change a property value of an existing filter bank. For example, if you have a filter bank `fb` for the `sym4` wavelet, you must create a second filter bank `fb2` for the `coif5` wavelet. You cannot assign a different Wavelet to `fb`.

Properties

SignalLength — Signal length

1024 (default) | positive integer greater than or equal to 2

Signal length, specified as a positive integer greater than or equal to 2.

Example: 'SignalLength',768

Data Types: double

Wavelet — Name of wavelet

'sym4' (default) | 'Custom' | character array | string scalar

Name of wavelet, specified as a character array or string scalar, used to construct the filter bank. **Wavelet** is an orthogonal or biorthogonal wavelet recognized by `wavemngr` or `Custom`.

To use a wavelet filter not recognized by `wavemngr`, set the **Wavelet** property to `Custom` and specify the `CustomWaveletFilter` and `CustomScalingFilter` properties.

Example: 'Wavelet','bior4.4'

Data Types: char | string

FilterType — Wavelet filter type

'Analysis' (default) | 'Synthesis'

Wavelet filter type, specified as one of 'Analysis' or 'Synthesis'. 'Analysis' uses the decomposition filters returned by `wfilters`. 'Synthesis' uses the reconstruction filters.

Level — Wavelet transform level

positive integer

Wavelet transform level, specified as a positive integer less than or equal to $\text{floor}(\log_2(\text{SignalLength}))$. By default the level is equal to $\text{floor}(\log_2(\text{SignalLength}/(L-1)))$ where L is the length of the wavelet filter associated with **Wavelet**. For wavelets recognized by `wavemngr`, this is equivalent to `wmaxlev(SignalLength,Wavelet)`. If $\text{floor}(\log_2(\text{SignalLength}/(L-1)))$ is less than or equal to 0, **Level** defaults to $\text{floor}(\log_2(\text{SignalLength}))$.

SamplingFrequency — Sampling frequency in hertz

1 (default) | positive scalar

Sampling frequency in hertz, specified as a positive scalar. If unspecified, frequencies are in cycles/sample and the Nyquist frequency is $\frac{1}{2}$.

Example: 'SamplingFrequency',5

Data Types: double

CustomWaveletFilter — Custom wavelet filter coefficients

even-length column vector | two-column matrix with even number of rows

Custom wavelet filter coefficients, specified as a real-valued column vector or matrix. `CustomWaveletFilter` must be an even-length column vector for an orthogonal wavelet or a two-column matrix with an even number of rows for a biorthogonal wavelet.

This property applies only when `Wavelet` is set to 'Custom'.

CustomScalingFilter — Custom scaling filter coefficients

even-length column vector | two-column matrix with even number of rows

Custom scaling filter coefficients, specified as a real-valued column vector or matrix. `CustomScalingFilter` must be an even-length column vector for an orthogonal wavelet or a two-column matrix with an even number of rows for a biorthogonal wavelet.

This property applies only when `Wavelet` is set to 'Custom'.

Object Functions

<code>dwtpassbands</code>	DWT filter bank passbands
<code>filters</code>	DWT filter bank filters
<code>framebounds</code>	DWT filter bank frame bounds
<code>freqz</code>	DWT filter bank frequency responses
<code>isBiorthogonal</code>	Determine if DWT filter bank is biorthogonal
<code>isOrthogonal</code>	Determine if DWT filter bank is orthogonal
<code>powerbw</code>	DWT filter bank power bandwidth
<code>qfactor</code>	DWT filter bank quality factor
<code>scalingfunctions</code>	DWT filter bank time-domain scaling functions
<code>wavelets</code>	DWT filter bank time-domain wavelets
<code>waveletsupport</code>	DWT filter bank time supports

Examples

Discrete Wavelet Transform Filter Bank with Default Values

Create a DWT filter bank using default values.

```
fb = dwtfilterbank
```

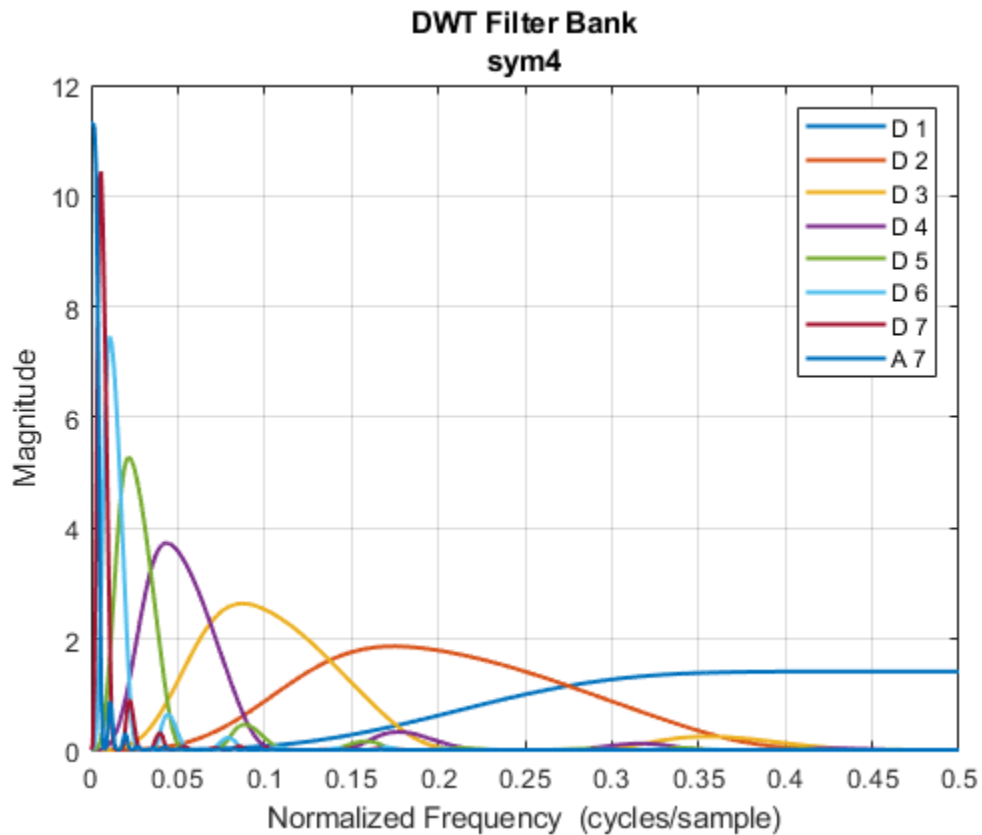
```
fb =
```

```
  dwtfilterbank with properties:
```

```
      Wavelet: 'sym4'  
    SignalLength: 1024  
        Level: 7  
    SamplingFrequency: 1  
      FilterType: 'Analysis'  
CustomWaveletFilter: []  
CustomScalingFilter: []
```

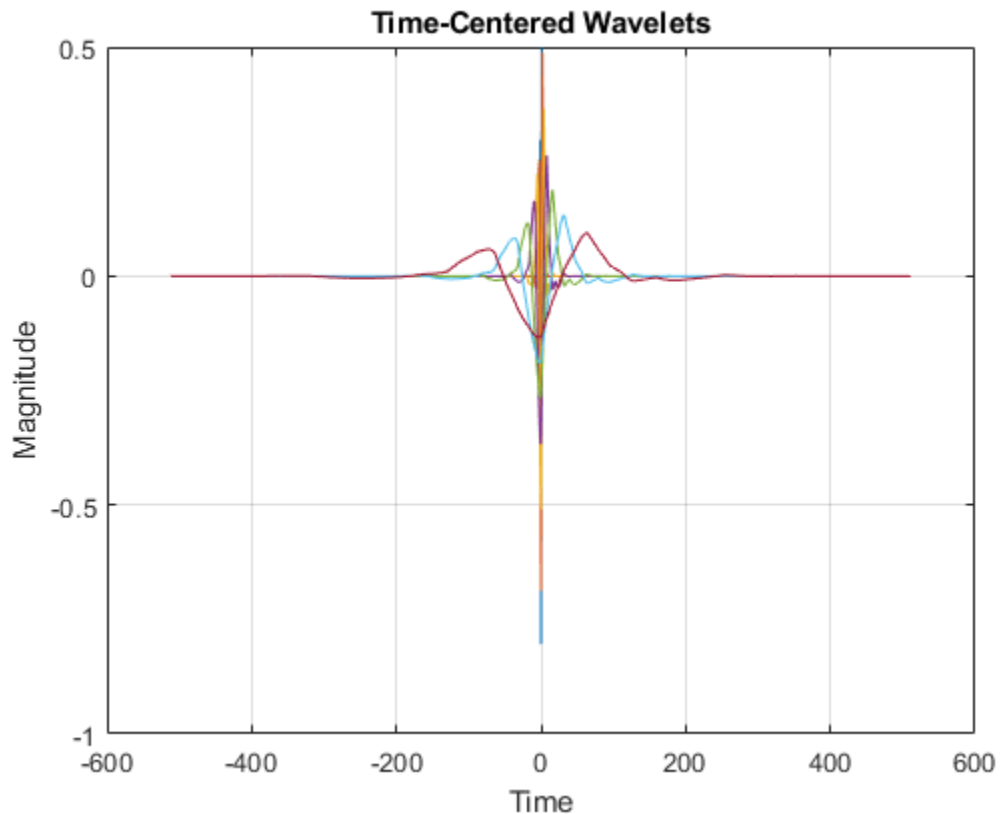
Plot the magnitude frequency responses of the wavelets and coarsest-scale scaling function. Open the plot in a separate figure window. The plot legend in the window is interactive. To hide a particular frequency response, click on its name.

```
freqz(fb)
```



Obtain and plot the time-centered wavelets corresponding to the wavelet bandpass filters.

```
[psi,t] = wavelets(fb);
plot(t,psi')
grid on
title('Time-Centered Wavelets')
xlabel('Time')
ylabel('Magnitude')
```



Create DWT Filter Bank Using Custom Filters

This example shows how to create a DWT filter bank using custom biorthogonal wavelet filters.

Two pairs of analysis (decomposition) and synthesis (reconstruction) filters are associated with a biorthogonal wavelet. Each pair consists of a lowpass and highpass filter. Specify the analysis and synthesis filters for the nearly-orthogonal biorthogonal wavelets based on the Laplacian pyramid scheme of Burt and Adelson (Table 8.4 on page 283 in [1]). Because the toolbox requires that all filters associated with a biorthogonal wavelet or an

orthogonal wavelet have the same even length, the filters are prepended and appended with 0s.

```
Hd = [0 -1 5 12 5 -1 0 0]/20*sqrt(2);
Gd = [0 3 -15 -73 170 -73 -15 3]/280*sqrt(2);
Hr = [0 -3 -15 73 170 73 -15 -3]/280*sqrt(2);
Gr = [0 -1 -5 12 -5 -1 0 0]/20*sqrt(2);
```

Hd and Gd are the lowpass and highpass decomposition filters, respectively. Hr and Gr are the lowpass and highpass reconstruction filters, respectively.

Construct analysis and synthesis DWT filter banks using the biorthogonal filters. Confirm the filter banks are biorthogonal and not orthogonal.

```
fbAna = dwtfilterbank('Wavelet','Custom',...
    'CustomScalingFilter',[Hd' Hr'],'CustomWaveletFilter',[Gd' Gr]);
fbSyn = dwtfilterbank('Wavelet','Custom',...
    'CustomScalingFilter',[Hd' Hr'],'CustomWaveletFilter',[Gd' Gr'],...
    'FilterType','Synthesis');
fprintf('fbAna: isOrthogonal = %d\tisBiorthogonal = %d\n',...
    isOrthogonal(fbAna),isBiorthogonal(fbAna));

fbAna: isOrthogonal = 0    isBiorthogonal = 1

fprintf('fbSyn: isOrthogonal = %d\tisBiorthogonal = %d\n',...
    isOrthogonal(fbSyn),isBiorthogonal(fbSyn));

fbSyn: isOrthogonal = 0    isBiorthogonal = 1
```

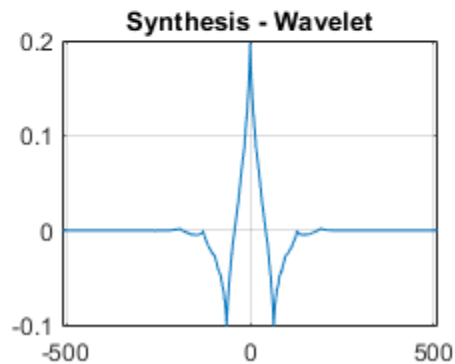
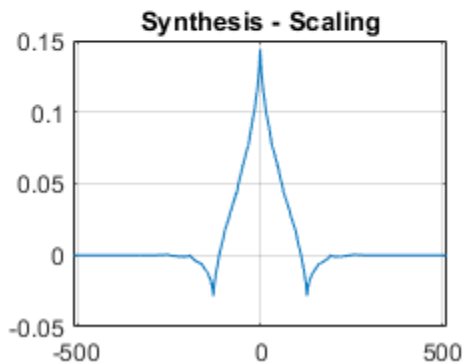
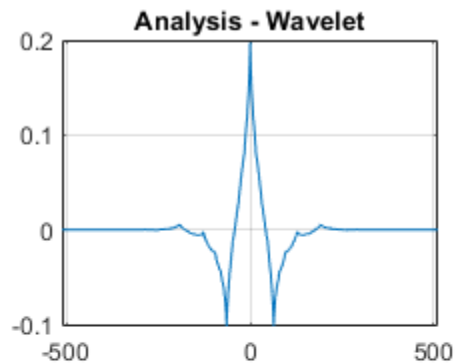
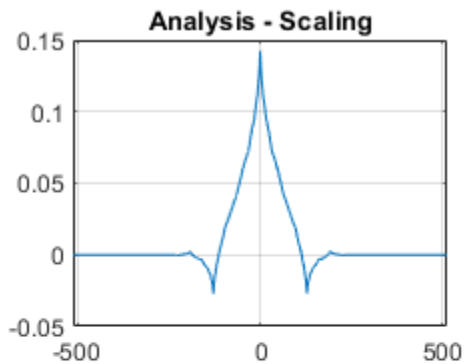
Obtain the wavelet and scaling functions of both filter banks. Plot the wavelet and scaling functions at the coarsest scales.

```
[fbAna_phi,t] = scalingfunctions(fbAna);
[fbAna_psi,~] = wavelets(fbAna);
[fbSyn_phi,~] = scalingfunctions(fbSyn);
[fbSyn_psi,~] = wavelets(fbSyn);
subplot(2,2,1)
plot(t,fbAna_phi(end,:))
grid on
title('Analysis - Scaling')
subplot(2,2,2)
plot(t,fbAna_psi(end,:))
grid on
title('Analysis - Wavelet')
subplot(2,2,3)
```

```

plot(t,fbSyn_phi(end,:))
grid on
title('Synthesis - Scaling')
subplot(2,2,4)
plot(t,fbSyn_psi(end,:))
grid on
title('Synthesis - Wavelet')

```



Compute the framebounds of the two filter banks. Since the filters are associated with biorthogonal wavelets, the framebounds will not equal 1.

```

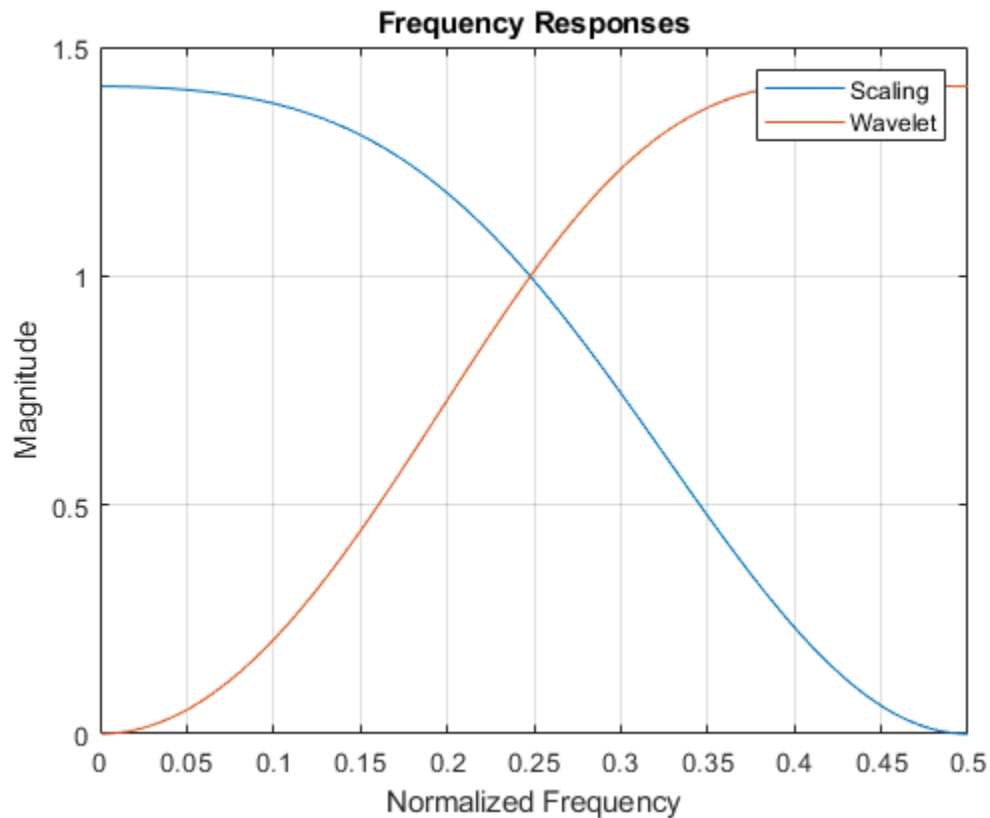
[a1,a2] = framebounds(fbAna)
a1 = 0.9505

```

```
a2 = 1.0211
[b1,b2] = framebounds(fbSyn)
b1 = 0.9800
b2 = 1.0528
```

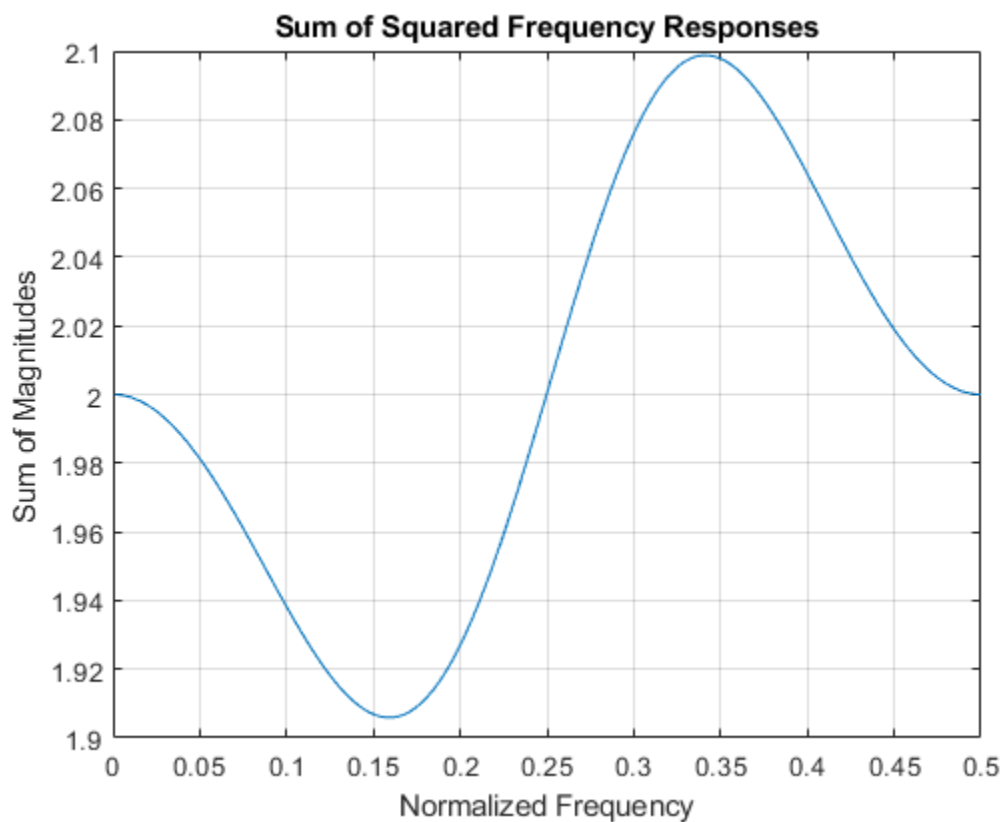
Obtain the frequency responses of the scaling and wavelets filters in the analysis filter bank. Plot up to Nyquist the magnitude frequency responses of the scaling and wavelet filters at the finest scale.

```
[psidft,f,phidft] = freqz(fbAna);
flen = length(f);
figure
plot(f(flen/2+1:end),abs(phidft(1,flen/2+1:end)))
hold on
plot(f(flen/2+1:end),abs(psidft(1,flen/2+1:end)))
grid on
legend('Scaling','Wavelet')
title('Frequency Responses')
xlabel('Normalized Frequency')
ylabel('Magnitude')
```



Zoom in and confirm the magnitude frequency responses at the point of intersection are not magnitude equal to 1. Plot the sum of the squared magnitudes of the frequency responses. Because the scaling (lowpass) and wavelet (highpass) filters do not form an orthogonal quadrature mirror filter pair, the sum does not equal to 2 at all frequencies.

```
figure
plot(f(flen/2+1:end),abs(phidft(1,flen/2+1:end)).^2 + abs(psidft(1,flen/2+1:end)).^2)
grid on
title('Sum of Squared Frequency Responses')
xlabel('Normalized Frequency')
ylabel('Sum of Magnitudes')
```



References

- [1] Daubechies, I. *Ten Lectures on Wavelets*. CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: Society for Industrial and Applied Mathematics, 1992.

See Also

dwt | modwt | wavedec | wavemngr

Topics

“Adding Quadrature Mirror and Biorthogonal Wavelet Filters”

Introduced in R2018a

dwtleader

Multifractal 1-D wavelet leader estimates

Syntax

```
[dh,h] = dwtleader(x)
[dh,h,cp] = dwtleader(x)
[dh,h,cp,tauq] = dwtleader(x)
[dh,h,cp,tauq,leaders] = dwtleader( ___ )
[dh,h,cp,tauq,leaders,structfunc] = dwtleader( ___ )

[ ___ ]= dwtleader(x,wname)
[ ___ ] = dwtleader( ___ ,Name,Value)
```

Description

[dh,h] = dwtleader(x) returns the singularity spectrum, dh, and the Holder exponents, h, for the 1-D real-valued data, x. The singularity spectrum and Holder exponents are estimated for the linearly-spaced moments of the structure functions from -5 to +5.

[dh,h,cp] = dwtleader(x) also returns the first three log cumulants, cp of the scaling exponents.

[dh,h,cp,tauq] = dwtleader(x) also returns the scaling exponents for the linearly spaced moments from -5 to 5. Wavelet leaders are not defined for the finest scale.

[dh,h,cp,tauq,leaders] = dwtleader(___) also returns the wavelet leaders by scale.

[dh,h,cp,tauq,leaders,structfunc] = dwtleader(___) also returns the multiresolution structure functions.

[___]= dwtleader(x,wname) uses the orthogonal or biorthogonal wavelet specified by wname to compute the wavelet leaders and the fractal estimates.

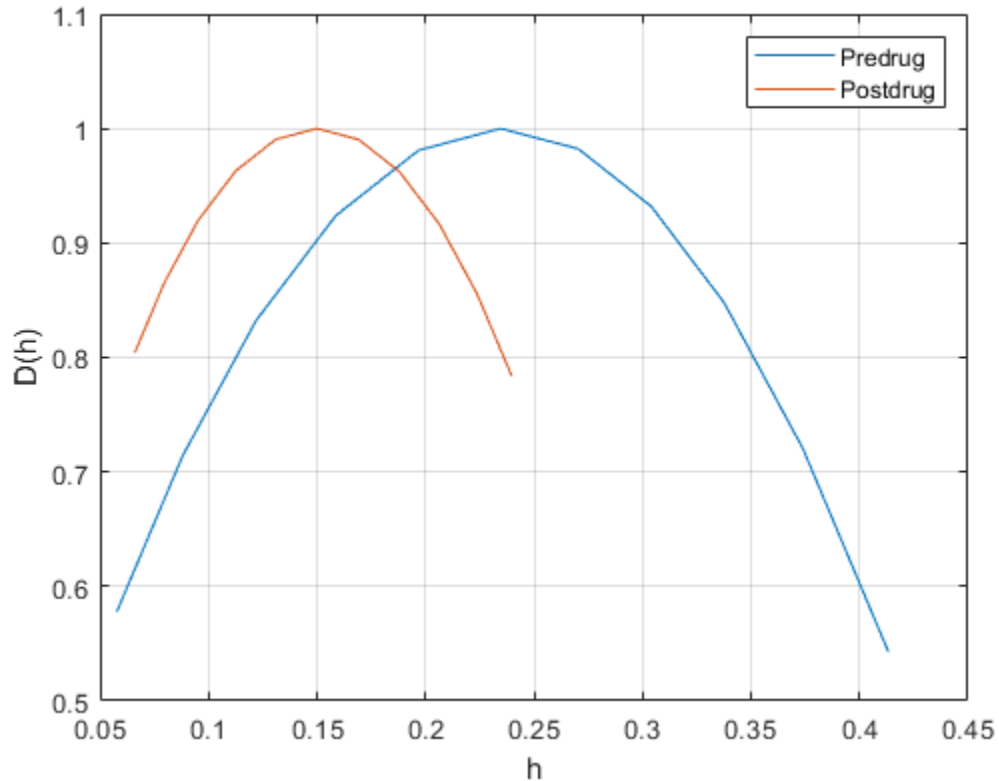
[___] = `dwtleader`(____, Name, Value) returns the wavelet leaders and other specified outputs with additional options specified by one or more Name, Value pair arguments.

Examples

Multifractal Spectrum of Heart-Rate Variability

Compare the multifractal spectrum of heart-rate variability data before and after application of a drug that reduces heart dynamics.

```
load hrvDrug;
predrug = hrvDrug(1:4642);
postdrug = hrvDrug(4643:end);
[dhpre,hpre] = dwtleader(predrug);
[dhpost,hpost] = dwtleader(postdrug);
plot(hpre,dhpre,hpost,dhpost)
xlabel('h')
ylabel('D(h)')
grid on
legend('Predrug','Postdrug')
```

The spread of the Holder exponent values before drug administration (approximately 0.08 to 0.55) is much larger than the spread of the values afterward (approximately 0.08 to 0.31). This indicates that the heart rate has become more monofractal.

Brownian Noise Singularity Spectrum

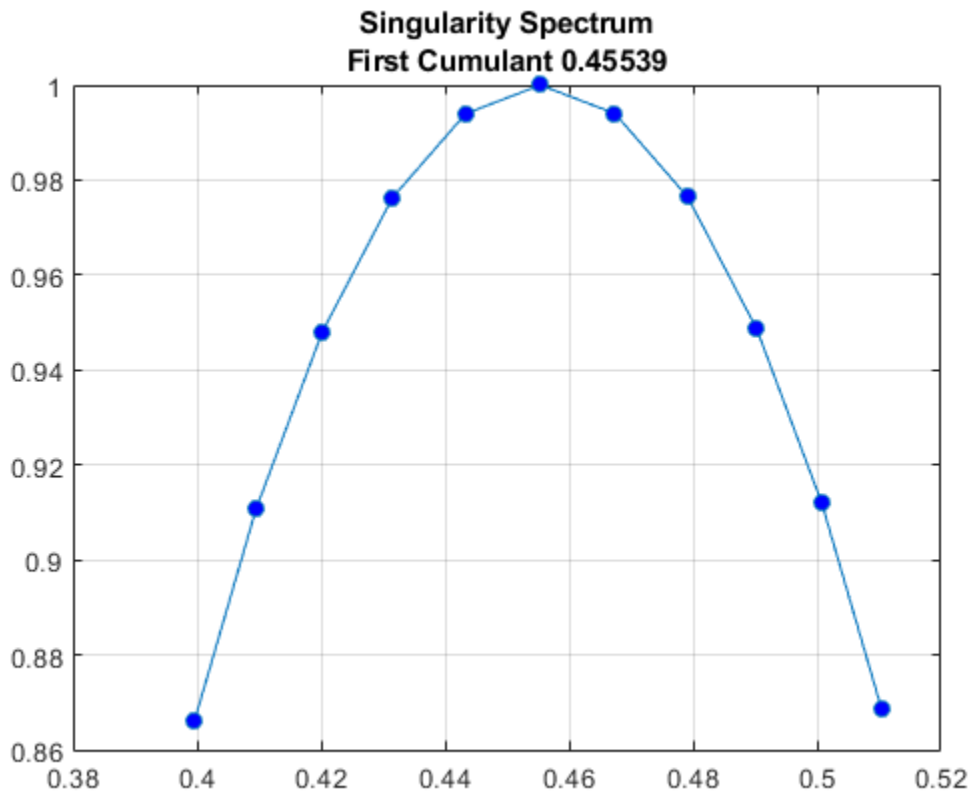
Compute the singularity spectrum and cumulants for a Brownian noise process.

Create the Brownian noise signal.

```
rng(100);
x = cumsum(randn(2^15,1));
```

Obtain and plot the singularity spectrum.

```
[dh,h,cp] = dwtleader(x);  
plot(h,dh,'o-','MarkerFaceColor','b')  
grid on  
title({'Singularity Spectrum'; ['First Cumulant ' num2str(cp(1))]}))
```



The small spread in the Holder exponents (approximately 0.472 to 0.512) indicates that this Brownian noise signal can be characterized by a global Holder exponent of 0.49875. The theoretical Holder exponent for Brownian motion is 0.5.

Obtain the cumulants.

```
cp
```

```
cp = 1×3
    0.4554    -0.0121    -0.0000
```

The first cumulant value is the slope of scaling exponents versus the moments. The second and third cumulants indicate the deviation from linearity. The first cumulant value and near-zero values of the second and third cumulants indicate that the scaling exponents are a linear function of the moments. Therefore, this Brownian motion signal is monofractal.

Multifractal Random Walk Cumulants

Compute the cumulants for a multifractal random walk. The multifractal random walk is a realization of a random process with a theoretical first cumulant of 0.75 and a second cumulant -0.05. The second cumulant value of -0.05 indicates that the scaling exponents deviate from a linear function with slope 0.75.

Load a random walk signal.

```
load mrw07505
```

Obtain and display the first and second cumulants.

```
[~,~,cp,tauq] = dwtleader(mrw07505);
cp([1 2])

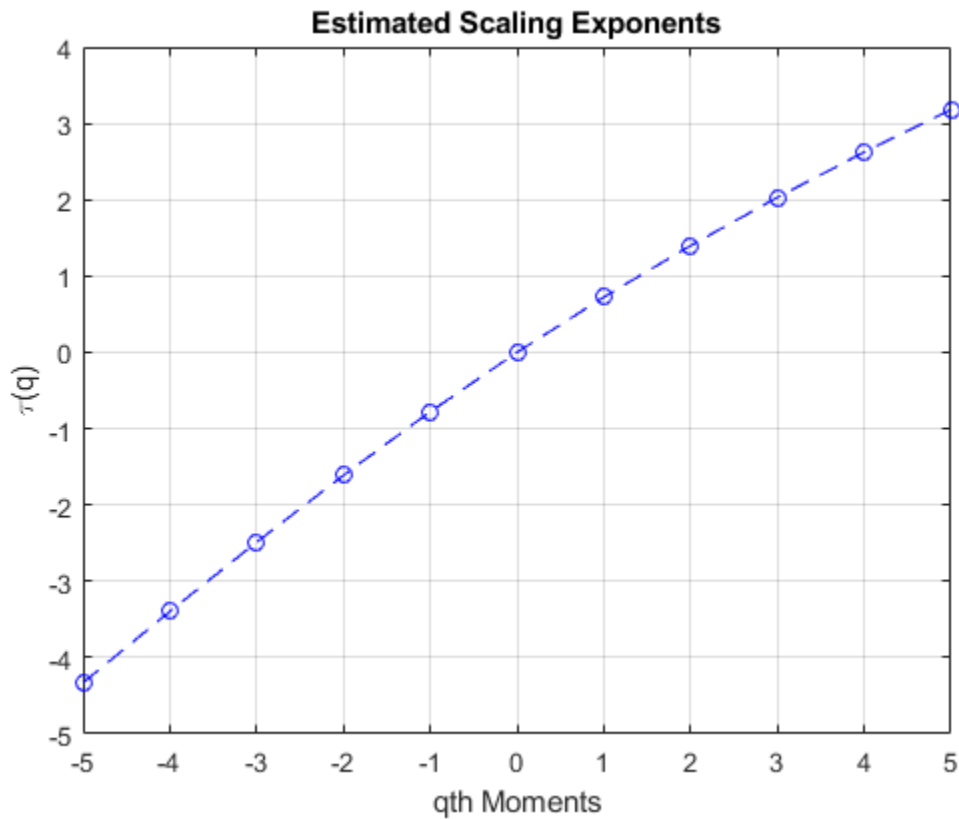
ans = 1×2
    0.7504    -0.0554
```

For monofractal processes, the scaling exponents are a linear function of the moments. Linearity is indicated by the second and third cumulants being close to zero. In this case, the nonzero second cumulant indicates that the process is multifractal.

Plot the scaling exponents for the q th moments.

```
plot(-5:5,tauq,'bo--')
title('Estimated Scaling Exponents')
grid on
```

```
xlabel('qth Moments')  
ylabel('\tau(q)')
```



The scaling exponents are a nonlinear function of the moments.

Input Arguments

x — Input signal

vector of real values

Input signal, specified as a 1-D vector of real values. For the default wavelet and minimum regression level, the time series must have at least 248 samples. For nondefault

values, the minimum-required data length depends on the wavelet filter and the levels used in the regression model. The wavelet leaders technique works best for data with 8000 or more samples.

wname — Wavelet name

'bior1.5' (default) | character vector | string scalar

Wavelet name, specified as a character vector or string scalar. `wname` is a wavelet family short name and filter number recognized by the wavelet manager, `wavemngr`.

To query valid wavelet family short names, use `wavemngr('read')`. To determine whether a particular wavelet is orthogonal or biorthogonal, use `waveinfo` with the wavelet family short name, for example, `waveinfo('db')`. Alternatively, use `wavemngr` with the 'type' option, for example, `wavemngr('type','fk4')`. A returned value of 1 indicates an orthogonal wavelet. A returned value of 2 indicates a biorthogonal wavelet.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1,Value1,...,NameN,ValueN`.

Example: `'MinRegressionLevel',5` sets the minimum regression level to 5.

RegressionWeight — Weight option

'uniform' (default) | 'scale'

Weight option to use in the weighted least-squares regression model to determine the singularity spectrum, Holder exponents, cumulants, and scaling exponents, specified as the comma-separated pair consisting of 'RegressionWeight' and either 'uniform' or 'scale'. The 'uniform' option applies equal weight to each scale. The 'scale' option uses the number of wavelet leaders by scale as weights.

Note To duplicate the behavior of `dwtleader` found in releases prior to R2018a, update all instances of `dwtleader` to include the name-value pair argument 'RegressionWeight' set to 'scale'.

MinRegressionLevel — Minimum regression level

3 (default) | positive integer

Minimum regression level, *minlev*, specified as the comma-separated pair consisting of 'MinRegressionLevel' and a positive integer greater than or equal to 2. Only levels greater than or equal to the specified minimum level are used in the multifractal estimates. `dwtLeader` requires at least 6 wavelet leaders at the maximum level and two levels to be used in the multifractal estimates. The scale in the discrete wavelet transform corresponding to the minimum level is 2^{minlev} . The smoother the data (that is, the closer the Holder exponents are to 1), the less likely that reducing the minimum regression level will degrade the results.

MaxRegressionLevel — Maximum regression level

positive integer

Maximum regression level, *maxlev*, specified as a positive integer greater than or equal to *minlev* + 1. The maximum level uses only levels less than or equal to *maxlev* in the multifractal estimates. The scale in the discrete wavelet transform corresponding to the maximum level is 2^{maxlev} . Specify a maximum regression level when you want to restrict the levels used in the regression to a value less than the default level. To determine the number of wavelet leaders by level, use the `leaders` output argument, or the `weights` field of the `structfunc` output argument. The default value is the largest level with at least six wavelet leaders

Output Arguments

dh — Singularity spectrum

vector

Singularity spectrum, returned as a vector. The singularity spectrum is estimated using structure functions determined for the linearly-spaced moments from -5 to 5. The structure functions are computed based on the wavelet leaders obtained using the biorthogonal spline wavelet filter. The biorthogonal spline wavelet filter that is used has one vanishing moment in the synthesis wavelet and five vanishing moments in the analysis wavelet ('`bior1.5`'). By default, multifractal estimates are derived from wavelet leaders at a minimum level of 3 and maximum level where there are at least six wavelet leaders.

h — Holder exponent estimates

1-by-11 vector of real scalars

Holder exponent estimates, returned as a 1-by-11 vector of scalars. Holder exponents characterize signal regularity. The closer a Holder exponent is to 1, the closer the

function is to differentiable. Conversely, the closer the Holder exponent is to zero, the closer the function is to discontinuous.

Data Types: double

cp — Cumulants

vector

Cumulants, returned as a 1-by-3 vector of scalars. The vector contains the first three log cumulants of the scaling exponents. The first cumulant characterizes the linear behavior in the scaling exponents. The second and third cumulants characterize the departure from linearity.

Data Types: double

tauq — Scaling exponents

column vector

Scaling exponents, returned as a column vector. The exponents are for the linearly-spaced moments from -5 to +5.

Leaders — Wavelet leaders

cell array

Leaders is a cell array with the i th element containing the wavelet leaders at level $i+1$, or scale $2^{(i+1)}$. Wavelet leaders are not defined at level 1.

structfunc — Multiresolution structure functions

struct

Multiresolution structure functions for the global Holder exponent estimates, returned as a struct. The structure function for data x is defined as

$$S(q, a) = \frac{1}{n_a} \sum_{k=1}^{n_a} |T_x(a, k)|^q \approx a^{\zeta(q)},$$

where a is the scale, q is the moment, T_x are the wavelet leaders by scale, n_a is the number of wavelet leaders at each scale, and $\zeta(q)$ is the scaling exponent. Expanding $\zeta(q)$ to a polynomial produces

$$\zeta(q) = c_1 q + c_2 q^2/2 + c_3 q^3/6 + \dots$$

The scaling exponents can be estimated from the log-cumulants of the wavelet leader coefficients. When $\zeta(q)$ is a linear function, the signal is monofractal. When it deviates from linear, the signal is multifractal.

`structfunc` is a structure array containing the following fields:

- `Tq` — Measurements of the input, x , at various scales. `Tq` is a matrix of multiresolution quantities that depend jointly on time and scale. Scaling phenomena in x imply a power-law relationship between the moments of `Tq` and the scale. For `dwtLeader`, the `Tq` field is an N_s -by-36 matrix, where N_s is the number of scales used in the multifractal estimates. The first 11 columns of `Tq` are the scaling exponent estimates by scale for each of the q th moments from -5 to 5. The next 11 columns contain the singularity spectrum estimates, `dh`, for each of the q th moments. Columns 23-33 contain the Holder exponent estimates, `h`. The last three columns contain the estimates for the first-order, second-order, and third-order cumulants, respectively.
- `weights` — Weights used in the regression. The weights are the number of wavelet leaders by scale. `weights` is an N_s -by-1 vector.
- `logscals` — Scales used as predictors in the regression. `logscals` is an N_s -by-1 vector with the base-2 logarithm of the scales.

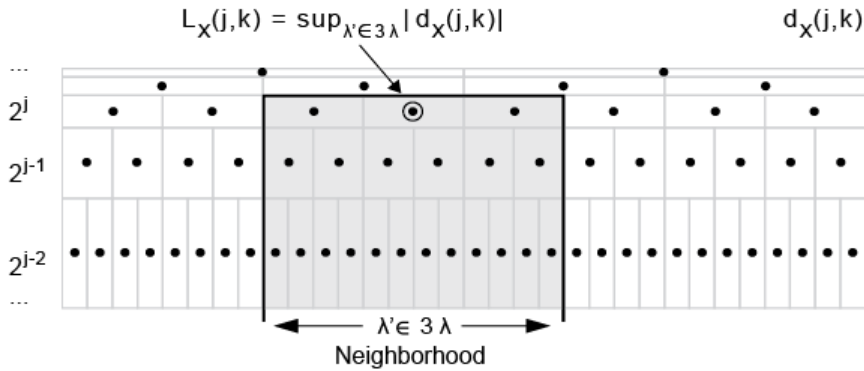
Algorithms

Wavelet leaders are derived from the critically sampled discrete wavelet transform (DWT) coefficients. Wavelet leaders offer significant theoretical advantages over wavelet coefficients in the multifractal formalism. Wavelet leaders are time- or space-localized suprema of the absolute value of the discrete wavelet coefficients. The time localization of the suprema requires that the wavelet coefficients are obtained using a compactly supported wavelet. The Holder exponents, which quantify the local regularity, are determined from these suprema. The singularity spectrum indicates the size of the set of Holder exponents in the data.

1-D wavelet leaders are defined as

$$L_x(j, k) = \sup_{\lambda' \subset 3\lambda_{j,k}} |d_x(j, k)|$$

where the scales are 2^j , translated to time positions $2^j k$. The time neighborhood is $3\lambda_{j,k} = \lambda_{j,k-1} \cup \lambda_{j,k} \cup \lambda_{j,k+1}$, where $\lambda_{j,k} = [k2^j, (k+1)2^j)$. The time neighborhood is taken over the scale and all finer scales. $d_x(j,k)$ are the wavelet coefficients.



To calculate the wavelet leaders, $L_x(j,k)$:

- 1 Compute the wavelet coefficients, $d_x(j,k)$, using the discrete wavelet transform and save the absolute value of each coefficient for each scale. Each finer scale has twice the number of coefficients than the next coarser scale. Each dyadic interval at scale 2^j can be written as a union of two intervals at a finer scale.

$$[2^j k, 2^j(k+1)) = [2^{j-1}(2k), 2^{j-1}(2k+2))$$

$$[2^{j-1}(2k), 2^{j-1}(2k+2)) = [2^{j-1}(2k), 2^{j-1}(2k+1)) \cup [2^{j-1}(2k+1), 2^{j-1}(2k+2))$$

- 2 Start at the scale that is one level coarser than the finest obtained scale.
- 3 Compare the first value to all its finer dyadic intervals and obtain the maximum value.
- 4 Go to the next value and compare its value to all of its finer scale values.
- 5 Continue comparing the values with their nested values and obtaining the maxima.
- 6 From the maximum values obtained for that scale, examine the first three values and obtain the maximum of those neighbors. That maximum value is a leader for that scale.
- 7 Continue comparing the maximum values to obtain the other leaders for that scale.
- 8 Move to the next coarser scale and repeat the process.

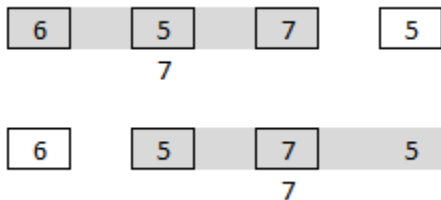
For example, assume that you have these absolute values of the coefficients at these scales:

6		2		7		5	
4	3	5	2	1	0	4	3

Starting with the top row, which is the next coarsest level from the finest scale (bottom row), compare each value to its dyadic intervals and obtain the maxima.

6		2		7		5	
4	3	5	2	1	0	4	3
6		5		7		5	

Then, look at the three neighboring values and obtain the maximum. Repeat for the next three neighbors. These maxima, 7 and 7, are the wavelet leaders for this level.



References

- [1] Wendt, H., and P. Abry. "Multifractality Tests Using Bootstrapped Wavelet Leaders." *IEEE Transactions on Signal Processing*. Vol. 55, No. 10, 2007, pp. 4811-4820.
- [2] Jaffard, S., B. Lashermes, and P. Abry. "Wavelet Leaders in Multifractal Analysis." *Wavelet Analysis and Applications*. T. Qian, M. I. Vai, and X. Yuesheng, Eds. 2006, pp. 219-264.

See Also

wfbm | wtmm

Topics

“Multifractal Analysis”

Introduced in R2016b

dwtmode

Discrete wavelet transform extension mode

Syntax

```
dwtmode(mode)
```

```
dwtmode  
dwtmode('status')  
st = dwtmode  
st = dwtmode('status')  
st = dwtmode('status','nodisp')
```

```
dwtmode('save',mode)  
dwtmode('save')  
dwtmode('save',CURRENTMODE)
```

Description

`dwtmode(mode)` sets the signal or image extension mode for both discrete wavelet and wavelet packet transforms to `mode`. All functions and Wavelet Analyzer app tools involving either the discrete wavelet transform (1-D and 2-D) or wavelet packet transform (1-D and 2-D), use the specified DWT extension mode.

The extension modes provide options for dealing with the problem of border distortion in signal or image analysis. For more information, see “Border Effects”.

`dwtmode` or `dwtmode('status')` display the current mode. If `DWTMODE.DEF` exists in the current path, the default mode is loaded from `DWTMODE.DEF` at the start of the MATLAB session. Otherwise, the file `DWTMODE.CFG` is used.

`st = dwtmode` or `st = dwtmode('status')` display and return the current mode in `st`.

`st = dwtmode('status','nodisp')` returns the current mode `st` and no status or warning text is displayed in the MATLAB command window.

`dwtmode('save',mode)` saves `mode` as the new default mode to the file `DWTMODE.DEF` in the current folder. If `DWTMODE.DEF` already exists in the current folder, the file is overwritten. The new default mode will be active as default mode in the next MATLAB session.

`dwtmode('save')` is equivalent to `dwtmode('save',CURRENTMODE)`, where `CURRENTMODE` represents the current extension mode.

Examples

Display and Change Signal Extension Mode

Clear the DWT extension mode global variable, and display the current DWT signal extension mode. If the DWT extension mode global variable does not exist, the default is half-point symmetrization.

```
clear global
dwtmode
```

```
*****
**  DWT Extension Mode: Symmetrization (half-point)  **
*****
```

Change the extension mode to periodized extension.

```
dwtmode('per')
```

```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!  WARNING: Change DWT Extension Mode  !
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

*****
**  DWT Extension Mode: Periodization  **
*****
```

Display the current DWT signal extension mode.

```
dwtmode
```

```
*****
**   DWT Extension Mode: Periodization  **
*****
```

Input Arguments

mode — Discrete wavelet transform extension mode

'zpd' | 'sp0' | 'spd' | ...

DWT extension mode used to extend the input, specified as one of the following values.

mode	DWT Extension Mode
'zpd'	Zero extension
'sp0'	Smooth extension of order 0
'spd' (or 'sp1')	Smooth extension of order 1
'sym' or 'symh'	Symmetric extension (half point): boundary value symmetric replication
'symw'	Symmetric extension (whole point): boundary value symmetric replication
'asym' or 'asymh'	Antisymmetric extension (half point): boundary value antisymmetric replication
'asymw'	Antisymmetric extension (whole point): boundary value antisymmetric replication
'ppd'	Periodized extension (1)
'per'	Periodized extension (2) If the signal length is odd, <code>wextend</code> adds to the right an extra sample that is equal to the last value, and performs the extension using the 'ppd' mode. Otherwise, 'per' reduces to 'ppd'. This rule also applies to images.

The DWT associated with the symmetric, smooth, zero, and periodic extension modes are slightly redundant. But the inverse DWT ensures a perfect reconstruction for the extensions mentioned.

Note `dwtmode` updates a global variable. Only use `dwtmode` to change the extension mode. Avoid changing the global variable directly.

Output Arguments

st — DWT extension mode

character array

DWT extension mode, returned as a character array.

Tips

- For most wavelet applications, either a periodic extension or symmetric extension works fine.

References

[1] Strang, G., and T. Nguyen. *Wavelets and Filter Banks*. Wellesley, MA: Wellesley-Cambridge Press, 1996.

See Also

Apps

Wavelet Analyzer

Functions

`dwt` | `dwt2` | `idwt` | `idwt2` | `wextend`

Introduced before R2006a

dwtpassbands

DWT filter bank passbands

Syntax

```
dwtbands = dwtpassbands(fb)
```

Description

`dwtbands = dwtpassbands(fb)` returns the theoretical discrete wavelet transform (DWT) passbands for the DWT filter bank, `fb`. `dwtbands` is a $L+1$ -by-2 matrix, where L is the filter bank `Level`. The first L rows of `dwtbands` contain the theoretical passband frequencies for the DWT listed in order of decreasing resolution (increasing scale). The first column of `dwtbands` contains the lower frequency limit and the second column contains the upper frequency limit. The final row of `dwtbands` contains the theoretical passband for the coarsest resolution scaling filter.

Examples

DWT Filter Bank Passbands

Obtain the theoretical DWT passbands for a four level wavelet transform using the Daubechies `db6` wavelet with a sampling frequency of 1 kHz.

```
wv = 'db6';  
Fs = 1e3;  
fb = dwtfilterbank('Wavelet',wv,'Level',4,'SamplingFrequency',Fs);  
dwtbands(fb)
```

```
ans = 5×2
```

```
250.0000 500.0000  
125.0000 250.0000  
62.5000 125.0000
```



```

31.2500    62.5000
           0    31.2500

```

Obtain the power bandwidths for the filter bank. Compare the theoretical passbands with the measured wavelet 3-dB bandwidths at all four levels.

```

ptable = powerbw(fb);
ptable(:,1:3)

```

```

ans=4x3 table
      Level      DWTBand      Wavelet3dBBandwidth
      _____  _____  _____
           1           250           500           250.98           500
           2           125           250           124.18           254.69
           3           62.5           125           62.578           127.76
           4           31.25          62.5           31.792           64.366

```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

dwtbands — Theoretical DWT passbands

real-valued matrix

Theoretical DWT passbands for the filter bank `fb`, returned as an $L+1$ -by-2 real-valued matrix, where L is the filter bank `Level`. The first L rows of `dwtbands` contain the theoretical passband frequencies for the DWT listed in order of decreasing resolution (increasing scale). The first column of `dwtbands` contains the lower frequency limit and the second column contains the upper frequency limit. The final row of `dwtbands` contains the theoretical passband for the coarsest resolution scaling filter.

See Also

`dwtfiltbank` | `powerbw`

Introduced in R2018a

dyaddown

Dyadic downsampling

Syntax

```

Y = dyaddown(X,EVENODD)
Y = dyaddown(X)
Y = dyaddown(X,EVENODD,'type')
Y = dyaddown(X,'type',EVENODD)
Y = dyaddown(X)
Y = dyaddown(X,'type')
Y = dyaddown(X,θ,'type')
Y = dyaddown(X,EVENODD)
Y = dyaddown(X,EVENODD,'c')

```

Description

$Y = \text{dyaddown}(X, \text{EVENODD})$ where X is a *vector*, returns a version of X that has been downsampled by 2. Whether Y contains the even- or odd-indexed samples of X depends on the value of positive integer EVENODD :

- If EVENODD is even, then $Y(k) = X(2k)$.
- If EVENODD is odd, then $Y(k) = X(2k+1)$.

$Y = \text{dyaddown}(X)$ is equivalent to $Y = \text{dyaddown}(X, \theta)$ (even-indexed samples).

$Y = \text{dyaddown}(X, \text{EVENODD}, 'type')$ or $Y = \text{dyaddown}(X, 'type', \text{EVENODD})$, where X is a *matrix*, returns a version of X obtained by suppressing one out of two:

Columns of X	If <i>'type'</i> = <i>'c'</i>
Rows of X	If <i>'type'</i> = <i>'r'</i>
Rows and columns of X	If <i>'type'</i> = <i>'m'</i>

according to the parameter EVENODD , which is as above.

If you omit the *EVENODD* or *'type'* arguments, *dyaddown* defaults to *EVENODD* = 0 (even-indexed samples) and *'type'* = *'c'* (columns).

$Y = \text{dyaddown}(X)$ is equivalent to $Y = \text{dyaddown}(X, 0, 'c')$.

$Y = \text{dyaddown}(X, 'type')$ is equivalent to $Y = \text{dyaddown}(X, 0, 'type')$.

$Y = \text{dyaddown}(X, \text{EVENODD})$ is equivalent to $Y = \text{dyaddown}(X, \text{EVENODD}, 'c')$.

Examples

```
% For a vector.
```

```
s = 1:10
```

```
s =
```

```
1 2 3 4 5 6 7 8 9 10
```

```
dse = dyaddown(s) % Downsample elements with even indices.
```

```
dse =
```

```
2 4 6 8 10
```

```
% or equivalently
```

```
dse = dyaddown(s,0)
```

```
dse =
```

```
2 4 6 8 10
```

```
dso = dyaddown(s,1) % Downsample elements with odd indices.
```

```
dso =
```

```
1 3 5 7 9
```

```
% For a matrix.
```

```
s = (1:3)'*(1:4)
```

```
s =
```

```
1 2 3 4
```

```
2 4 6 8
```

```
3 6 9 12
```

```
dec = dyaddown(s,0,'c') % Downsample columns with even indices.
```

```
dec =
```

```
2 4
```

```
4 8
```

```
6 12
```

```
der = dyaddown(s,1,'r') % Downsample rows with odd indices.
```

```
der =
```

```
1 2 3 4
```

```
3 6 9 12
```

```
dem = dyaddown(s,1,'m') % Downsample rows and columns
                                % with odd indices.
dem =
     1     3
     3     9
```

References

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

See Also

dyadup

Introduced before R2006a

dyadup

Dyadic upsampling

Syntax

```
Y = dyadup(X, EVENODD)
Y = dyadup(X)
Y = dyadup(X, EVENODD, 'type')
Y = dyadup(X, 'type', EVENODD)
Y = dyadup(X)
Y = dyadown(X, 1, 'c')
Y = dyadup(X, 'type')
Y = dyadup(X, 1, 'type')
Y = dyadup(X, EVENODD)
Y = dyadup(X, EVENODD, 'c')
```

Description

`dyadup` implements a simple zero-padding scheme very useful in the wavelet reconstruction algorithm.

$Y = \text{dyadup}(X, \text{EVENODD})$, where X is a *vector*, returns an extended copy of vector X obtained by inserting zeros. Whether the zeros are inserted as even- or odd-indexed elements of Y depends on the value of positive integer EVENODD :

- If EVENODD is even, then $Y(2k-1) = X(k)$, $Y(2k) = 0$.
- If EVENODD is odd, then $Y(2k-1) = 0$, $Y(2k) = X(k)$.

$Y = \text{dyadup}(X)$ is equivalent to $Y = \text{dyadup}(X, 1)$ (odd-indexed samples).

$Y = \text{dyadup}(X, \text{EVENODD}, \text{'type'})$ or $Y = \text{dyadup}(X, \text{'type'}, \text{EVENODD})$, where X is a *matrix*, returns extended copies of X obtained by inserting

Columns in X	If $\text{'type'} = \text{'c'}$
----------------	---------------------------------

Rows in X	If ' <i>type</i> ' = 'r'
Rows and columns in X	If ' <i>type</i> ' = 'm'

according to the parameter *EVENODD*, which is as above.

If you omit the *EVENODD* or '*type*' arguments, *dyadup* defaults to *EVENODD* = 1 (zeros in odd-indexed positions) and '*type*' = 'c' (insert columns).

$Y = \text{dyadup}(X)$ is equivalent to $Y = \text{dyaddown}(X, 1, 'c')$.

$Y = \text{dyadup}(X, 'type')$ is equivalent to $Y = \text{dyadup}(X, 1, 'type')$.

$Y = \text{dyadup}(X, \text{EVENODD})$ is equivalent to $Y = \text{dyadup}(X, \text{EVENODD}, 'c')$.

Examples

```
% For a vector.
```

```
s = 1:5
```

```
s =
```

```
1 2 3 4 5
```

```
dse = dyadup(s) % Upsample elements at odd indices.
```

```
dse =
```

```
0 1 0 2 0 3 0 4 0 5 0
```

```
% or equivalently
```

```
dse = dyadup(s,1)
```

```
dse =
```

```
0 1 0 2 0 3 0 4 0 5 0
```

```
dso = dyadup(s,0) % Upsample elements at even indices.
```

```
dso =
```

```
1 0 2 0 3 0 4 0 5
```

```
% For a matrix.
```

```
s = (1:2)'+(1:3)
```

```
s =
```

```
1 2 3
```

```
2 4 6
```

```
der = dyadup(s,1,'r') % Upsample rows at even indices.
```

```
der =
```

```
0 0 0
```

```
1 2 3
0 0 0
2 4 6
0 0 0

doc = dyadup(s,0,'c') % Upsample columns at odd indices.
doc =
    1 0 2 0 3
    2 0 4 0 6
dem = dyadup(s,1,'m') % Upsample rows and columns
                    % at even indices.
dem =
    0     0     0     0     0     0     0
    0     1     0     2     0     3     0
    0     0     0     0     0     0     0
    0     2     0     4     0     6     0
    0     0     0     0     0     0     0

% Using default values for dyadup and dyaddown, we have:
% dyaddown(dyadup(s)) = s.
s = 1:5
s =
    1 2 3 4 5

uds = dyaddown(dyadup(s))
uds =
    1 2 3 4 5

% In general reversed identity is false.
```

References

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- If X is empty, generated code returns X and MATLAB returns `[]`.
- Suppose that all of the following conditions are true:
 - X is a variable-size array.
 - X is not a variable-length column vector (`:-by-1`).
 - X is a column vector at run time.
 - `'type'` is not supplied.

In generated code, the output for `y = dyadup(X,k)`, where k is optional, matches the output for `y = dyadup(X,k,'c')`. In MATLAB, the output for `y = dyadup(X,k)` matches the output for `y = dyadup(X,k,'r')`.

For code generation, when you do not specify `'type'`, if you want `dyadup` to treat X as a column vector, X must be a variable-length vector (`:-by-1`).

See Also

`dyaddown`

Introduced before R2006a

editLabelDefinition

Edit label definition properties

Syntax

```
editLabelDefinition(lss, lblname, propname, val)
```

Description

`editLabelDefinition(lss, lblname, propname, val)` changes the `propname` property of the label or sublabel definition `lblname` to `val`.

The function can edit only the “Name” on page 1-0 , “DefaultValue” on page 1-0 , “Tag” on page 1-0 , “Description” on page 1-0 , and “Categories” on page 1-0 properties. To change any other property of the label definition, remove the definition using `removeLabelDefinition` and add a definition with the desired property values using `addLabelDefinitions`.

- If you edit the “DefaultValue” on page 1-0 property, all existing label values remain unchanged. The new default value applies only to new members, new regions, or new points.
- You can edit the “Categories” on page 1-0 property only when the “LabelDataType” on page 1-0 of the target label or sublabel definition is 'Categorical'.

New specified categories do not replace any existing categories. They are appended to the existing values.

Examples

Edit Label Definition

Load a labeled signal set containing recordings of whale songs. Get the names of the labels.

```
load whales
lss

lss =
  labeledSignalSet with properties:
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

```
getLabelNames(lss)
```

```
ans = 3x1 string array
    "WhaleType"
    "MoanRegions"
    "TrillRegions"
```

The first label corresponds to the type of whale. Get the types available in the set.

```
lbldefs = getLabelDefinitions(lss);
types = lbldefs(1)

types =
  signalLabelDefinition with properties:
      Name: "WhaleType"
      LabelType: "attribute"
      LabelDataType: "categorical"
      Categories: [3x1 string]
      DefaultValue: []
      Sublabels: [0x0 signalLabelDefinition]
      Tag: ""
      Description: "Whale type"
```

Use `labeledSignalSet` to create a labeled signal set.

```
types = types.Categories
```

```
types = 3x1 string array
    "blue"
    "humpback"
    "white"
```

Modify the label to incorporate sperm whales and killer whales. Verify that the labeled signal set includes the two new whale types.

```
editLabelDefinition(lss, 'WhaleType', ...
    'Categories', {'sperm', 'killer'})
```

```
lbldefs = getLabelDefinitions(lss);
types = lbldefs(1).Categories
```

```
types = 5x1 string array
    "blue"
    "humpback"
    "white"
    "sperm"
    "killer"
```

The definition for trill regions has a sublabel that identifies peaks.

```
lbldefs(3).Sublabels
```

```
ans =
    signalLabelDefinition with properties:
        Name: "TrillPeaks"
        LabelType: "point"
        LabelDataType: "numeric"
        ValidationFunction: []
        PointLocationsDataType: "double"
        DefaultValue: []
        Sublabels: [0x0 signalLabelDefinition]
        Tag: ""
        Description: "Trill peaks"
```

Use `labeledSignalSet` to create a labeled signal set.

Change the description of the sublabel.

```
editLabelDefinition(lss, ["TrillRegions" "TrillPeaks"], 'Description', 'Peaks of trill re
```

```

lbldefs = getLabelDefinitions(lss);
lbldefs(3).Sublabels

ans =
    signalLabelDefinition with properties:
        Name: "TrillPeaks"
        LabelType: "point"
        LabelDataType: "numeric"
        ValidationFunction: []
        PointLocationsDataType: "double"
        DefaultValue: []
        Sublabels: [0x0 signalLabelDefinition]
        Tag: ""
        Description: "Peaks of trill regions"

```

Use `labeledSignalSet` to create a labeled signal set.

Input Arguments

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1),randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep",'LabelType','roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: {'Asleep' 'REM'} or ["Asleep" "REM"] specifies a region of a signal in which a patient undergoes REM sleep.

propname — Property name

'Name' | 'DefaultValue' | 'Tag' | 'Description' | 'Categories'

Property name, specified as 'Name', 'DefaultValue', 'Tag', 'Description', or 'Categories'.

Data Types: char | string

val — Property value

numeric value | logical value | character vector | string | vector of strings | cell array of character vectors

Label values, specified as a numeric or logical value, a character vector or string, a vector of strings, or a cell array of character vectors. `val` must be of the data type specified for `propname`.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

emd

Empirical mode decomposition

Syntax

```
[imf,residual] = emd(X)
[imf,residual,info] = emd(X)
[ ___ ] = emd( ___ ,Name,Value)

emd( ___ )
```

Description

`[imf,residual] = emd(X)` returns intrinsic mode functions `imf` and residual signal `residual` corresponding to the empirical mode decomposition of `X`. Use `emd` to decompose and simplify complicated signals into a finite number of intrinsic mode functions required to perform Hilbert-spectral analysis.

`[imf,residual,info] = emd(X)` returns additional information `info` on IMFs and residual signal for diagnostic purposes.

`[___] = emd(___ ,Name,Value)` estimates `emd` with additional options specified by one or more `Name,Value` pair arguments.

`emd(___)` plots the original signal, IMFs, and residual signal as subplots in the same figure.

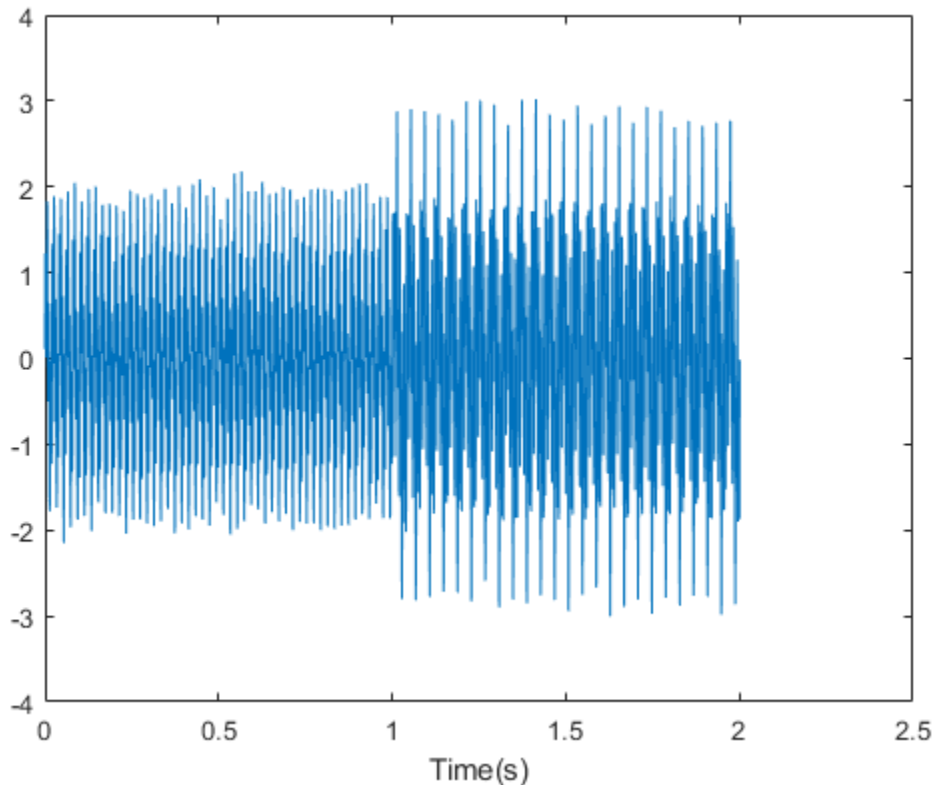
Examples

Perform Empirical Mode Decomposition and Visualize Hilbert Spectrum of Signal

For this example, consider a nonstationary continuous signal composed of sinusoidal waves with a distinct change in frequency. The vibration of a jackhammer and the sound of fireworks are examples of nonstationary continuous signals.

Load the nonstationary signal data with the sampling frequency `fs`, and visualize the mixed sinusoidal signal.

```
load('sinusoidalSignalExampleData.mat','X','fs')
t = (0:length(X)-1)/fs;
plot(t,X)
xlabel('Time(s)')
```



The mixed signal contains sinusoidal waves with different amplitude and frequency values.

To create the Hilbert spectrum plot, you need the intrinsic mode functions (IMFs) of the signal. Perform empirical mode decomposition to compute the IMFs and residuals of the signal. Since the signal is not smooth, specify 'pchip' as the Interpolation method.


```
[imf,residual,info] = emd(X,'Interpolation','pchip');
```

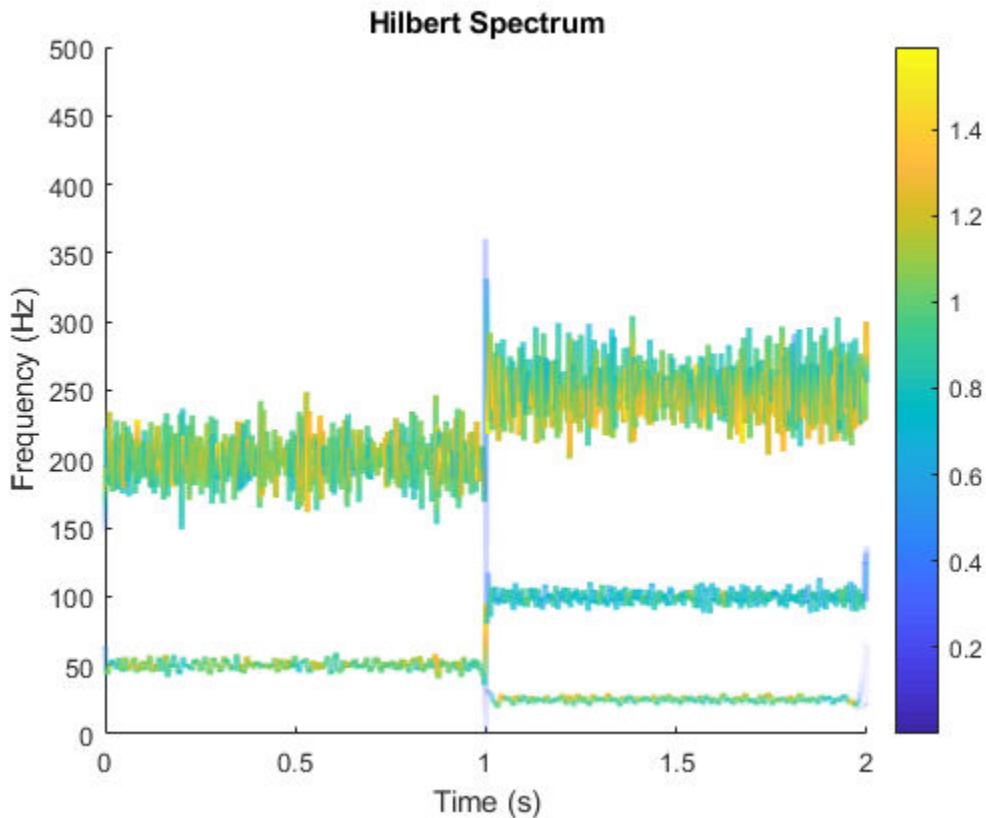
Current IMF	#Sift Iter	Relative Tol	Stop Criterion Hit
1	2	0.026352	SiftMaxRelativeTolerance
2	2	0.0039573	SiftMaxRelativeTolerance
3	1	0.024838	SiftMaxRelativeTolerance
4	2	0.05929	SiftMaxRelativeTolerance
5	2	0.11317	SiftMaxRelativeTolerance
6	2	0.12599	SiftMaxRelativeTolerance
7	2	0.13802	SiftMaxRelativeTolerance
8	3	0.15937	SiftMaxRelativeTolerance
9	2	0.15923	SiftMaxRelativeTolerance

The decomposition stopped because the number of extrema of the residual signal is less

The table generated in the command window indicates the number of sift iterations, the relative tolerance, and the sift stop criterion for each generated IMF. This information is also contained in `info`. You can hide the table by adding the `'Display',0` name value pair.

Create the Hilbert spectrum plot using the `imf` components obtained using empirical mode decomposition.

```
hht(imf,fs)
```



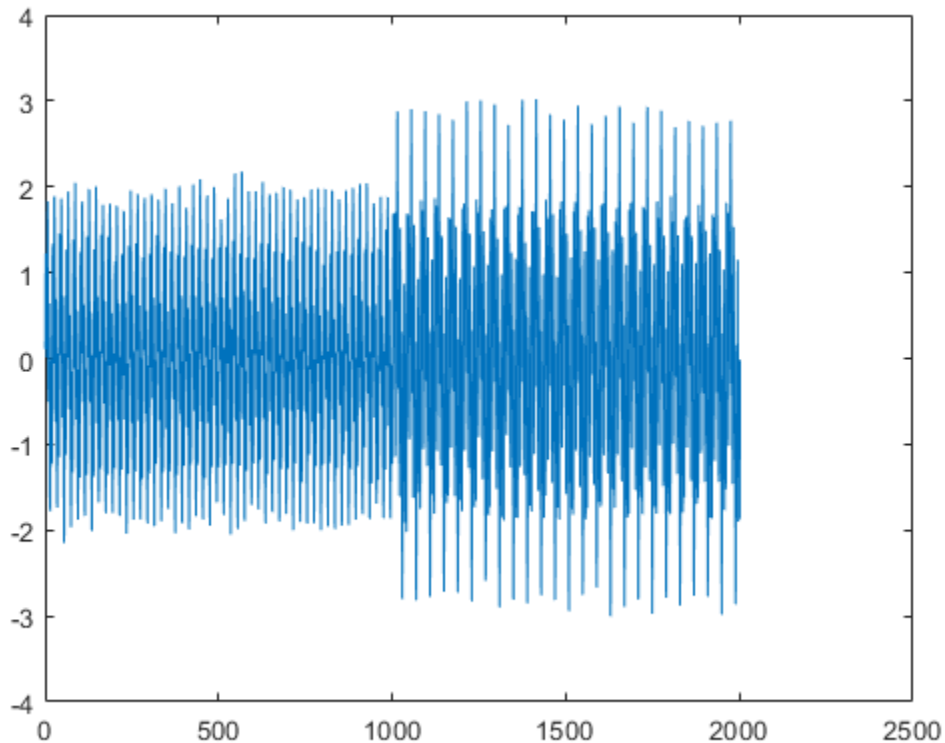
The frequency versus time plot is a sparse plot with a vertical color bar indicating the instantaneous energy at each point in the IMF. The plot represents the instantaneous frequency spectrum of each component decomposed from the original mixed signal. Three IMFs appear in the plot with a distinct change in frequency at 1 second.

Visualize Residual and Intrinsic Mode Functions of Signal

For this example, consider a nonstationary continuous signal composed of sinusoidal waves with a distinct change in frequency. The vibration of a jackhammer, or the sound of fireworks are examples of nonstationary continuous signals.

Load the nonstationary signal data, and visualize the mixed sinusoidal signal.

```
load('sinusoidalSignalExampleData.mat', 'X', 'fs');  
plot(X);
```



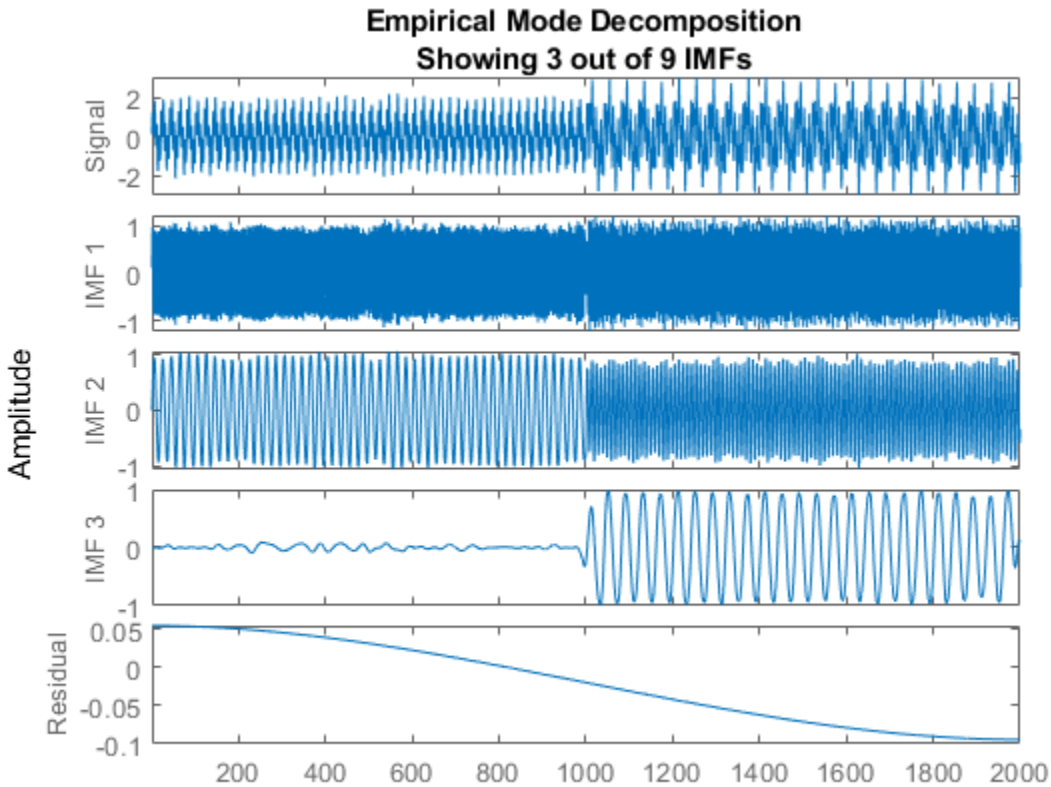
Observe that the mixed signal contains sinusoidal waves with different amplitude and frequency values.

Perform empirical mode decomposition to plot the intrinsic mode functions and residual of the signal. Since the signal is not smooth, specify 'pchip' as the Interpolation method.

```
emd(X, 'Interpolation', 'pchip');
```

Current IMF	#Sift Iter	Relative Tol	Stop Criterion Hit
1	2	0.026352	SiftMaxRelativeTolerance
2	2	0.0039573	SiftMaxRelativeTolerance
3	1	0.024838	SiftMaxRelativeTolerance
4	2	0.05929	SiftMaxRelativeTolerance
5	2	0.11317	SiftMaxRelativeTolerance
6	2	0.12599	SiftMaxRelativeTolerance
7	2	0.13802	SiftMaxRelativeTolerance
8	3	0.15937	SiftMaxRelativeTolerance
9	2	0.15923	SiftMaxRelativeTolerance

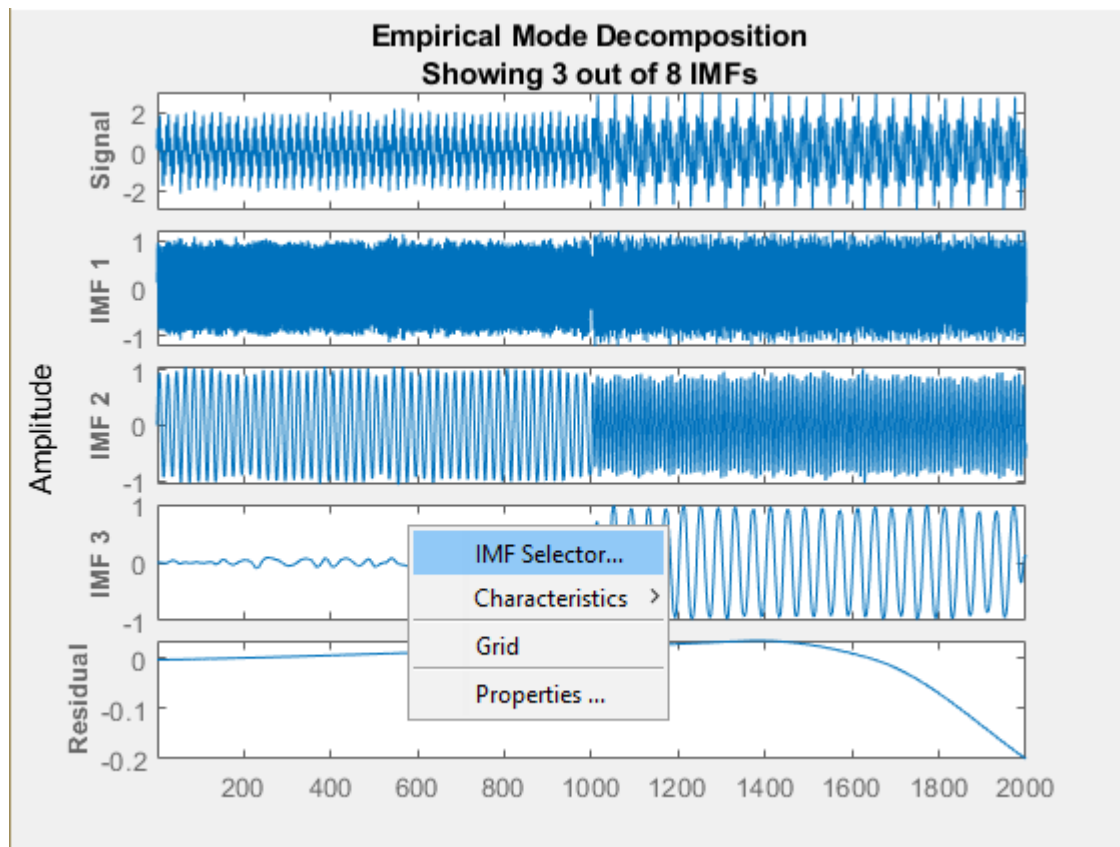
The decomposition stopped because the number of extrema of the residual signal is less



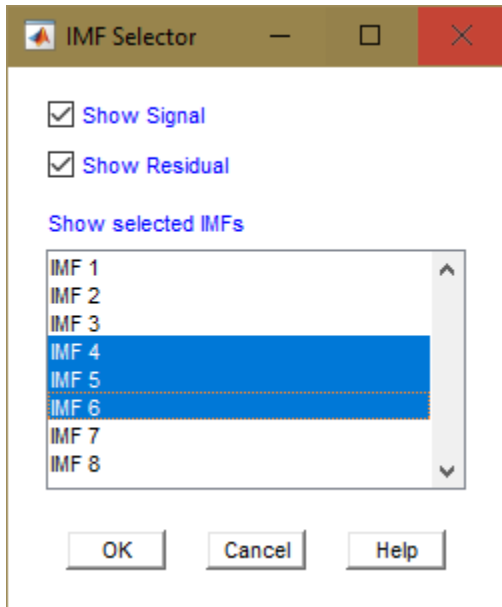
An interactive plot with the original signal, the first 3 IMFs, and the residual is generated. The table generated in the command window indicates the number of sift iterations, the

relative tolerance, and the sift stop criterion for each generated IMF. You can hide the table by specifying `Display` as `0`.

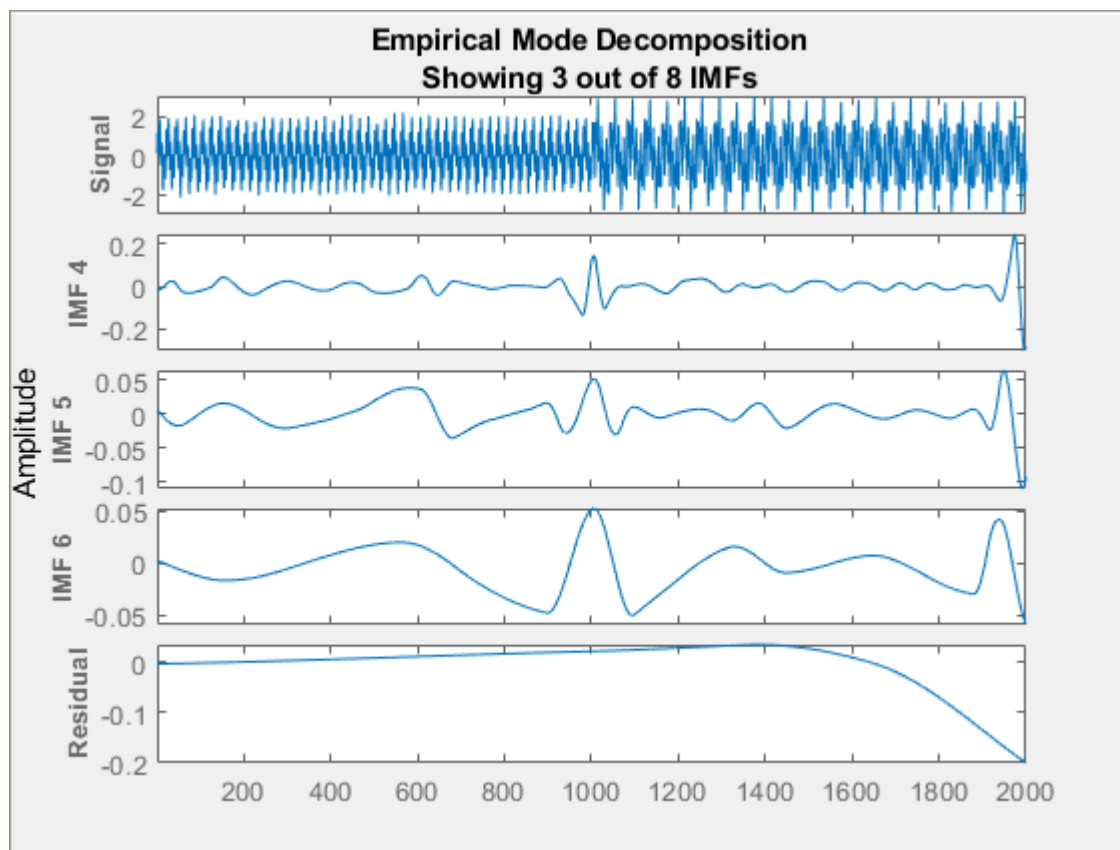
Right-click on the white space in the plot to open the **IMF selector** window. Use **IMF selector** to selectively view the generated IMFs, the original signal, and the residual.



Select the IMFs to be displayed from the list. Choose whether to display the original signal and residual on the plot.



The selected IMFs are now displayed on the plot.



Use the plot to visualize individual components decomposed from the original signal along with the residual. Note that the residual is computed for the total number of IMFs, and does not change based on the IMFs selected in the **IMF selector** window.

Input Arguments

X — Uniformly sampled time-domain signal

vector | single data column timetable

Uniformly sampled time-domain signal, specified as either a vector or single data column timetable.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

Example: `'MaxNumIMF', 5`

SiftRelativeTolerance — Cauchy type convergence criterion

0.2 (default) | positive scalar

Cauchy type convergence criterion, specified as the comma-separated pair consisting of `'SiftRelativeTolerance'` and a positive scalar. `SiftRelativeTolerance` is one of the sifting stop criteria, that is, sifting stops when current relative tolerance is less than `SiftRelativeTolerance`.

SiftMaxIterations — Maximum number of sifting iterations

100 (default) | positive scalar integer

Maximum number of sifting iterations, specified as the comma-separated pair consisting of `'SiftMaxIterations'` and a positive scalar integer. `SiftMaxIterations` is one of the sifting stop criteria, that is, sifting stops when the current number of iterations is larger than `SiftMaxIterations`.

`SiftMaxIterations` can be specified using only positive whole numbers.

MaxNumIMF — Maximum number of IMFs extracted

10 (default) | positive scalar integer

Maximum number of IMFs extracted, specified as the comma-separated pair consisting of `'MaxNumIMF'` and a positive scalar integer. `MaxNumIMF` is one of the decomposition stop criteria, that is, decomposition stops when number of IMFs generated is equal to `MaxNumIMF`.

`MaxNumIMF` can be specified using only positive whole numbers.

MaxNumExtrema — Maximum number of extrema in the residual signal

1 (default) | positive scalar integer

Maximum number of extrema in the residual signal, specified as the comma-separated pair consisting of `'MaxNumExtrema'` and a positive scalar integer. `MaxNumExtrema` is one

of the decomposition stop criteria, that is, decomposition stops when number of extrema is less than `MaxNumExtrema`.

`MaxNumExtrema` can be specified using only positive whole numbers.

MaxEnergyRatio — Signal to residual energy ratio

20 (default) | scalar

Signal to residual energy ratio, specified as the comma-separated pair consisting of `'MaxEnergyRatio'` and a scalar. `MaxEnergyRatio` is the ratio of the energy of the signal at the beginning of sifting and the average envelope energy. `MaxEnergyRatio` is one of the decomposition stop criteria, that is, decomposition stops when current energy ratio is larger than `MaxEnergyRatio`.

Interpolation — Interpolation method for envelope construction

'spline' (default) | 'pchip' | string

Interpolation method for envelope construction, specified as the comma-separated pair consisting of `'Interpolation'` and either `'spline'` or `'pchip'`.

Specify `Interpolation` as:

- `'spline'`, if `X` is a smooth signal
- `'pchip'`, if `X` is a nonsmooth signal

`'spline'` interpolation method uses cubic spline, while `'pchip'` uses piecewise cubic Hermite interpolating polynomial method.

Display — Toggle information display in the command window

1 (default) | 0 | scalar

Toggle information display in the command window, specified as the comma-separated pair consisting of `'Display'` and either 1 or 0. The table generated in the command window indicates the number of sift iterations, the relative tolerance, and the sift stop criterion for each generated IMF. Specify `Display` as 1 to show the table or 0 to hide the table.

Output Arguments

imf — Intrinsic mode function

matrix | timetable

Intrinsic mode function, returned as a matrix or timetable. `imf` is any function whose envelope is symmetric with respect to zero and whose numbers of extrema and zero crossings differ by at most one. Use `imf` to apply Hilbert-Huang transform to perform spectral analysis on the signal.

`imf` is returned as:

- A matrix whose each column is an `imf`, when `X` is a vector
- A timetable, when `X` is a single data column timetable

residual — Residual of the signal

column vector | single data column timetable

Residual of the signal, returned as a column vector or a single data column timetable. `residual` represents the portion of the original signal `X` not decomposed by `emd`.

`residual` is returned as:

- A column vector, when `X` is a vector.
- A single data column timetable, when `X` is a single data column timetable.

info — Additional information for diagnostics

structure

Additional information for diagnostics, returned as a structure with the following fields:

- `NumIMF` - Number of IMFs extracted from the signal
- `NumExtrema` - Number of extrema in each IMF
- `NumZeroCrossing` - Number of zero crossings for each IMF
- `NumSifting` - Number of siftings performed for each IMF
- `MeanEnvelopeEnergy` - Energy of the mean of upper and lower envelope obtained in each IMF computation
- `RelativeTolerance` - Relative tolerance attained for each IMF

Algorithms

Empirical Mode Decomposition

emd decomposes a signal $X(t)$ into k number of intrinsic mode functions (IMF), and residual $r_k(t)$ using the sifting process. A brief overview of the sifting process, listed in [1] and [2], is as follows:

- 1 Find local maxima and minima for signal $X(t)$ to construct an upper envelope $s_+(t)$, and a lower envelope $s_-(t)$.
- 2 Compute mean envelope for i th iteration, $m_{k,i}(t)$,

$$m_{k,i}(t) = \frac{1}{2}[s_+(t) + s_-(t)]$$

- 3 With $c_k(t) = X(t)$ for the first iteration, subtract mean envelope from residual signal,

$$c_k(t) = c_k(t) - m_{k,i}(t)$$

If $c_k(t)$ does not match the criteria of an IMF, steps 4 and 5 are skipped. The procedure is iterated again at step 1 with the new value of $c_k(t)$.

- 4 If $c_k(t)$ matches the criteria of an IMF, a new residual is computed. To update the residual signal, subtract the k th IMF from the previous residual signal,

$$r_k(t) = r_{k-1}(t) - c_k(t)$$

- 5 Then begin from step 1, using the residual obtained as a new signal $r_k(t)$, and store $c_k(t)$ as an intrinsic mode function.

For N intrinsic mode functions, the original signal is represented as,

$$X(t) = \sum_{i=1}^N c_i(t) + r_N(t)$$

For more information about the sifting process, see [1] and [2].

SiftRelativeTolerance

SiftRelativeTolerance is a Cauchy type stop criterion proposed in [4]. Sifting stops when current relative tolerance is less than **SiftRelativeTolerance**. The current relative tolerance is defined as

$$\text{RelativeTolerance} \triangleq \frac{\|c(t)_{\text{previous}} - c(t)_{\text{current}}\|^2}{\|c(t)_{\text{current}}\|^2}.$$

Because the Cauchy criterion does not directly count the number of zero crossings and local extrema, it is possible that the IMFs returned by the decomposition do not satisfy

the strict definition of an intrinsic mode function. In those cases, you can try reducing the value of the `SiftRelativeTolerance` from its default value. See [4] for a detailed discussion of stopping criteria. The reference also discusses the advantages and disadvantages of insisting on strictly defined IMFs in empirical mode decomposition.

MaxEnergyRatio

Energy ratio is the ratio of the energy of the signal at the beginning of sifting and the average envelope energy [3]. Decomposition stops when current energy ratio is larger than `MaxEnergyRatio`. For k IMFs, `EnergyRatio` is defined as

$$\text{EnergyRatio} \triangleq 10\log_{10}\left(\frac{\|X(t)\|^2}{\|r_k(t)\|^2}\right).$$

References

- [1] Huang, Norden E., Zheng Shen, Steven R. Long, Manli C. Wu, Hsing H. Shih, Qunan Zheng, Nai-Chyuan Yen, Chi Chao Tung, and Henry H. Liu. "The empirical mode decomposition and the Hilbert spectrum for nonlinear and non-stationary time series analysis." *Proceedings of the Royal Society of London. Series A: Mathematical, Physical and Engineering Sciences*. Vol. 454, 1998, pp. 903-995.
- [2] Rilling, G., Patrick Flandrin, and Paulo Gonçalves. "On empirical mode decomposition and its algorithms." *IEEE-EURASIP Workshop on Nonlinear Signal and Image Processing* 2003. NSIP-03. Grado, Italy. 8-11.
- [3] Rato, R.T., Manuel Ortigueira, and Arnaldo Batista. "On the HHT, its problems, and some solutions." *Mechanical Systems and Signal Processing* Vol. 22, 2008, pp. 1374-1394.
- [4] Wang, Gang, Xian-Yao Chen, Fang-Li Qiao, Zhaohua Wu, and Norden Huang. "On Intrinsic Mode Function." *Advances in Adaptive Data Analysis*. Vol. 2, Number 3, 2010, pp. 277-293.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

hht

Introduced in R2018a

entrupd

Entropy update (wavelet packet)

Syntax

```
T = entrupd(T,ENT)
T = entrupd(T,ENT,PAR)
```

Description

entrupd is a one- or two-dimensional wavelet packet utility.

T = entrupd(T,ENT) or T = entrupd(T,ENT,PAR) returns for a given wavelet packet tree T, the updated tree using the entropy function ENT with the optional parameter PAR (see wenergy for more information).

Examples

```
% The current extension mode is zero-padding (see dwtmode).

% Load signal.
load noisdopp; x = noisdopp;

% Decompose x at depth 2 with db1 wavelet packets
% using shannon entropy.
t = wptdec(x,2,'db1','shannon');

% Read entropy of all the nodes.
nodes = allnodes(t);
ent = read(t,'ent',nodes);
ent'
ent =
    1.0e+04 *
    -5.8615 -6.8204 -0.0350 -7.7901 -0.0497 -0.0205 -0.0138

% Update nodes entropy.
```

```
t = entrupe(t, 'threshold', 0.5);
nent = read(t, 'ent');
nent'
```

937 488 320 241 175 170 163

See Also

wenergy | wpdec | wpdec2

Introduced before R2006a

featureMatrix

Scattering feature matrix

Syntax

```
smat = featureMatrix(sf,x)
smat = featureMatrix(sf,s)
smat = featureMatrix( ____,Name,Value)
```

Description

`smat = featureMatrix(sf,x)` returns the scattering features for the scattering decomposition framework `sf` and the input signal `x`. `x` is a real-valued vector or matrix. If `x` is a vector, `smat` is an M -by- N matrix, where M is the number of resolutions across all orders of the scattering transform, and N is the resolution of the scattering coefficients. If `x` is a matrix, `smat` is an M -by- N -by- P array, where P is the number of columns in `x`.

The precision of `smat` depends on the precision specified in the framework `sf`.

`smat = featureMatrix(sf,s)` returns the scattering feature matrix for the cell array of scattering coefficients `s`. `s` is the output of `scatteringTransform` applied to an input signal for the scattering decomposition framework `sf`.

`smat = featureMatrix(____,Name,Value)` returns the scattering feature matrix with additional options specified by one or more `Name,Value` pair arguments.

Examples

Obtain Scattering Feature Matrix

This example shows how to obtain the scattering feature matrix for a scattering decomposition framework and how to compare the matrix with scattering coefficients.

Load an ECG signal sampled at 180 Hz. Create a scattering decomposition framework that can be used with the signal.

```
load wecg
Fs = 180;
sf = waveletScattering('SignalLength', numel(wecg), ...
    'SamplingFrequency', Fs);
```

Calculate the scattering feature matrix using the log transformation. Display the dimensions of the matrix.

```
smat = featureMatrix(sf, wecg, 'Transform', 'Log');
size(smat)
```

```
ans = 1x2
    154     8
```

Now calculate the scattering transform of the signal. Obtain the scattering coefficients. The output is a cell array with three elements. Each element is a table. Confirm the total number of rows in the tables is equal to the number of rows in the matrix.

```
S = scatteringTransform(sf, wecg);
t1rows = size(S{1},1);
t2rows = size(S{2},1);
t3rows = size(S{3},1);
disp(['Total Number of Rows: ', num2str(t1rows+t2rows+t3rows)])
```

```
Total Number of Rows: 154
```

Display the base-2 log resolution of the zeroth-order scattering coefficients.

```
disp(['Resolution: ', num2str(S{1}.resolution(1))])
```

```
Resolution: -8
```

Obtain the natural logarithm of the zeroth-order scattering coefficients. Compare the scattering coefficients with the first row in the feature matrix. The number of coefficients in each equals the absolute value of the base-2 log resolution.

```
logS = log(sf, S);
logScat = logS{1}.signals{1};
[smat(1,:) ' logScat]
```

```
ans = 8x2

-1.2914 -1.2914
-2.4682 -2.4682
-1.6368 -1.6368
-1.2716 -1.2716
-1.6818 -1.6818
-4.3701 -4.3701
-1.3199 -1.3199
-1.0542 -1.0542
```

Input Arguments

sf — Scattering decomposition framework

waveletScattering object

Scattering decomposition framework, specified as a `waveletScattering` object.

x — Input data

real-valued vector | real-valued matrix

Input data, specified as a real-valued vector or matrix. If `x` is a vector, the number of samples in `x` must equal the `SignalLength` value of `sf`. If `x` is a matrix, the number of rows in `x` must equal the `SignalLength` value of `sf`.

Data Types: `double`

s — Scattering coefficients

cell array

Scattering coefficients, specified as a cell array. `s` is obtained from the scattering transform of the scattering decomposition framework `sf`. For more information, see `scatteringTransform`.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

```
Example: smat =  
featureMatrix(sf,x,'Transform','log','Normalization','parent')
```

Normalization — Type of normalization

'none' (default) | 'parent'

Type of normalization to apply to the scattering coefficients, specified as 'none' or 'parent'. If specified as 'parent', scattering coefficients of order greater than 0 are normalized by their parents along the scattering path.

Transform — Type of transformation

'none' (default) | 'log'

Type of transformation to apply to the scattering coefficients, specified as 'none' or 'log'.

Output Arguments

smat — Scattering features

real-valued matrix | real-valued array

Scattering features for the scattering decomposition framework, `sf`, returned as a real-valued matrix or array. If `x` is a vector, `smat` is an M -by- N matrix, where M is the number of resolutions across all orders of the scattering transform and N is the resolution of the scattering coefficients. If `x` is a matrix, `smat` is an M -by- N -by- P array, where P is the number of columns in `x`.

The precision of `smat` depends on the precision specified in the framework `sf`.

See Also

`scatteringTransform` | `waveletScattering`

Introduced in R2018b

featureMatrix

Image scattering feature matrix

Syntax

```
smat = featureMatrix(sf,im)
smat = featureMatrix(sf,sc)
smat = featureMatrix( ____, 'Transform', transformtype)
```

Description

`smat = featureMatrix(sf,im)` returns the scattering feature matrix for the image scattering decomposition framework, `sf`, and the input image, `im`. `im` is a real-valued 2-D (M -by- N) or 3-D matrix (M -by- N -by-3). If `im` is a 3-D matrix, the size of the third dimension must be 3. If `im` is a 2-D matrix, `smat` is Np -by- Ms -by- Ns , where Np is the number of scattering paths and Ms -by- Ns is the resolution of the scattering coefficients. If `im` is a 3-D matrix, `smat` is Np -by- Ms -by- Ns -by-3.

`smat = featureMatrix(sf,sc)` returns the scattering feature matrix for the cell array of scattering coefficients, `sc`. `sc` is obtained from the `scatteringTransform` method of the image scattering decomposition framework.

`smat = featureMatrix(____, 'Transform', transformtype)` applies the transformation specified by `transformtype` to the scattering coefficients. Valid options for `transformtype` are 'log' and 'none'. If unspecified, `transformtype` defaults to 'none'. You can use this syntax with any of the previous syntaxes.

Examples

Obtain Feature Matrix for Wavelet Image Scattering Framework

This example shows how to obtain the feature matrix for a wavelet image scattering framework.

Load the xbox image. Create an image scattering framework suitable for the image.

```
load xbox
sf = waveletScattering2('ImageSize',size(xbox))

sf =
    waveletScattering2 with properties:
        ImageSize: [128 128]
        InvarianceScale: 64
        NumRotations: [6 6]
        QualityFactors: [1 1]
        Precision: "single"
        OversamplingFactor: 0
        OptimizePath: 1
```

Obtain the feature matrix.

```
smat = featureMatrix(sf,xbox);
```

Input Arguments

sf — Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

im — Input image

real-valued matrix

Input image, specified as real-valued 2-D matrix or 3-D matrix. If im is 3-D, im is assumed to be a color image in the RGB color space, and the size of the third dimension must equal 3. The row and column sizes of im must match the ImageSize property of sf.

sc — Scattering coefficients

cell array

Scattering coefficients, specified as a cell array. sc is obtained from the scatteringTransform method of the image scattering decomposition framework.

transformtype — Transformation

'none' (default) | 'log'

Transformation to apply to the scattering coefficients:

- 'none': No transformation is applied to the scattering coefficients.
- 'log': The natural logarithm is applied to the scattering coefficients.

Output Arguments

smat — Scattering feature matrix

real-valued array

Scattering feature matrix for the 2-D scattering decomposition framework `sf`, returned as a real-valued array. If `im` is a 2-D matrix, `smat` is Np -by- Ms -by- Ns , where Np is the number of scattering paths and Ms -by- Ns is the resolution of the scattering coefficients. If `im` is a 3-D matrix, `smat` is Np - Ms -by- Ns -by-3.

See Also

`scatteringTransform` | `waveletScattering2`

Introduced in R2019a

fbspwavf

Complex frequency B-spline wavelet

Syntax

```
[PSI,X] = fbspwavf(LB,UB,N,M,FB,FC)
```

Description

[PSI,X] = fbspwavf(LB,UB,N,M,FB,FC) returns values of the complex frequency B-Spline wavelet defined by the order parameter M (M is an integer such that $1 \leq M$), a bandwidth parameter FB , and a wavelet center frequency FC .

The function PSI is computed using the explicit expression

$$\text{PSI}(X) = (FB^{0.5}) * (\text{sinc}(FB*X/M).^M) .* \exp(2*i*pi*FC*X)$$

on an N point regular grid in the interval [LB,UB].

FB and FC must be such that $FC > 0$ and $FB > 0$.

Output arguments are the wavelet function PSI computed on the grid X .

Examples

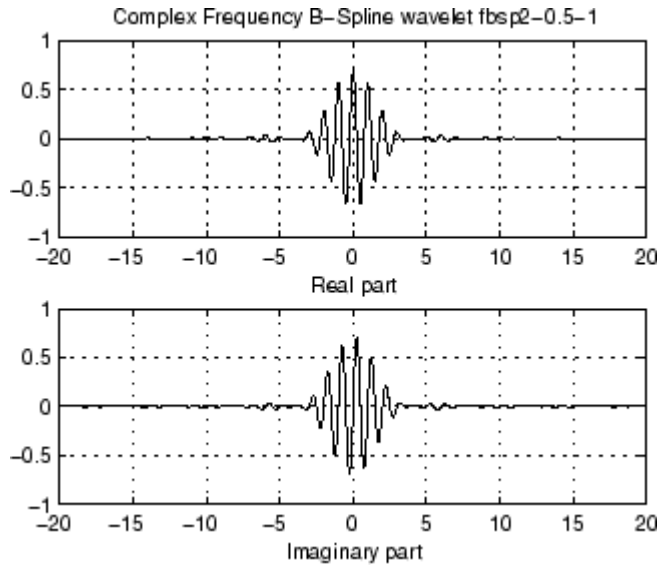
```
% Set order, bandwidth and center frequency parameters.
m = 2; fb = 0.5; fc = 1;

% Set effective support and grid parameters.
lb = -20; ub = 20; n = 1000;

% Compute complex Frequency B-Spline wavelet fbsp2-0.5-1.
[psi,x] = fbspwavf(lb,ub,n,m,fb,fc);

% Plot complex Frequency B-Spline wavelet.
subplot(211)
```

```
plot(x,real(psi))
title('Complex Frequency B-Spline wavelet fbsp2-0.5-1')
xlabel('Real part'), grid
subplot(212)
plot(x,imag(psi))
xlabel('Imaginary part'), grid
```



References

Teolis, A. (1998), *Computational signal processing with wavelets*, Birkhauser, p. 63.

See Also

waveinfo

Introduced before R2006a

fejerkorovkin

Fejér-Korovkin wavelet filters

Syntax

`Lo = fejerkorovkin(wname)`

Description

`Lo = fejerkorovkin(wname)` returns the Fejér-Korovkin scaling filter specified by `wname`. Valid entries for `wname` are `'fk4'`, `'fk6'`, `'fk8'`, `'fk14'`, `'fk18'`, and `'fk22'`. For information on the Fejér-Korovkin filters, see Nielson[1].

Examples

Fejer-Korovkin Filters

Construct and plot the Fejer-Korovkin (14) scaling function and wavelet.

Obtain the Fejer-Korovkin scaling filter and display its 14 coefficients.

```
Lo = fejerkorovkin('fk14')
```

```
Lo = 1×14
```

```
    0.2604    0.6869    0.6116    0.0514   -0.2456   -0.0486    0.1243    0.0222   -0.0
```

Use the scaling filter to obtain the wavelet filter and display its wavelet filter coefficients.

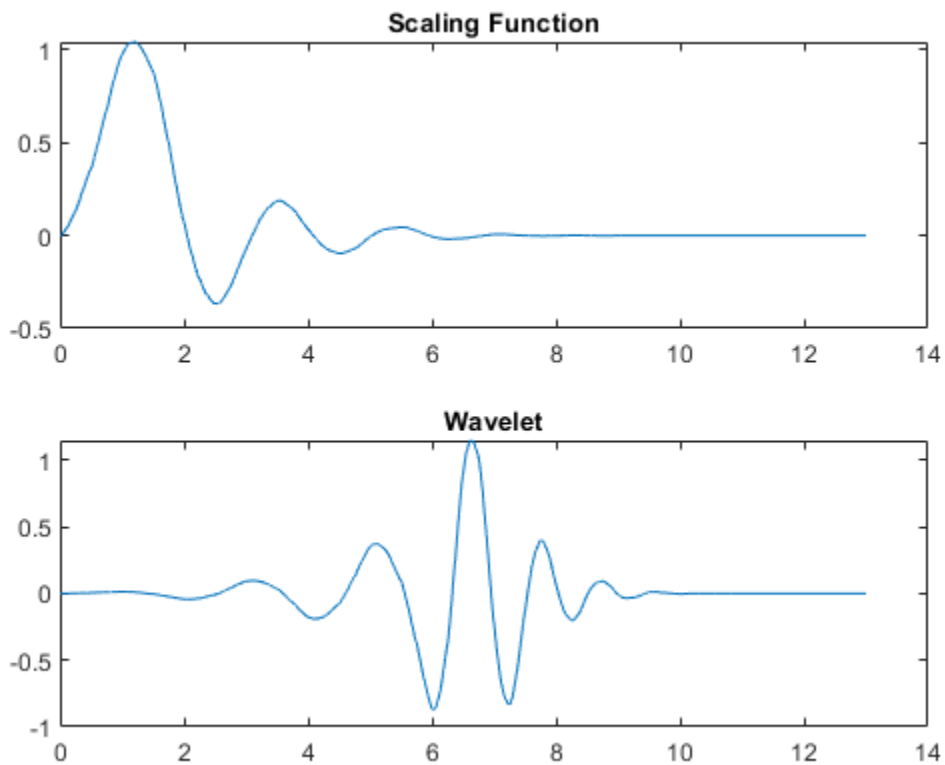
```
Hi = qmf(Lo)
```

```
Hi = 1×14
```

```
    0.0035    0.0093   -0.0033   -0.0298   -0.0051    0.0640    0.0222   -0.1243   -0.0
```

wavefun provides an efficient way to construct and plot the scaling function and wavelet.

```
[phi,psi,xval] = wavefun('fk14');  
subplot(2,1,1)  
plot(xval,phi)  
title('Scaling Function')  
subplot(2,1,2)  
plot(xval,psi)  
title('Wavelet')
```



Input Arguments

wname — Name

'fk4' | 'fk6' | 'fk8' | 'fk14' | 'fk18' | 'fk22'

Name of desired Fejér-Korovkin scaling filter. The numeric value in each name is the number of Fejér-Korovkin filter coefficients. For example, if wname is 'fk14', Lo has 14 coefficients.

Output Arguments

Lo — Scaling filter

vector

Scaling filter, returned as a vector.

References

- [1] Nielsen, M. "On the construction and frequency localization of finite orthogonal quadrature filters." *Journal of Approximation Theory*. Vol. 108, Number 1, 2001, pp. 36-52.

See Also

coifwavf | dbwavf | symwavf

Introduced in R2015b

filt2ls

Transform quadruplet of filters to lifting scheme

Syntax

```
LS = filt2ls(LoD,HiD,LoR,HiR)
```

Description

LS = filt2ls(LoD,HiD,LoR,HiR) returns the lifting scheme LS associated with the four input filters LoD, HiD, LoR, and HiR that verify the perfect reconstruction condition.

Examples

```
[LoD,HiD,LoR,HiR] = wfilters('db2')  
  
LoD =  
    -0.1294    0.2241    0.8365    0.4830  
  
HiD =  
    -0.4830    0.8365   -0.2241   -0.1294  
  
LoR =  
    0.4830    0.8365    0.2241   -0.1294  
  
HiR =  
    -0.1294   -0.2241    0.8365   -0.4830  
  
LS = filt2ls(LoD,HiD,LoR,HiR);  
displs(LS);  
  
LS = {...
```

```
'd'          [ -1.73205081] [0]
'p'          [ -0.06698730  0.43301270] [1]
'd'          [  1.00000000] [-1]
[  1.93185165] [  0.51763809] []
};
```

```
LSref = liftwave('db2');
displs(LSref);
```

```
LSref = {...
'd'          [ -1.73205081] [0]
'p'          [ -0.06698730  0.43301270] [1]
'd'          [  1.00000000] [-1]
[  1.93185165] [  0.51763809] []
};
```

See Also

ls2filt | lsinfo

Introduced before R2006a

filterbank

Wavelet time scattering filter banks

Syntax

```
filters = filterbank(sf)
[filters,f] = filterbank(sf)
[filters,f,filterparams] = filterbank(sf)
[___] = filterbank(sf,order)
```

Description

`filters = filterbank(sf)` returns the filter banks used in the computation of the scattering coefficients. `filters` is a cell array of structure arrays with *norder* elements, where *norder* is the number of scattering orders. The first element of `filters` contains the scaling filter, `phift`, used in the computation of the 0th-order scattering coefficients. Subsequent elements of `filters` contain the wavelet filters, `psift`, and scaling filter, `phift`, for the corresponding filter banks of the scattering decomposition.

The precision of `phift` and `psift` depends on the precision specified in the framework `sf`.

`[filters,f] = filterbank(sf)` returns the frequencies corresponding to the DFT bins in the `psift` and `phift` fields of `filters`. If you specify a sampling frequency in the construction of `sf`, `f` is measured in hertz. Otherwise, `f` is measured in cycles/sample.

`[filters,f,filterparams] = filterbank(sf)` returns the filter parameters for each element of `filters`. `filterparams` is a cell array with *norder* elements. Each element of `filterparams` is a MATLAB table.

`[___] = filterbank(sf,order)` returns the filter banks used to compute the specified `order` scattering coefficients. `order` is an integer between 0 and *nfilters* inclusive, where *nfilters* is the number of filter banks in the scattering decomposition inclusive. These input arguments can be used with any of the output syntaxes shown previously.

Examples

Plot Scattering Framework Filter Banks

Create a scattering decomposition framework for a signal sampled at 25 Hz.

```
sf = waveletScattering('SamplingFrequency',25)
```

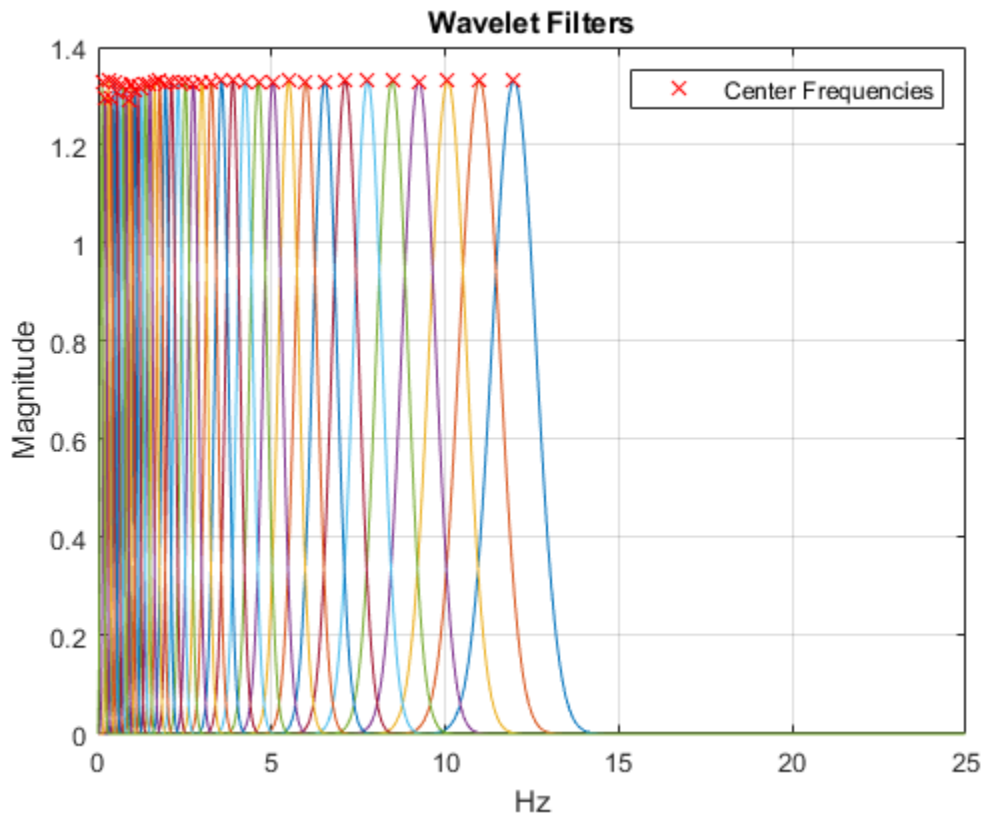
```
sf =
  waveletScattering with properties:
    SignalLength: 1024
    InvarianceScale: 20.4800
    QualityFactors: [8 1]
    Boundary: "periodic"
    SamplingFrequency: 25
    Precision: "double"
    OversamplingFactor: 0
```

Obtain the filter banks, DFT frequency bins, and filter bank parameters.

```
[filters,f,fparams] = filterbank(sf);
```

Plot the wavelet filters used in computing the first-order coefficients. Plot the wavelet center frequencies as well.

```
coefOrder = 1;
wvFilters = filters{coefOrder+1}.psift;
wvcenFrq = fparams{coefOrder+1}.omegapsi;
plot(f,wvFilters)
hold on
cf = plot(wvcenFrq,max(wvFilters),'rx');
grid on
title('Wavelet Filters')
xlabel('Hz')
ylabel('Magnitude')
legend(cf,'Center Frequencies')
```



Input Arguments

sf — Scattering decomposition framework

waveletScattering object

Scattering decomposition framework, specified as a `waveletScattering` object.

order — Order of scattering coefficients

positive integer

Order of scattering coefficients, specified as a positive integer between 0 and $nfilters$ inclusive, where $nfilters$ is the number of filter banks in the scattering decomposition `sf`.

Data Types: double

Output Arguments

filters — Filter banks

cell array

Filter banks used in the computation of the scattering coefficients, returned as a cell array of structure arrays. `filters` has *norder* elements, where *norder* is the number of scattering orders. The first element of `filters` is a structure with the single field `phift`. `phift` contains the scaling filter used in the computation of the 0th-order scattering coefficients. Subsequent elements of `filters` contain the wavelet filters, `psift`, and the scaling filter, `phift`, used for the corresponding filter banks of the scattering decomposition in the structure fields.

The precision of `phift` and `psift` depends on the precision specified in the framework `sf`.

f — Frequencies

real-valued vector

Frequencies corresponding to the DFT bins in the `psift` and `phift` fields of `filters`. If you specify a sampling frequency in the construction of `sf`, `f` is measured in hertz. Otherwise, `f` is measured in cycles/sample.

Data Types: double

filterparams — Filter bank parameters

cell array

Filter bank parameters for each element of `filters`, returned as a cell array. `filterparams` has *norder* elements, where *norder* is the number of scattering orders.

The first element of `filterparams` is a MATLAB table with the following variables:

- `boundary` — The signal extension used in the filters, returned as either `'periodic'` or `'reflection'`.
- `precision` — The precision used in the filters, returned as `'double'` or `'single'`.
- `sigmaphi` — The time standard deviation of the scaling function, returned as a scalar. If you specify a sampling frequency, `sigmaphi` is in seconds. Otherwise, `sigmaphi` is in samples.

- `freqsigmaphi` — The frequency standard deviation of the scaling function, returned as a scalar. If you specify a sampling frequency, `freqsigmaphi` is in hertz. Otherwise, `freqsigmaphi` is in cycles/sample.
- `phiftdsupport` — The frequency support of the scaling function, returned as a scalar. If you specify a sampling frequency, `phiftdsupport` is in hertz. Otherwise, `phiftdsupport` is in cycles/sample.
- `phi3dBbw` — The 3-dB bandwidth of the scaling function, returned as a scalar.

Subsequent elements of `filterparams` include additional variables for the wavelet parameters:

- `J` — The integer number of logarithmically spaced wavelet filters in the scattering filter bank.
- `omegappsi` — The center frequencies for the wavelet filters in descending order (highest to lowest), returned as a vector. The `omegappsi` variable includes the center frequencies for any linearly spaced filters.
- `freqsigmapsi` — The wavelet frequency standard deviations, returned as a vector.
- `timesigmapsi` — The wavelet time standard deviations, returned as a vector.
- `psi3dBbw` — The wavelet 3-dB bandwidths, returned as a vector.
- `psiftdsupport` — The wavelet frequency supports, returned as a vector.

See Also

`littlewoodPaleySum` | `waveletScattering`

Introduced in R2018b

filterbank

Wavelet and scaling filters

Syntax

```
phif = filterbank(sf)
[phif,psifilters] = filterbank(sf)
[phif,psifilters,f] = filterbank(sf)
[phif,psifilters,f,filterparams] = filterbank(sf)
[ ___ ] = filterbank(sf,fb)
```

Description

`phif = filterbank(sf)` returns the Fourier transform of the scaling filter for the 2-D wavelet scattering framework, `sf`. `phif` is a single or double-precision matrix depending on the value of the `Precision` property of the scattering framework. `phif` has dimensions M -by- N , where M and N are the padded row and column sizes of the scattering framework.

`[phif,psifilters] = filterbank(sf)` returns the Fourier transforms for the wavelet filters in `psifilters`. `psifilters` is an Nfb -by-1 cell array, where Nfb is the number of filter banks in the scattering framework. Each element of `psifilters` is a 3-D array. The 3-D arrays are M -by- N -by- L , where M and N are the padded row and column sizes of the wavelet filters and L is the number of wavelet filters for each filter bank. The wavelet filters are ordered by increasing scale with `NumRotations` wavelet filters for each scale. Within a scale, the wavelet filters are ordered by rotation angle.

`[phif,psifilters,f] = filterbank(sf)` returns the center spatial frequencies for the wavelet filters in `psifilters`. `f` is an Nfb -by-1 cell array, where Nfb is the number of filter banks in `sf`. The j th element of `f` contains the center frequencies for the j th wavelet filter bank in `psifilters`. Each element of `f` is a L -by-2 matrix with each row containing the center frequencies of the corresponding L th wavelet.

`[phif,psifilters,f,filterparams] = filterbank(sf)` returns the filter parameters for the 2-D scattering framework. `filterparams` is an Nfb -by-1 cell array of

MATLAB tables, where the j th element of `filterparams` is a MATLAB table containing the filter parameters for the j th filter bank

`[___] = filterbank(sf,fb)` returns the desired outputs for the filter banks specified in `fb`. `fb` is a scalar or vector of integers between 1 and `numfilterbanks(sf)` inclusive. If `fb` is a scalar, `psifilters` is an M -by- N -by- L matrix, and `filterparams` is a MATLAB table.

Examples

Plot Wavelet Center Frequencies

This example shows how to plot the scaling filter and the wavelet filter center frequencies for a two-filter bank wavelet image scattering framework.

Create a wavelet image scattering framework with two filter banks. The first filter bank has a quality factor of 1, and the second filter bank has a quality factor of 2.

```
sf = waveletScattering2('QualityFactors',[2 1])
```

```
sf =  
  waveletScattering2 with properties:
```

```
      ImageSize: [128 128]  
  InvarianceScale: 64  
    NumRotations: [6 6]  
  QualityFactors: [2 1]  
      Precision: "single"  
OversamplingFactor: 0  
    OptimizePath: 1
```

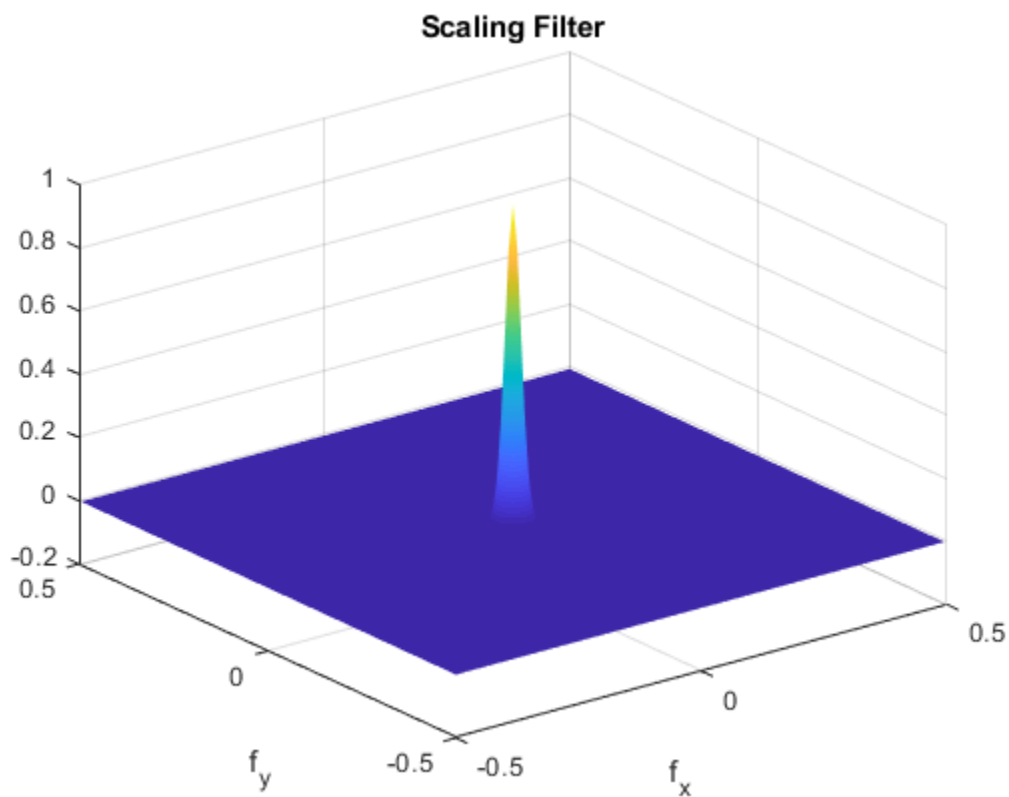
Obtain the scaling filter, wavelet filters, and wavelet center frequencies for the framework.

```
[phif,psifilters,f] = filterbank(sf);
```

Make a surface plot of the scaling filter.

```
Nx = size(phif,1);  
Ny = size(phif,2);
```

```
fx = -1/2:1/Nx:1/2-1/Nx;  
fy = -1/2:1/Ny:1/2-1/Ny;  
surf(fx,fy,fftshift(phif))  
shading interp  
title('Scaling Filter')  
xlabel('f_x')  
ylabel('f_y')
```



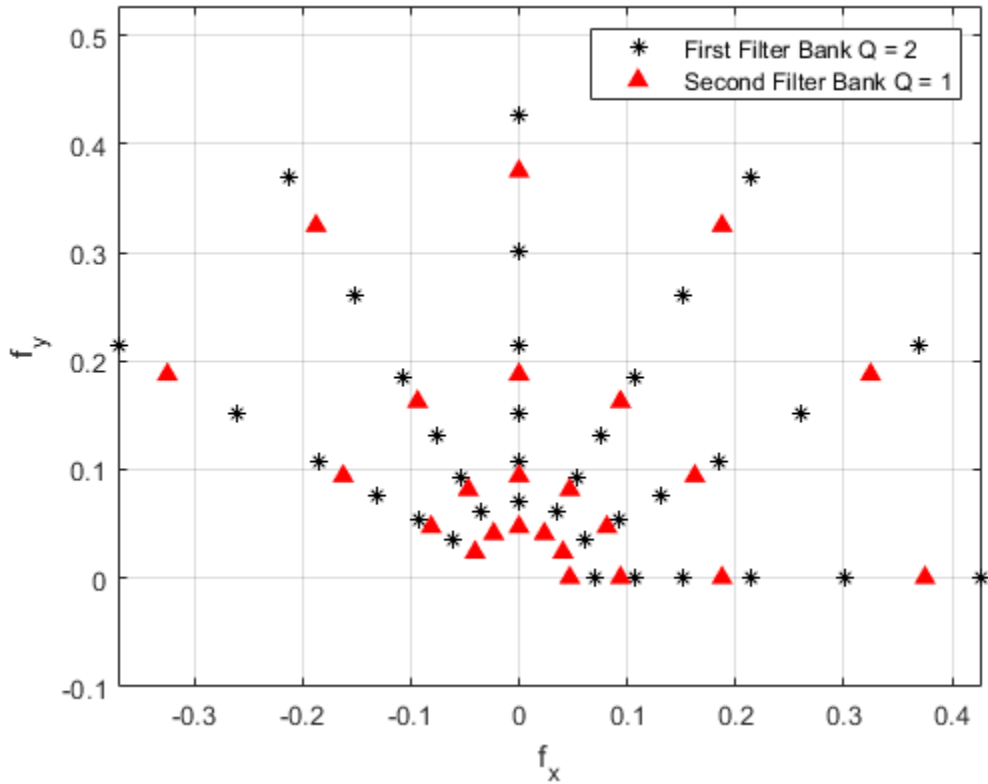
Plot the wavelet center frequencies for the two filter banks.

```
figure  
plot(f{1}(:,1),f{1}(:,2),'k*')  
hold on  
grid on
```

```

plot(f{2}(:,1),f{2}(:,2),'r^','MarkerFaceColor',[1 0 0])
axis equal
xlabel('f_x')
ylabel('f_y')
legend('First Filter Bank Q = 2','Second Filter Bank Q = 1')

```



Plot Specific Wavelet of Image Scattering Framework

This example shows how to obtain and plot a specific wavelet of a wavelet image scattering framework.

Create a wavelet image scattering framework with two filter banks. The first filter bank has a quality factor of 2 and seven rotations per wavelet. The second filter bank has a quality factor of 1 and five rotations per wavelet.

```
sf = waveletScattering2('QualityFactors',[2 1],'NumRotations',[7 5])
sf =
    waveletScattering2 with properties:
        ImageSize: [128 128]
        InvarianceScale: 64
        NumRotations: [7 5]
        QualityFactors: [2 1]
        Precision: "single"
        OversamplingFactor: 0
        OptimizePath: 1
```

Obtain the wavelet filters and center frequencies for the framework. Return the dimensions of the two cell arrays.

```
[~,psifilters,f] = filterbank(sf);
psifilters

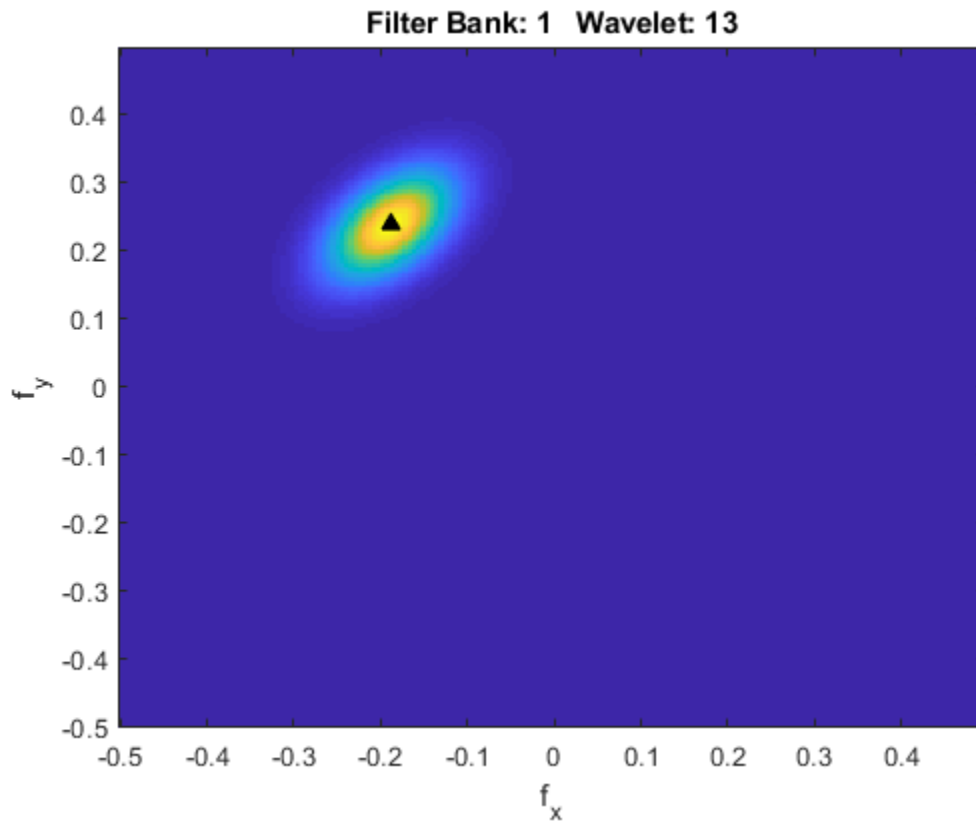
psifilters = 2x1 cell array
    {192x192x42 single}
    {192x192x20 single}

f

f = 2x1 cell array
    {42x2 double}
    {20x2 double}
```

The first filter bank has 42 wavelet filters, and the second filter bank has 20 filters. The number of filters in each filter bank is a multiple of the corresponding value in `NumRotations`. Use the helper function `helperPlotWavelet` to plot a specific wavelet and mark its center frequency.

```
figure
whichFilterBank = 1;
whichWavelet = 13;
helperPlotWavelet(psifilters,f,whichFilterBank,whichWavelet)
```



Appendix

The following helper function is used in this example.

```
function helperPlotWavelet(psiFilters,psiFreq,filBank,wvFilter)
Nx = size(psiFilters{filBank},2);
Ny = size(psiFilters{filBank},1);
fx = -1/2:1/Nx:1/2-1/Nx;
fy = -1/2:1/Ny:1/2-1/Ny;
imagesc(fx,fy,fftshift(psiFilters{filBank}(:,:,wvFilter)))
axis xy
hold on
xlabel('f_x')
ylabel('f_y')
```



```

plot(psiFreq{filBank}(wvFilter,1),psiFreq{filBank}(wvFilter,2),...
     'k^','markerfacecolor',[0 0 0])
str = sprintf('Filter Bank: %d Wavelet: %d',filBank,wvFilter);
title(str)
end

```

Determine Wavelet Semi-Major Axis

This example shows how to determine the semi-major axis of a wavelet filter in a 2-D wavelet scattering framework.

Create a 2-D wavelet scattering framework. The framework has two filter banks with quality factors of 2 and 1, respectively. There are seven rotations per wavelet in the first filter bank and five rotations per wavelet in the second filter bank. Return the Fourier transforms of the wavelet filters and their center spatial frequencies, and the filter bank parameters.

```

sf = waveletScattering2('QualityFactors',[2 1],'NumRotations',[7 5]);
[~,psif,f,fparams] = filterbank(sf);

```

The wavelet filters in `psif` are ordered by increasing scale, with `NumRotations` wavelet filters for each scale. Within a scale, the wavelet filters are ordered by rotation.

Return the reported 3 dB bandwidths, rotation angles, and slant parameter of the first filter bank. Return the dimensions of the matrix containing the wavelet filters of the first filter bank. Confirm that (number of rotations) \times (number of bandwidths) equals the size of the third dimension of the matrix. The product is the number of wavelet filters in the filter bank. The row and column sizes are the dimensions of the padded wavelet filters. Note that the slant parameter is less than 1.

```
fparams{1}.psi3dBbw
```

```
ans = 1×6
```

```
    0.1464    0.1036    0.0732    0.0518    0.0366    0.0366
```

```
fparams{1}.rotations
```

```
ans = 1×7
```

```
0 0.4488 0.8976 1.3464 1.7952 2.2440 2.6928
```

```
fparams{1}.slant
```

```
ans = 0.5817
```

```
size(psif{1})
```

```
ans = 1×3
```

```
192 192 42
```

The vector `fparams{1}.psi3dBbw` has six elements. The number of elements is equal to the number of wavelet scales in the first filter bank.

From the first filter bank, obtain the unrotated wavelet filter from the second finest scale. Obtain the center spatial frequency of the wavelet. Use the helper function `helperPlotWaveletFT` to plot the wavelet and mark its center frequency. The code for `helperPlotWaveletFT` is shown at the end of this example.

```
whichFilterBank = 1;
```

```
whichScale = 2;
```

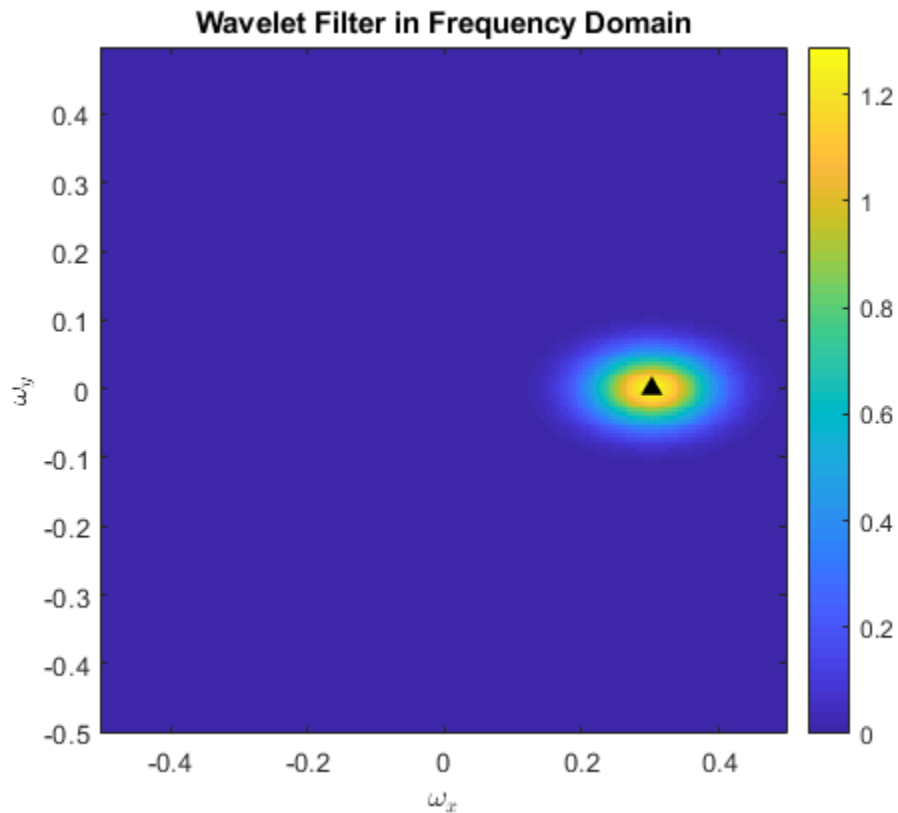
```
whichRotAngle = 1;
```

```
numRot = sf.NumRotations(whichFilterBank);
```

```
wvf = psif{whichFilterBank}(:, :, 1+(whichScale-1)*numRot+(whichRotAngle-1));
```

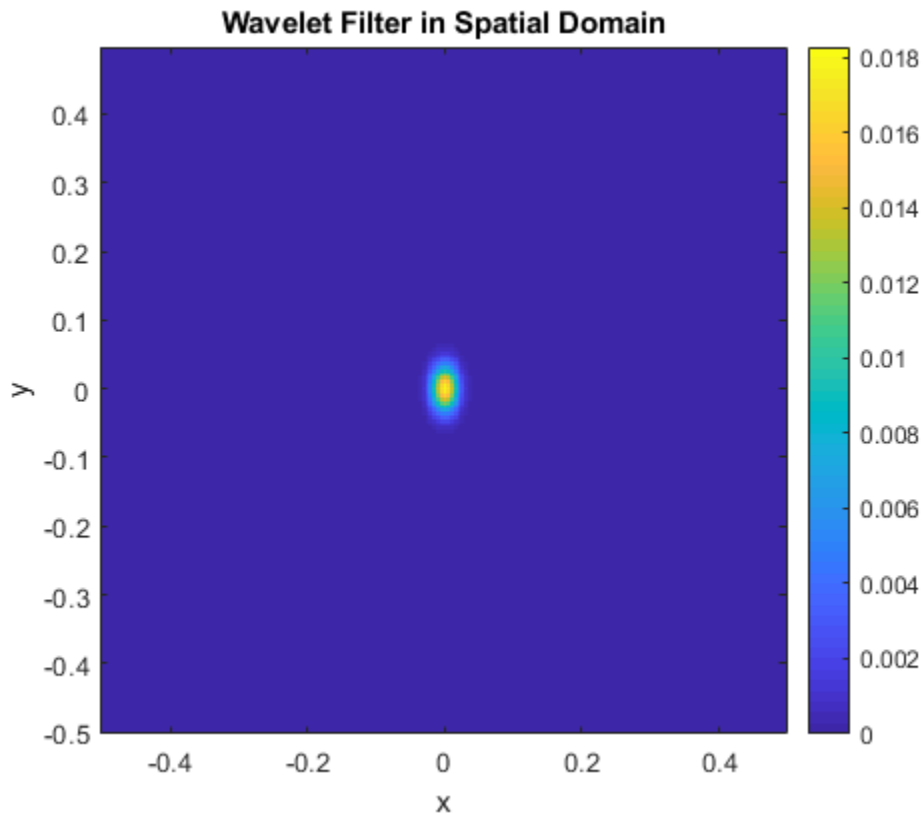
```
wvfCenFrq = f{whichFilterBank}(1+(whichScale-1)*numRot+(whichRotAngle-1), :);
```

```
helperPlotWaveletFT(wvf, wvfCenFrq)
```



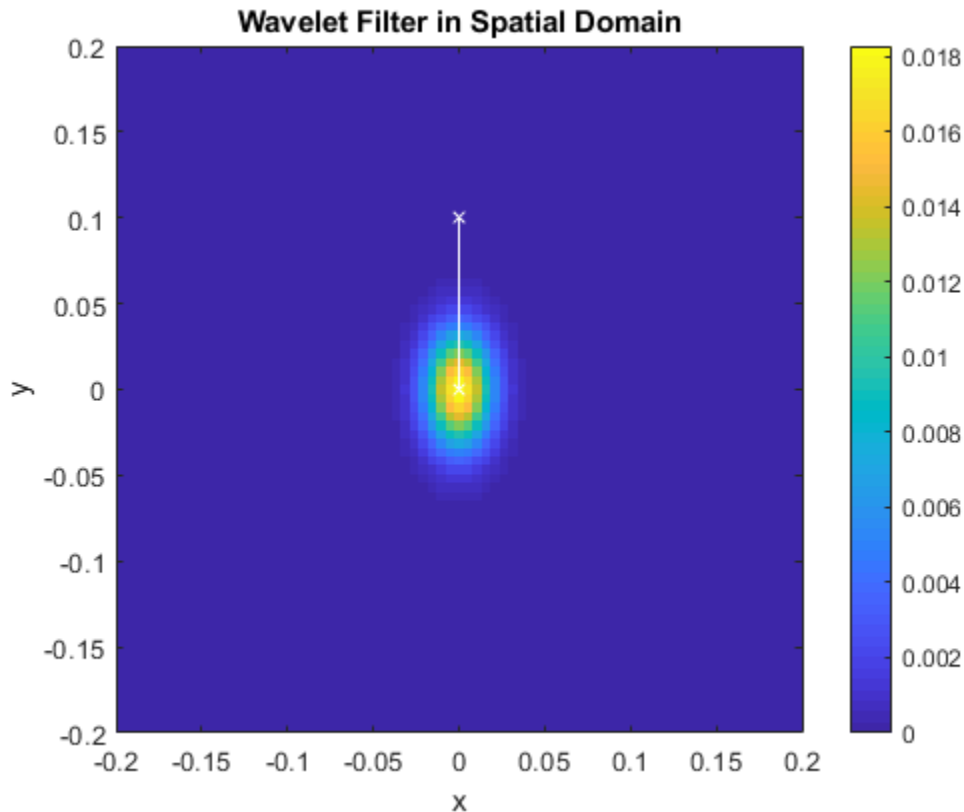
Take the inverse Fourier transform of the wavelet filter. The filter is strictly real, and the inverse Fourier transform is complex-valued. Use the helper function `helperPlotWavelet` to plot the absolute value of the wavelet. The code for `helperPlotWavelet` is shown at the end of this example.

```
wvf_ifft = ifft2(wvf);  
figure  
helperPlotWavelet(wvf_ifft)
```



Note that the semi-major axis of the wavelet support is in the y -direction. This is consistent with a slant parameter whose value is less than 1. The vector $(0, 0.1)$ coincides with the semi-major axis. Plot the vector in the previous figure.

```
vec = [0 0.1];  
hold on  
plot([0 vec(1)], [0 vec(2)], 'wx-')  
xlim([-0.2 0.2])  
ylim([-0.2 0.2])
```



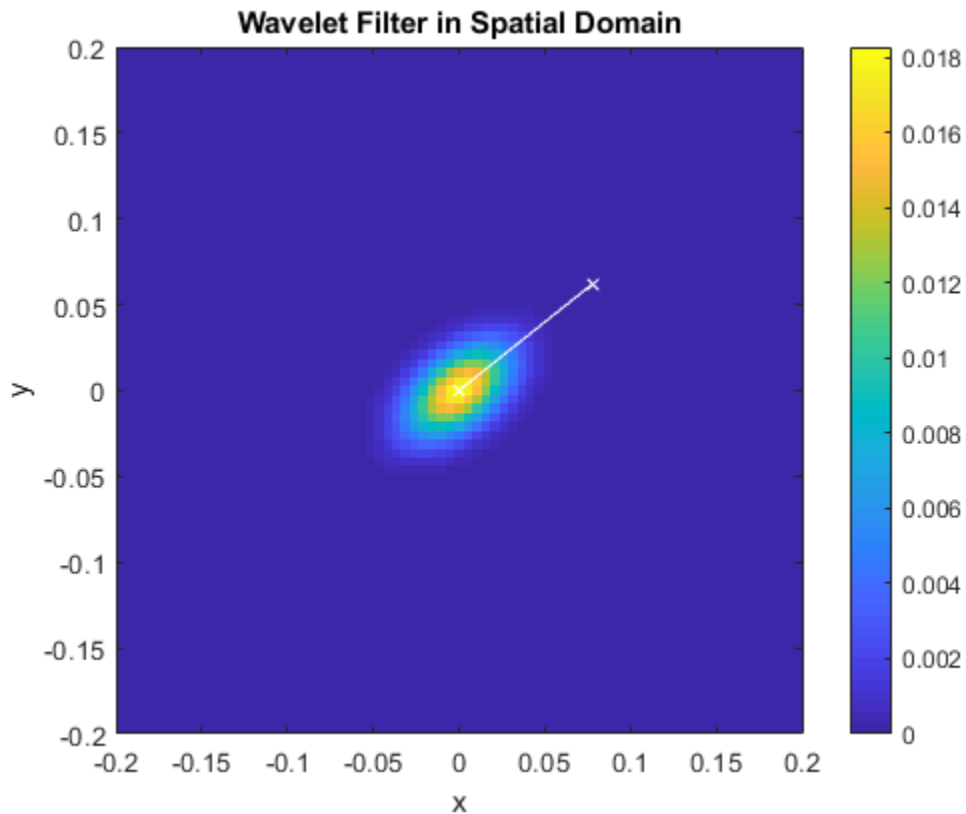
From the same filter bank and scale, choose a rotated wavelet filter. Plot the absolute value of the wavelet in the spatial domain. Use the associated rotation angle in `fparams{1}.rotations`, and rotate clockwise the vector $(0, 0.1)$ by that amount. Plot the rotated vector in the figure. Confirm that the vector is aligned with the semi-major axis of the rotated wavelet.

```
whichRotAngle = 3;
rotAngle = fparams{whichFilterBank}.rotations(whichRotAngle);
rmat = [cos(rotAngle) sin(rotAngle) ; -sin(rotAngle) cos(rotAngle)];

wvf = psif{whichFilterBank}(:, :, 1+(whichScale-1)*numRot+(whichRotAngle-1));
wvf_ifft = ifft2(wvf);

rvec = rmat*vec';
```

```
figure
helperPlotWavelet(wvf_iff)
hold on
plot([0 rvec(1)], [0 rvec(2)], 'wx-')
xlim([-0.2 0.2])
ylim([-0.2 0.2])
```



Appendix

The following helper functions are used in this example.

```
function helperPlotWaveletFT(wavelet, cenFreq)
Nx = size(wavelet,2);
```

```

Ny = size(wavelet,1);
fx = -1/2:1/Nx:1/2-1/Nx;
fy = -1/2:1/Ny:1/2-1/Ny;
imagesc(fx,fy,fftshift(wavelet))
colorbar
axis xy
hold on
xlabel('\omega_x','Interpreter','latex')
ylabel('\omega_y','Interpreter','latex')
axis equal
axis tight
plot(cenFreq(1),cenFreq(2),'k^','markerfacecolor',[0 0 0])
title('Wavelet Filter in Frequency Domain')
end

```

```

function helperPlotWavelet(wavelet)
Nx = size(wavelet,2);
Ny = size(wavelet,1);
fx = -1/2:1/Nx:1/2-1/Nx;
fy = -1/2:1/Ny:1/2-1/Ny;
imagesc(fx,fy,abs(fftshift(wavelet)))
colorbar
axis xy
hold on
xlabel('x')
ylabel('y')
axis equal
axis tight
title('Wavelet Filter in Spatial Domain')
end

```

Input Arguments

sf — Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

fb — Filter banks

positive integer | vector of positive integers

Filter banks, specified as an integer or vector of integers between 1 and `numfilterbanks(sf)` inclusive. If `fb` is a scalar, `psifilters` is an M -by- N -by- L matrix and `filterparams` is a MATLAB table.

Output Arguments

phif — Fourier transform of scaling filter

real-valued matrix

Fourier transform of scaling filter, returned as a real-valued 2-D matrix. The precision of `phif` depends on the value of the `Precision` property of the scattering framework. `phif` has dimensions M -by- N , where M and N are the padded row and column sizes of the scattering framework.

psifilters — Fourier transforms of the wavelet filters

cell array

Fourier transforms of the wavelet filters, returned as an Nfb -by-1 cell array, where Nfb is the number of filter banks in the scattering framework. Each element of `psifilters` is a 3-D array. The 3-D arrays are M -by- N -by- L , where M and N are the padded row and column sizes of the wavelet filters and L is the number of wavelet filters for each filter bank. The wavelet filters are ordered by increasing scale with `NumRotations` wavelet filters for each scale.

Example: Note that `size(psifilters,3)` is equal to `size(f,1)`.

f — Center spatial frequencies

cell array

Center spatial frequencies of the wavelet filters, returned as a Nfb -by-1 cell array where Nfb is the number of filter banks in the scattering framework. The j th element of `f` contains the center frequencies for the j th wavelet filter bank in `psifilters`. Each element of `f` is an L -by-2 matrix with each row containing the center frequencies of the corresponding L th wavelet. The spatial frequencies are in cycles per pixel.

filterparams — Filter parameters

cell array

Filter parameters for the 2-D scattering framework, `sf`. `filterparams` is an Nfb -by-1 cell array of MATLAB tables, where the j th element of `filterparams` is a MATLAB table

containing the filter parameters for the j th filter bank. Each table contains these variables:

- `Q` — The quality factor of the filter bank, returned as an integer.
- `J` — The highest factor used in the dilation of the Morlet wavelets, $2^{J/Q}$, returned as an integer.
- `precision` — The precision of the scattering framework, returned as `'single'` or `'double'`.
- `omegapsi` — The wavelet center frequencies in descending order (highest to lowest), returned as a vector.
- `freqsigmapsi` — The wavelet frequency standard deviations, returned as a vector.
- `slant` — The slant parameter for the spatial vertical semi-major axis of the wavelet, returned as a real number. The slant parameter, also known as the spatial aspect ratio, characterizes the shape of the support of the wavelet.
- `spatialsigmapsi` — The wavelet spatial standard deviations, returned as a vector.
- `spatialsigmaphi` — The scaling filter spatial standard deviation, returned as a real number.
- `psi3dBbw` — The wavelet 3 dB bandwidths, returned as a vector.
- `psiftsupport` — The wavelet frequency support, returned as a vector.
- `phiftsupport` — The scaling filter frequency support, returned as a real number.
- `phi3dBbw` — The scaling filter 3 dB bandwidth, returned as a real number.
- `rotations` — The wavelet orientation angles in radians, returned as a vector. The length of `rotations` equals the `NumRotations` value associated with the filter bank.

The following vectors in the table have equal length: `omegapsi`, `freqsigmapsi`, `spatialsigmapsi`, `psi3dBbw`, and `psiftsupport`.

The total number of wavelet filters in a filter bank is `length(omegapsi) × length(rotations)`. See “Determine Wavelet Semi-Major Axis” on page 1-443.

Definitions

Slant Parameter

The *slant parameter* or *spatial aspect ratio* controls the shape of the elliptical support of the Morlet wavelet.

The Morlet wavelet is of the form

$$\psi(x, y) = e^{-(x^2 + \nu^2 y^2)/2\sigma^2} e^{i\omega\lambda x}$$

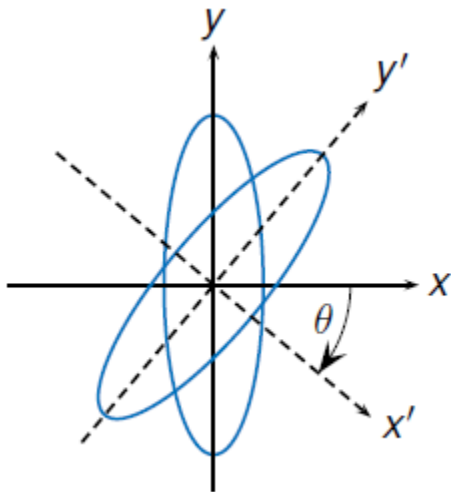
where ν is the *slant parameter*. Typically, $\nu < 1$, so that the ellipse $\frac{x^2}{\sigma^2} + \frac{y^2}{\sigma^2/\nu^2}$ is

elongated spatially in the y -direction. The wavelet is rotated in a clockwise direction:

$$\begin{pmatrix} x' \\ y' \end{pmatrix} = \begin{pmatrix} \cos \theta & \sin \theta \\ -\sin \theta & \cos \theta \end{pmatrix} \begin{pmatrix} x \\ y \end{pmatrix}.$$

The rotated Morlet wavelet is of the form

$$\psi(x', y') = e^{-(x'^2 + \nu^2 y'^2)/2\sigma^2} e^{i\omega\lambda x'}$$



If $g(x,y)$ and $G(\omega_x,\omega_y)$ form a Fourier pair: $g(x,y) \xleftrightarrow{FT} G(\omega_x,\omega_y)$, then so do their rotations:

$$g(x\cos\theta + y\sin\theta, -x\sin\theta + y\cos\theta) \xleftrightarrow{FT} G(\omega_x\cos\theta + \omega_y\sin\theta, -\omega_x\sin\theta + \omega_y\cos\theta)$$

The Fourier transform of the Morlet wavelet is

$$\widehat{\psi}(\omega_x, \omega_y) = \frac{2\pi\sigma^2}{\nu} e^{-\frac{\sigma^2}{\nu} \left((\omega_x - \omega_\lambda)^2 + \frac{\omega_y^2}{\nu^2} \right)}$$

A given wavelet $\psi(x,y)$ has a reported bandwidth bw . The reported bandwidth is dependent on the scale, but not the rotation angle. For a reported 3 dB bandwidth bw , the bandwidth along the semi-major spatial axis of the ellipse that describes the wavelet support is $bw \times \text{slant}$.

The semi-major spatial axis depends on the rotation angle. The semi-major spatial axis can be computed from the vector rotations in the output argument `filterparams`.

As ordered in the output arguments `f` and `psifilters`, the wavelet filter `psifilters(:, :, 1 + k × NumRotations)` for an integer k is the first, unrotated wavelet at a given scale. The wavelet has a center spatial frequency of `f(1 + k × NumRotations, :)`, which is of the form (ω_x, θ) . See “Determine Wavelet Semi-Major Axis” on page 1-443.

See Also

`waveletScattering2`

Introduced in R2019a

filters

DWT filter bank filters

Syntax

```
[Lo,Hi] = filters(fb)
```

Description

`[Lo,Hi] = filters(fb)` returns the lowpass (scaling) and highpass (wavelet) filters, `Lo` and `Hi` respectively, for the DWT filter bank `fb`. `Lo` and `Hi` are both L -by-2 matrices. L is an even positive integer. The first column of `Lo` and the first column of `Hi` are the analysis filters. The second column of `Lo` and the second column of `Hi` are the synthesis filters.

Examples

DWT Filter Bank Filters

Obtain the lowpass and highpass filters for the order 4 symlet. Confirm the filter bank is orthogonal.

```
fb = dwtfilterbank('Wavelet','sym4');  
[Lo,Hi] = filters(fb)
```

```
Lo = 8×2
```

```
-0.0758    0.0322  
-0.0296   -0.0126  
 0.4976   -0.0992  
 0.8037    0.2979  
 0.2979    0.8037  
-0.0992    0.4976  
-0.0126   -0.0296
```

```
0.0322 -0.0758
```

```
Hi = 8x2
```

```
-0.0322 -0.0758
-0.0126  0.0296
 0.0992  0.4976
 0.2979 -0.8037
-0.8037  0.2979
 0.4976  0.0992
 0.0296 -0.0126
-0.0758 -0.0322
```

```
isOrthogonal(fb)
```

```
ans = logical
      1
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

Lo — Lowpass (scaling) filters

real-valued matrix

Lowpass (scaling) filters for the DWT filter bank, returned as an L-by-2 matrix. L is an even positive integer. The first column of Lo is the analysis filter and the second column is the synthesis filter.

For orthogonal wavelets, the lowpass synthesis and lowpass analysis filters are time-reversed versions of each other.

Hi — Highpass (wavelet) filters

real-valued matrix

Highpass (wavelet) filters for the DWT filter bank, returned as an L-by-2 matrix. L is an even positive integer. The first column of **Hi** is the analysis filter and the second column is the synthesis filter.

For orthogonal wavelets, the highpass synthesis and highpass analysis filters are time-reversed versions of each other.

See Also

`dwtfilterbank` | `wfilters`

Introduced in R2018a

framebounds

DWT filter bank frame bounds

Syntax

```
[a,b] = framebounds(fb)
```

Description

`[a,b] = framebounds(fb)` returns the frame bounds for the DWT filter bank, `fb`. For an orthogonal wavelet filter bank, the theoretical frame bounds `a` and `b` are equal to 1.

Examples

DWT Filter Bank Frame Bounds

Obtain the frame bounds for the orthogonal Daubechies db6 wavelet.

```
wv = 'db6';  
fb = dwtfilterbank('Wavelet',wv)
```

```
fb =  
    dwtfilterbank with properties:
```

```
        Wavelet: 'db6'  
    SignalLength: 1024  
         Level: 6  
    SamplingFrequency: 1  
        FilterType: 'Analysis'  
    CustomWaveletFilter: []  
    CustomScalingFilter: []
```

```
[a,b] = framebounds(fb)
```

```
a = 1.0000
```

```
b = 1.0000
```

The filter bank has the default filter type `Analysis`. Create a second filter bank using the same orthogonal wavelet but with the filter type `Synthesis`. Obtain the frame bounds of this filter bank. Observe there is no change.

```
fbSynthesis = dwtfilterbank('Wavelet',wv,'FilterType','Synthesis');  
[a2,b2] = framebounds(fbSynthesis)
```

```
a2 = 1.0000
```

```
b2 = 1.0000
```

Create a filter bank for the biorthogonal `bior3.9` wavelet. Obtain the frame bounds. The frame bounds will not be equal to 1.

```
wv = 'bior3.9';  
fbA = dwtfilterbank('Wavelet',wv);  
[c,d] = framebounds(fbA)
```

```
c = 0.6250
```

```
d = 3.2982
```

Create a second filter bank using the same biorthogonal wavelet but with the filter type `Synthesis`. Obtain the frame bounds of this filter bank. Since the wavelet is biorthogonal, the frame bounds change.

```
fbASynthesis = dwtfilterbank('Wavelet',wv,'FilterType','Synthesis');  
[c2,d2] = framebounds(fbASynthesis)
```

```
c2 = 0.5502
```

```
d2 = 2.0015
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

a — Lower frame bound

positive real number

Lower frame bound of the DWT filter bank `fb`, returned as a positive real number.

b — Upper frame bound

positive real number

Upper frame bound of the DWT filter bank `fb`, returned as a positive real number.

See Also

`dwtfilterbank` | `isBiorthogonal` | `isOrthogonal`

Introduced in R2018a

freqz

CWT filter bank frequency responses

Syntax

```
[psidft,f] = freqz(fb)  
freqz(fb)
```

Description

`[psidft,f] = freqz(fb)` returns the frequency responses for the wavelet filters, `psidft`, and the frequency vector, `f`, for the continuous wavelet transform (CWT) filter bank, `fb`. Frequencies are in cycles/sample or Hz. If you specify a sampling period, the frequencies are in cycles/unit time where the time unit is the unit of the duration `SamplingPeriod`.

The frequency responses, `psidft`, are one-sided frequency responses for the positive frequencies. For the analytic wavelets supported by `cwtfilterbank`, the frequency responses are real-valued and are equivalent to the magnitude frequency response.

`freqz(fb)` with no output arguments plots the magnitude frequency responses for the CWT filter bank, `fb`.

Examples

Frequency Responses of a Continuous Wavelet Transform Filter Bank

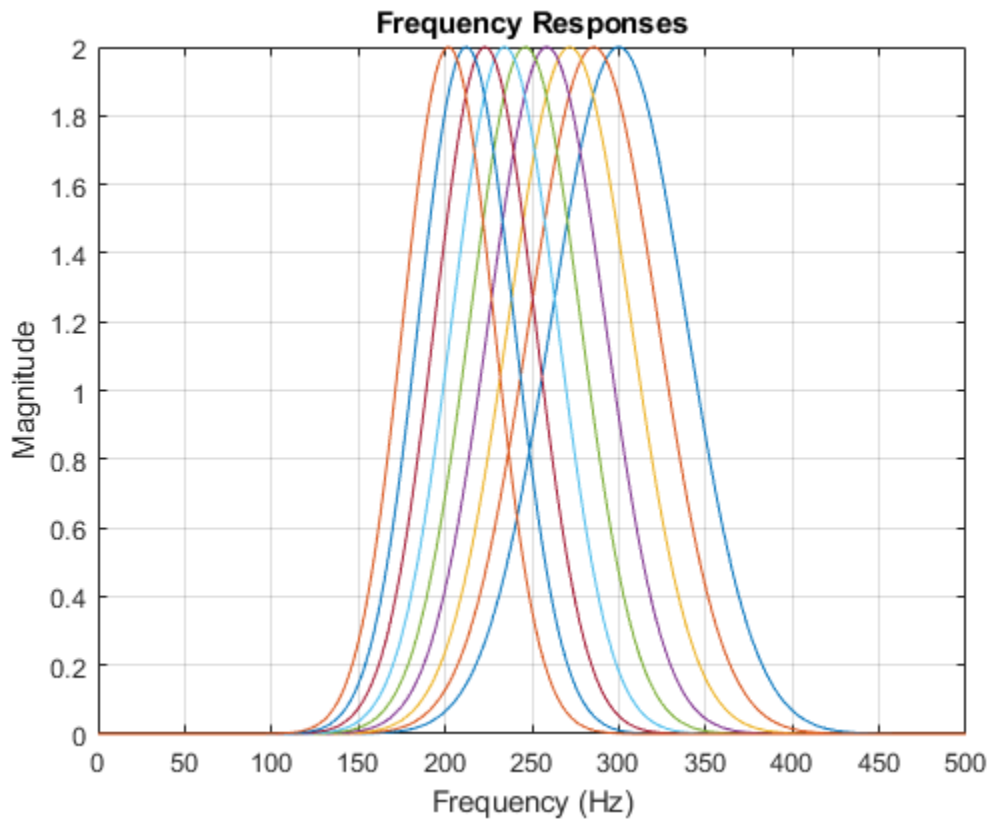
Create a CWT filter bank. Set the voices per octave to 14, the sampling frequency to 1000 Hz, and frequency limits to range from 200 Hz to 300 Hz.

```
fb = cwtfilterbank('VoicesPerOctave',14,'SamplingFrequency',1000,'FrequencyLimits',[200  
fb =  
    cwtfilterbank with properties:
```

```
VoicesPerOctave: 14
    Wavelet: 'morse'
SamplingFrequency: 1000
    SamplingPeriod: []
    PeriodLimits: []
    SignalLength: 1024
    FrequencyLimits: [200 300]
    TimeBandwidth: 60
WaveletParameters: []
    Boundary: 'reflection'
```

Obtain the frequency responses for the filter bank. Plot the frequency responses.

```
[psidft,f] = freqz(fb);
plot(f,psidft)
grid on
title('Frequency Responses')
xlabel('Frequency (Hz)')
ylabel('Magnitude')
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

psidft — Frequency responses

real-valued 2-D matrix

Frequency responses of a CWT filter bank, returned as a real-valued N_s -by- M matrix where N_s is the number of scales. If the filter bank `Boundary` is 'reflection', M is equal to $2 \times \text{floor}(\text{SignalLength}/2) + 1$. If the filter bank `Boundary` is 'periodic', M is equal to $\text{floor}(\text{SignalLength}/2) + 1$.

The frequency responses, `psidft`, are one-sided frequency responses for the positive frequencies. For the analytic wavelets supported by `cwtfilterbank`, the frequency responses are real-valued and are equivalent to the magnitude frequency response.

f — Frequencies

real-valued vector

Frequencies, in cycles/sample or Hz, returned as a real-valued vector. If the filter bank `Boundary` is 'reflection', `f` has length $2 \times \text{floor}(\text{SignalLength}/2) + 1$. If the filter bank `Boundary` is 'periodic', `f` has length $\text{floor}(\text{SignalLength}/2) + 1$.

If you specify a sampling period, the frequencies are in cycles/unit time where the time unit is the unit of the duration `SamplingPeriod`.

Data Types: `double`

See Also

`centerFrequencies` | `centerPeriods` | `cwtfilterbank` | `powerbw`

Introduced in R2018a

freqz

DWT filter bank frequency responses

Syntax

```
[psidft,f] = freqz(fb)  
[psidft,f,phidft] = freqz(fb)  
freqz(fb)
```

Description

`[psidft,f] = freqz(fb)` returns the complex-valued frequency responses for the wavelet filters, `psidft`, and the frequency vector, `f`, for the discrete wavelet transform (DWT) filter bank `fb`. `psidft` is an L-by-N matrix where L is the filter bank `Level` and N is the filter bank `SignalLength`. `f` is a real-valued vector of length L. Frequencies are in cycles/sample or in Hz if a sampling frequency is defined in `fb`. The frequency responses are centered so that the zero frequency is in the middle.

`[psidft,f,phidft] = freqz(fb)` returns the complex-valued frequency responses for the scaling filters, `phidft`, for the DWT filter bank `fb` at all levels of the decomposition.

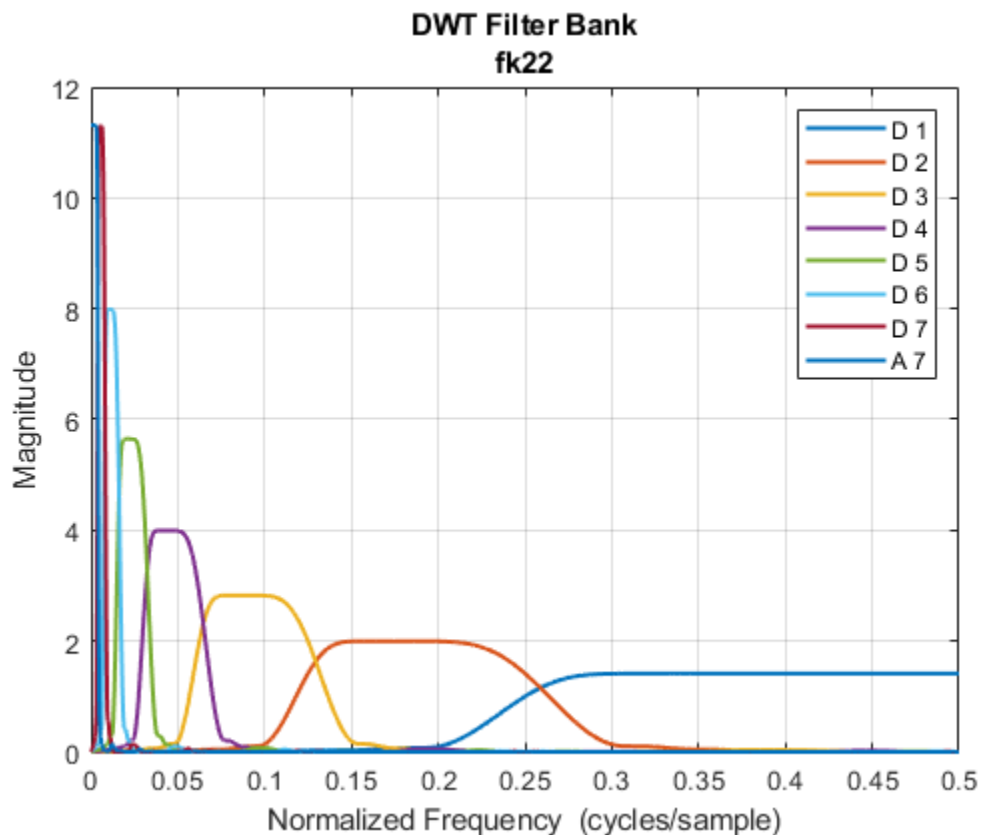
`freqz(fb)` plots the one-sided magnitude frequency responses for the wavelet filter bank, `fb`. Magnitude frequency responses are plotted for all wavelet bandpass filters and the coarsest resolution scaling filter. The legend is interactive. To toggle the visibility of the filter magnitude response, click on the corresponding line in the legend.

Examples

DWT Filter Bank Frequency Responses

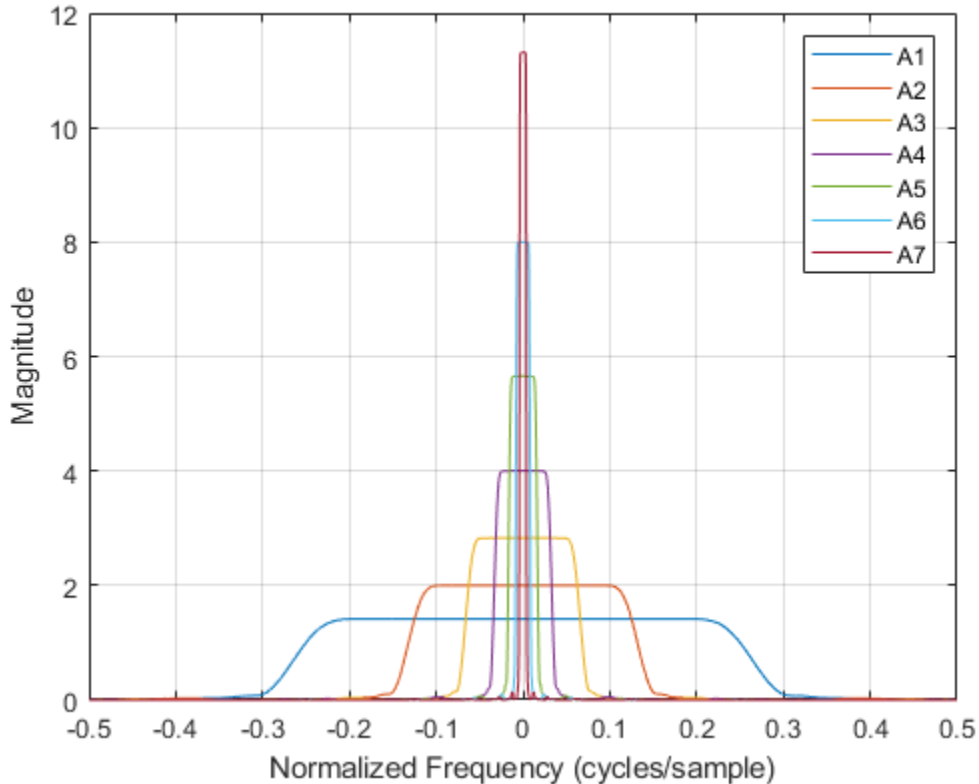
Create a DWT filter bank for a length 4096 signal and the Fejér-Korovkin `fk22` wavelet. Plot the magnitude frequency responses of the wavelet filters and final resolution scaling filter.

```
len = 4096;
fb = dwtfilterbank('Wavelet','fk22','SignalLength',len);
freqz(fb)
```



Obtain the frequency responses for the wavelet and scaling filters. Plot the magnitude frequency responses of the scaling filters at all levels of decomposition.

```
[psidft,f,phidft] = freqz(fb);
plot(f,abs(phidft))
grid on
xlabel('Normalized Frequency (cycles/sample)')
ylabel('Magnitude')
legend('A1','A2','A3','A4','A5','A6','A7')
```

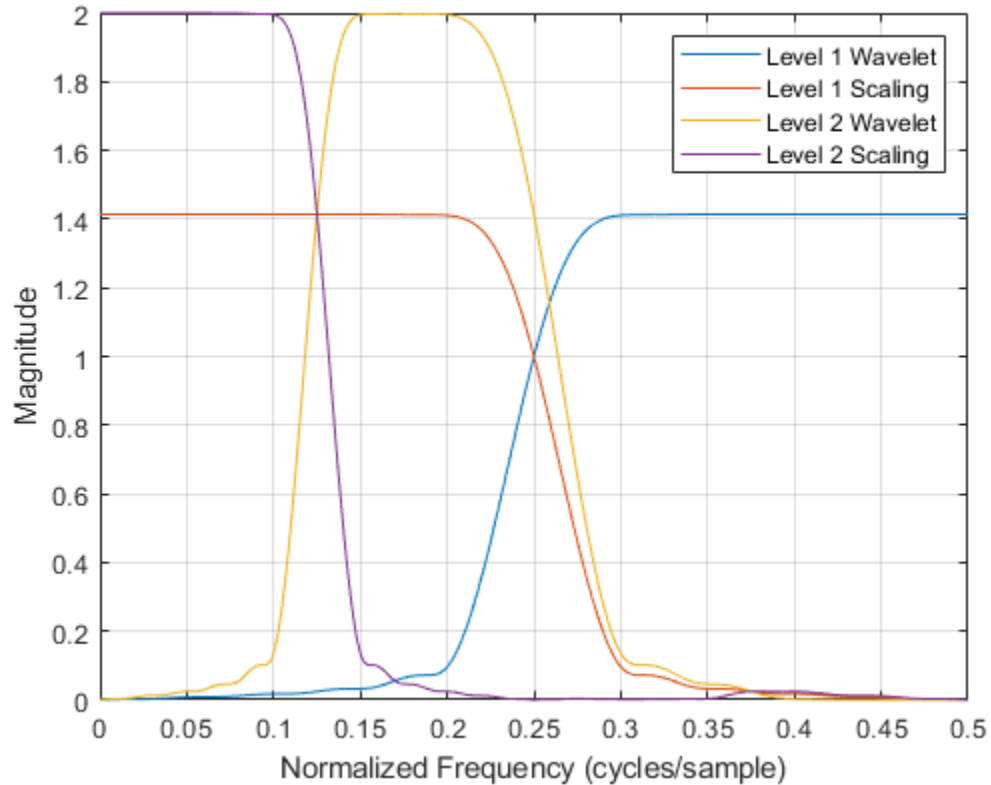


Plot the one-sided magnitude frequency responses of the wavelet and scaling filters at the first two levels of decomposition. Note how the second level frequency responses overlap the magnitude response of the first level scaling filter.

```

plot(f(len/2:end),abs(psidft(1,len/2:end))')
hold on
plot(f(len/2:end),abs(phidft(1,len/2:end))')
plot(f(len/2:end),abs(psidft(2,len/2:end))')
plot(f(len/2:end),abs(phidft(2,len/2:end))')
grid on
xlabel('Normalized Frequency (cycles/sample)')
ylabel('Magnitude')
legend('Level 1 Wavelet','Level 1 Scaling','Level 2 Wavelet','Level 2 Scaling')

```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

psidft — Wavelet filter frequency responses

complex-valued matrix

Wavelet filter frequency responses for the DWT filter bank `fb`, returned as an L-by-N matrix where L is the filter bank `Level` and N is the filter bank `SignalLength`. The frequency responses are centered so that the zero frequency is centered in the middle.

f — Frequencies

real-valued vector

Frequencies, in cycles/sample or Hz, returned as a real-valued vector of length N where N is the filter bank `SignalLength`. If a sampling frequency is specified in `fb`, frequencies are in Hz.

Data Types: `double`

phidft — Scaling function frequency responses

complex-valued matrix

Scaling function frequency responses for the DWT filter bank `fb`, returned as an L-by-N matrix where L is the filter bank `Level` and N is the filter bank `SignalLength`. The frequency responses are centered so that the zero frequency is centered in the middle.

See Also

`dwtfilterbank`

Introduced in R2018a

gauswavf

Gaussian wavelet

Syntax

```
[psi,x] = gauwavf(lb,ub,n)
[psi,x] = gauwavf(lb,ub,n,p)
[psi,x] = gauwavf(lb,ub,n,wname)
```

Description

`[psi,x] = gauwavf(lb,ub,n)` returns the 1st order derivative of the Gaussian wavelet, `psi`, on an `n`-point regular grid, `x`, for the interval `[lb,ub]`. The effective support of the Gaussian wavelets is `[-5, 5]`.

`[psi,x] = gauwavf(lb,ub,n,p)` returns the `p`th derivative. `p` is an integer from 1 through 8.

The Gaussian function is defined as $C_p e^{-x^2}$. C_p is such that the 2-norm of the `p`th derivative of `psi` is equal to 1.

Note For visualizing the second or third order derivative of Gaussian wavelets, the convention is to use the negative of the normalized derivative. In the case of the second derivative, scaling by -1 produces a wavelet with its main lobe in the positive `y` direction. This scaling also makes the Gaussian wavelet resemble the Mexican hat, or Ricker, wavelet. The validity of the wavelet is not affected by the -1 scaling factor.

`[psi,x] = gauwavf(lb,ub,n,wname)` used the valid wavelet family short name `wname` plus the order of the derivative in a character vector or string scalar, such as `'gaus4'`. To see valid character vectors for Gaussian wavelets, use `waveinfo('gaus')` or use `wavemngr('read',1)` and refer to the Gaussian section.

Examples

Create Gaussian Wavelet

This example shows how to create and plot a Gaussian wavelet of order 8.

Set the initial effective support and grid parameters.

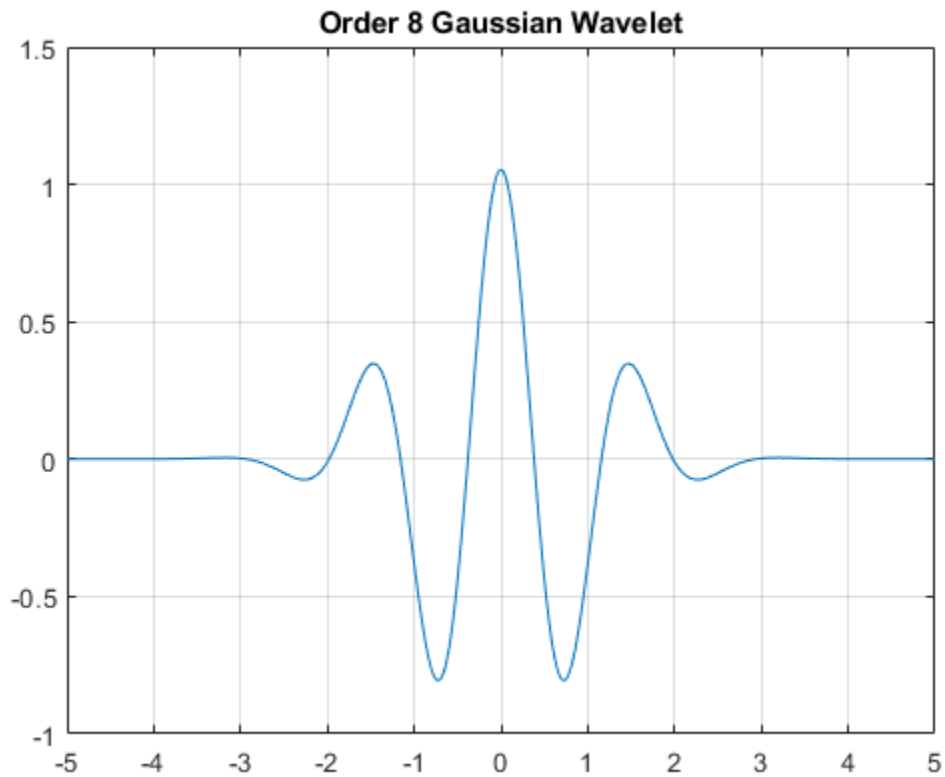
```
lb = -5;  
ub = 5;  
n = 1000;
```

Compute the Gaussian wavelet of order 8.

```
[psi,x] = gauswavf(lb,ub,n,8);
```

Now plot the wavelet.

```
plot(x,psi)  
title('Order 8 Gaussian Wavelet')  
grid on
```



Input Arguments

lb — Left endpoint

real number

Left endpoint of the closed interval, specified as a real number. `lb` is strictly less than `ub`.

Data Types: double

ub — Right endpoint

real number

Right endpoint of the closed interval, specified as a real number. `ub` is strictly greater than `lb`.

Data Types: `double`

n — Number of regularly spaced points

positive integer

Number of regularly spaced points in the interval `[lb,ub]`, specified as a positive integer. The derivative of the Gaussian is evaluated at these points.

Data Types: `double`

p — Derivative

positive integer

Positive integer defining the order of the derivative of the Gaussian wavelet, specified as a positive integer. `p` is an integer from 1 through 8.

wname — Gaussian wavelet

character vector | string scalar

Gaussian wavelet to evaluate, specified as a character vector or string scalar. `wname` is of the form `'cgauN'` where `N` is an integer that denotes the order of the derivative of the Gaussian wavelet. `N` is an integer from 1 through 8.

Example: `'gaus4'` denotes the fourth derivative of the Gaussian wavelet.

Output Arguments

psi — Derivative of Gaussian wavelet

real-valued vector

Derivative of the Gaussian wavelet, returned as a real-valued 1-by-`N` vector.

x — Sample points

real-valued vector

Sample points where the derivative of the Gaussian wavelet is evaluated, returned as a real-valued 1-by-`N` vector. The sample points are evenly distributed between `lb` and `ub`.

See Also

waveinfo | wavemngr

Introduced before R2006a

get

WPTREE contents

Syntax

```
[FieldValue1,FieldValue2, ...] =  
get(T,'FieldName1','FieldName2', ...)  
[FieldValue1,FieldValue2, ...] = get(T)
```

Description

[FieldValue1,FieldValue2, ...] = get(T,'FieldName1','FieldName2', ...) returns the content of the specified fields for the WPTREE object T.

For the fields that are objects or structures, you can get the subfield contents, giving the name of these subfields as '*FieldName*' values. (See “Examples” below.)

[FieldValue1,FieldValue2, ...] = get(T) returns all the field contents of the tree T.

The valid choices for '*FieldName*' are

'dtree'	DTREE parent object
'wavInfo'	Structure (wavelet information)

The fields of the wavelet information structure, 'wavInfo', are also valid for '*FieldName*':

'wavName'	Wavelet name
'Lo_D'	Low Decomposition filter
'Hi_D'	High Decomposition filter
'Lo_R'	Low Reconstruction filter

'Hi_R'	High Reconstruction filter
'entInfo'	Structure (entropy information)

The fields of the entropy information structure, 'entInfo', are also valid for 'FieldName':

'entName'	Entropy name
'entPar'	Entropy parameter

Or fields of DTREE parent object:

'ntree'	NTREE parent object
'allNI'	All nodes information
'terNI'	Terminal nodes information

Or fields of NTREE parent object:

'wtbo'	WTBO parent object
'order'	Order of the tree
'depth'	Depth of the tree
'spsch'	Split scheme for nodes
'tn'	Array of terminal nodes of the tree

Or fields of WTBO parent object:

'wtboInfo'	Object information
'ud'	Userdata field

Examples

```
% Compute a wavelet packets tree
x = rand(1,1000);
t = wpdec(x,2,'db2');
o = get(t,'order');
[o,tn] = get(t,'order','tn');
```

```
[o,allNI,tn] = get(t,'order','allNI','tn');  
[o,wavInfo,allNI,tn] = get(t,'order','wavInfo','allNI','tn');  
[o,tn,Lo_D,EntName] = get(t,'order','tn','Lo_D','EntName');  
[wo,nt,dt] = get(t,'wtbo','ntree','dtree');
```

See Also

disp | read | set | write

Introduced before R2006a

getLabelDefinitions

Get label definitions in labeled signal set

Syntax

```
lbldefs = getLabelDefinitions(lss)
```

Description

`lbldefs = getLabelDefinitions(lss)` returns a vector of `signalLabelDefinition` objects with the labels of the labeled signal set `lss`.

Changing `lbldefs` does not affect the labeled set. To modify label definitions, use `editLabelDefinition`, `addLabelDefinitions`, and `removeLabelDefinition`.

Examples

Get Label Definitions

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
      NumMembers: 2  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [2x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.

Use `setLabelValue` to add data to the set.

Retrieve the definitions of the labels in the set.

```
dfs = getLabelDefinitions(lss);  
  
for k = 1:length(dfs)  
    dfs(k)  
end  
  
ans =  
    signalLabelDefinition with properties:  
        Name: "WhaleType"  
        LabelType: "attribute"  
        LabelDataType: "categorical"  
        Categories: [3x1 string]  
        DefaultValue: []  
        Sublabels: [0x0 signalLabelDefinition]  
        Tag: ""  
        Description: "Whale type"
```

Use `labeledSignalSet` to create a labeled signal set.

```
ans =  
    signalLabelDefinition with properties:  
        Name: "MoanRegions"  
        LabelType: "roi"  
        LabelDataType: "logical"  
        ValidationFunction: []  
        ROIlimitsDataType: "double"  
        DefaultValue: []  
        Sublabels: [0x0 signalLabelDefinition]  
        Tag: ""  
        Description: "Regions where moans occur"
```

Use `labeledSignalSet` to create a labeled signal set.

```
ans =  
    signalLabelDefinition with properties:  
        Name: "TrillRegions"  
        LabelType: "roi"
```

```

    LabelDataType: "logical"
    ValidationFunction: []
    ROIlimitsDataType: "double"
    DefaultValue: []
    Sublabels: [1x1 signalLabelDefinition]
    Tag: ""
    Description: "Regions where trills occur"

```

Use `labeledSignalSet` to create a labeled signal set.

Input Arguments

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

Output Arguments

lbldefs — Signal label definitions

`signalLabelDefinition` object

Signal label definitions, returned as a `signalLabelDefinition` object or a vector of such objects.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

getLabeledSignal

Get labeled signals from labeled signal set

Syntax

```
[t,info] = getLabeledSignal(lss)
[t,info] = getLabeledSignal(lss,midX)
```

Description

[t,info] = getLabeledSignal(lss) returns a table with all the signals and labeled data in the labeled signal set lss.

[t,info] = getLabeledSignal(lss,midX) returns a table with the signals specified in midX.

Examples

Get Labeled Signal

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss
```

```
lss =
  labeledSignalSet with properties:
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Get a table with all the signals in `lss`.

```
t = getLabeledSignal(lss)
```

```
t=2x4 table
```

	Signal	WhaleType	MoanRegions	TrillRegions
Member{1}	[79572x1 double]	blue	[3x2 table]	[1x3 table]
Member{2}	[76579x1 double]	blue	[3x2 table]	[1x3 table]

Identify the sublabels of the trill regions.

```
d = getLabelNames(lss, 'TrillRegions')
```

```
d =  
"TrillPeaks"
```

Get the labeled signal corresponding to the second member of the set.

```
[lbs,info] = getLabeledSignal(lss,2)
```

```
lbs=1x4 table
```

	Signal	WhaleType	MoanRegions	TrillRegions
Member{2}	[76579x1 double]	blue	[3x2 table]	[1x3 table]

```
info = struct with fields:  
TimeInformation: "sampleRate"  
SampleRate: 4000
```

Extract the signal. Determine its sample rate and use the sample rate to compute the time vector.

```
fs = info.SampleRate;  
sg = getSignal(lss,2);  
t = (0:length(sg)-1)/fs;
```

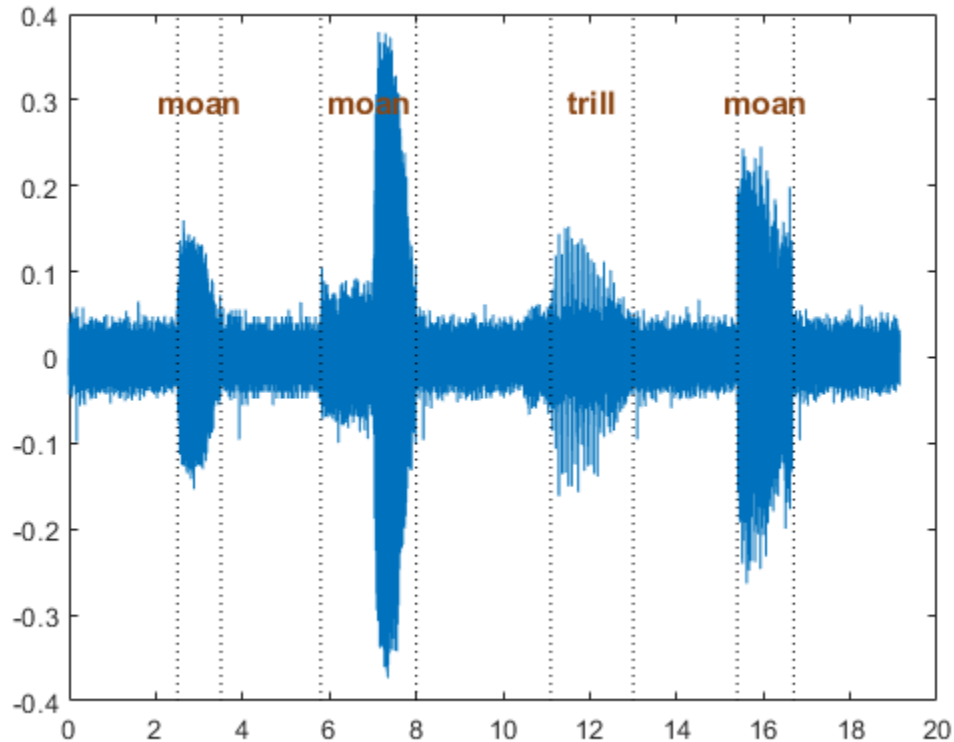
Identify the moan and trill regions of interest and the peaks of the trill region.

```
mvals = getLabelValues(lss,2,'MoanRegions');  
tvals = getLabelValues(lss,2,'TrillRegions');
```

```
peaks = getLabelValues(lss,2,{'TrillRegions','TrillPeaks'});
```

Plot the signal. Highlight the regions of interest and the peaks.

```
plot(t,sg)  
  
hold on  
[X,Y] = meshgrid([mvals.ROIlimits;tvals.ROIlimits],ylim);  
plot(X,Y,':k')  
topts = {'HorizontalAlignment','center','FontWeight','bold', ...  
        'FontSize',12,'Color',[139 69 19]/255};  
text((X(1,1:4)+X(1,5:end))/2,Y(2,5:end)-0.1, ...  
     ["moan" "moan" "moan" "trill"],topts{:})  
hold off
```

Input Arguments

lss — Labeled signal set
labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

Output Arguments

t — Labeled signal

table

Labeled signal, specified as a table.

info — Time information

structure

Time information, returned as a structure.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

getLabelNames

Get label names in labeled signal set

Syntax

```
lblnames = getLabelNames(lss)
sublblnames = getLabelNames(lss, lblname)
```

Description

`lblnames = getLabelNames(lss)` returns a string array containing the label names in the labeled signal set `lss`.

`sublblnames = getLabelNames(lss, lblname)` returns a string array containing the sublabel names for the label named `lblname` in the labeled signal set `lss`.

Examples

Get Label Names

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss

lss =
    labeledSignalSet with properties:
        Source: {2x1 cell}
        NumMembers: 2
        TimeInformation: "sampleRate"
        SampleRate: 4000
        Labels: [2x3 table]
        Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Get the names of the labels in the set.

```
str = getLabelNames(lss)
```

```
str = 3x1 string array  
    "WhaleType"  
    "MoanRegions"  
    "TrillRegions"
```

Verify that only the 'TrillRegions' label has sublabels.

```
for kj = 1:length(str)  
    sbstr = str{kj};  
    sbl = [sbstr getLabelNames(lss,sbstr)]  
end
```

```
sbl =  
'WhaleType'
```

```
sbl =  
'MoanRegions'
```

```
sbl = 1x2 string array  
    "TrillRegions"    "TrillPeaks"
```

Input Arguments

lss — **Labeled signal set**
labeledSignalSet object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

lblname — Label name

character vector | string scalar

Label name, specified as a character vector or a string scalar.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Output Arguments

lblnames — Label names

string array

Label names, returned as a string array.

sublblnames — Sublabel names

string array

Sublabel names, returned as a string array.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

getLabelValues

Get label values from labeled signal set

Syntax

```
val = getLabelValues(lss)
val = getLabelValues(lss, midx)

[val, sublbltbl] = getLabelValues(lss, midx, lblname)

[ ___ ] = getLabelValues( ___, 'LabelRowIndex', ridx)
[ ___ ] = getLabelValues( ___, 'SublabelRowIndex', sridx)
```

Description

`val = getLabelValues(lss)` returns a table containing the label values for all members of the labeled signal set `lss`.

`val = getLabelValues(lss, midx)` returns a table containing the label values for the member specified by `midx`.

`[val, sublbltbl] = getLabelValues(lss, midx, lblname)` returns the value of the label named `lblname`. If `lblname` has sublabels, then the table `sublbltbl` shows the structure of the label value and its sublabel variables.

`[___] = getLabelValues(___, 'LabelRowIndex', ridx)` specifies the row index, `ridx`, of an ROI or point label whose value you want to get.

`[___] = getLabelValues(___, 'SublabelRowIndex', sridx)` specifies the row index, `sridx`, of an ROI or point sublabel whose value you want to get.

Examples

Get Label Values

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss
```

```
lss =
  labeledSignalSet with properties:
```

```
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use labelDefinitionsHierarchy to see a list of labels and sublabels.
Use setLabelValue to add data to the set.

Get the values of the labels.

```
lbls = getLabelValues(lss)
```

```
lbls=2x3 table
```

	WhaleType	MoanRegions	TrillRegions
Member{1}	blue	[3x2 table]	[1x3 table]
Member{2}	blue	[3x2 table]	[1x3 table]

Display the moan ROI limits for the second signal of the set.

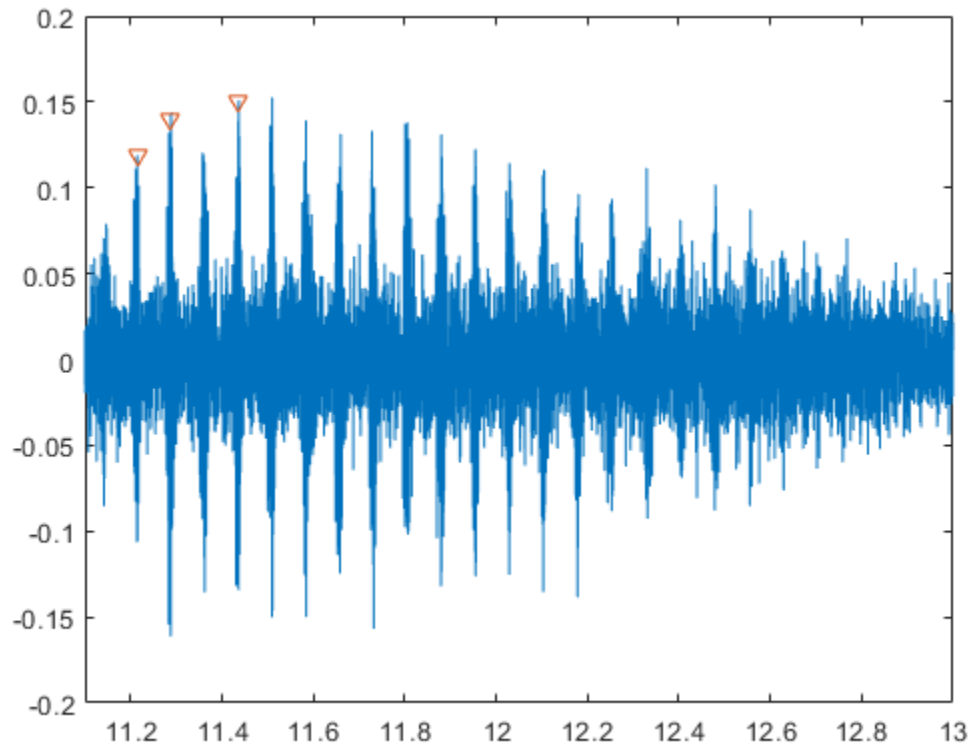
```
lbb = getLabelValues(lss,2,'MoanRegions')
```

```
lbb=3x2 table
```

ROIlimits	Value
2.5 3.5	[1]
5.8 8	[1]
15.4 16.7	[1]

Plot the trill region of the signal between the ROI limits. Display the labeled trill peaks.

```
tvals = getLabelValues(lss,2,'TrillRegions');  
peaks = getLabelValues(lss,2,{'TrillRegions','TrillPeaks'});  
  
sg = getSignal(lss,2);  
plot((0:length(sg)-1)/lss.SampleRate,sg)  
xlim(tvals.ROIlimits)  
hold on  
plot(peaks.Location,cell2mat(peaks.Value),'v')  
hold off
```



Display the coordinates of the third trill peak.


```
pcoor = getLabelValues(lss,2,{'TrillRegions','TrillPeaks'}, ...
    'LabelRowIndex',1,'SublabelRowIndex',3)
```

```
pcoor=1x2 table
    Location      Value
    _____  _____
    11.437        [0.1500]
```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: labeledSignalSet({randn(100,1),randn(10,1)},signalLabelDefinition('female')) specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. midx specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: signalLabelDefinition("Asleep",'LabelType','roi') specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

ridx — Label row index

positive integer

Label row index, specified as a positive integer. This argument applies only for ROI and point labels.

sridx — Sublabel row index

positive integer

Sublabel row index, specified as a positive integer. This argument applies only when a label and sublabel pair has been specified in `lblname` and the sublabel is of type ROI or point.

Output Arguments

val — Label values

table

Label values, returned as a table.

sublbltbl — Sublabel values

table

Sublabel values, returned as a table showing the structure of the label value and its sublabel variables.

- If `lblname` has no sublabels, then `sublbltbl` is empty.
- If you specify `lblname` as a string or cell array, then `sublbltbl` is empty.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

getMemberNames

Get member names in labeled signal set

Syntax

```
mnames = getMemberNames(lss)
```

Description

`mnames = getMemberNames(lss)` returns a string array containing the member names in the order in which they are stored in the labeled signal set `lss`.

Examples

Get Member Names

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
      NumMembers: 2  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [2x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Return a string array with the names of the members.

```
getMemberNames(lss)
```

```
ans = 2x1 string array  
    "Member{1}"  
    "Member{2}"
```

Set the names of the set members to the whales' nicknames.

```
setMemberNames(lss, {'Brutus' 'Lucy'})
```

Verify that the members have the nicknames as names.

```
getMemberNames(lss)
```

```
ans = 2x1 string array  
    "Brutus"  
    "Lucy"
```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1), randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

Output Arguments

mnames — Member names

string array

Member names, returned as a string array.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

getSignal

Get signals from labeled signal set

Syntax

```
[s,info] = getSignal(lss,midX)
```

Description

[s,info] = getSignal(lss,midX) returns the values for the signals contained in member midX of the labeled signal set lss.

Examples

Get Signal

Load a labeled signal set containing recordings of whale songs.

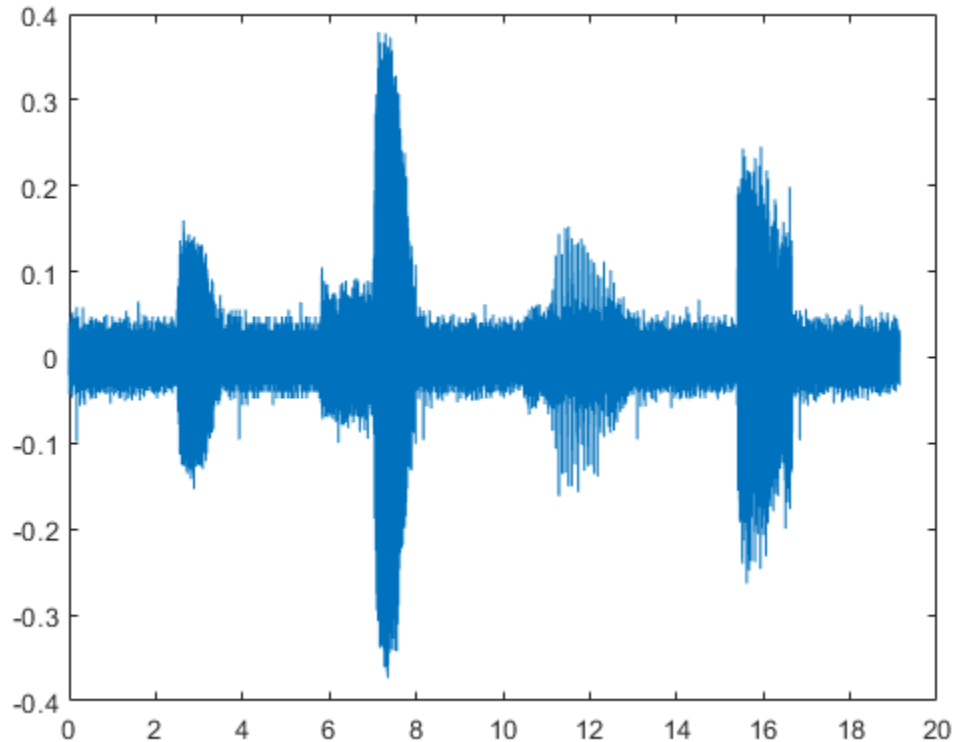
```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
    NumMembers: 2  
  TimeInformation: "sampleRate"  
    SampleRate: 4000  
      Labels: [2x3 table]  
  Description: "Characterize wave song regions"
```

Use labelDefinitionsHierarchy to see a list of labels and sublabels.
Use setLabelValue to add data to the set.

Retrieve the second member of the set and plot it.

```
[song,tinfo] = getSignal(lss,2);  
t = (0:length(song)-1)/tinfo.SampleRate;  
plot(t,song)
```



Input Arguments

lss — Labeled signal set
labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

Output Arguments

s — Signal values

vector | matrix | timetable | cell array

Signal values, returned as vector, matrix, timetable, or cell array.

info — Time information

structure

Time information, returned as a structure.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

haart

Haar 1-D wavelet transform

Syntax

```
[a,d] = haart(x)
[a,d] = haart(x,level)
[a,d] = haart( ____,integerflag)
```

Description

`[a,d] = haart(x)` returns the approximation coefficients, `a`, and detail coefficients, `d`, of a 1-D Haar discrete wavelet transform. The input `x` can be univariate or multivariate data. The default `level` depends on the length of `x`.

`[a,d] = haart(x,level)` obtains the Haar transform down to the specified level.

`[a,d] = haart(____,integerflag)` specifies how the Haar transform handles integer-valued data, using any of the previous syntaxes.

Examples

Haar Transform of ECG Data

Obtain the Haar transform down to the default maximum level.

```
load wecg;
[a,d] = haart(wecg);
```

Haar Transform of Electricity Consumption Data Down to Specified Level

Obtain the Haar transform of a multivariate time series dataset of electricity consumption data down to level 4. The `signals` data is transposed so that each time series is in a column, rather than a row.

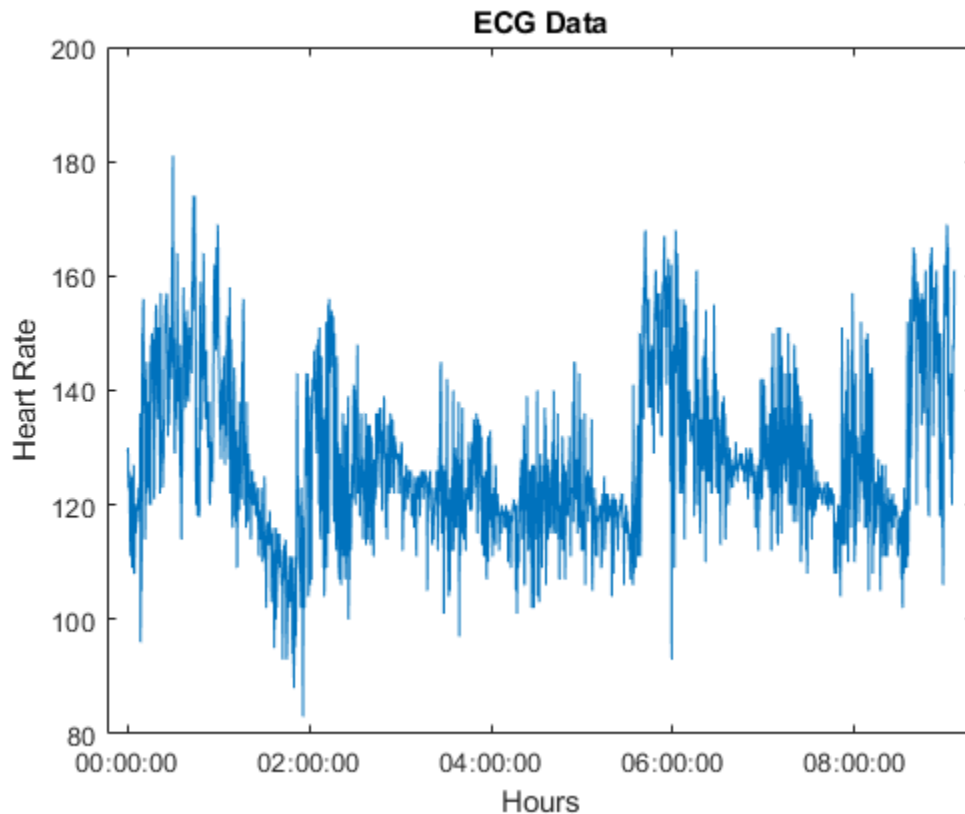
```
load elec35_nor;  
signals = signals';  
[a,d] = haart(signals,4);
```

Haar Transform of Integer Data Series

Obtain the Haar transform and inverse Haar transform of ECG heart rate data. The data is made up of integers only.

Load and plot the ECG data.

```
load BabyECGData;  
plot(times,HR)  
xlabel('Hours')  
ylabel('Heart Rate')  
title('ECG Data')
```

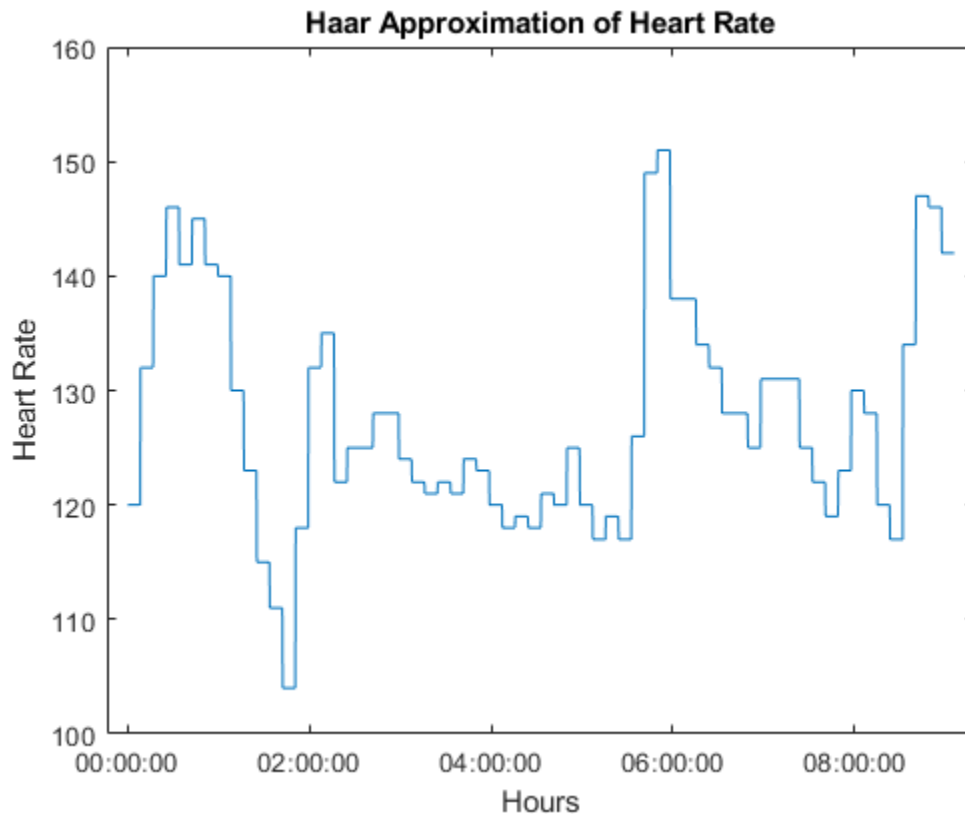


Obtain the Haar transform. Then, obtain the inverse Haar transform approximated at level 5. The scale for this level is 512 seconds, which is 2^5 times the sampling interval (16 seconds).

```
[a,d] = haart(HR,'integer');
HaarHR = ihaart(a,d,5,'integer');
```

Compare the reconstructed data to the original data.

```
figure;
plot(times,HaarHR)
xlabel('Hours')
ylabel('Heart Rate')
title('Haar Approximation of Heart Rate')
```



Input Arguments

x — Input signal

vector of real values | matrix of real values

Input signal, specified as vector or matrix of real values. If x is a vector, it must be even length. If x is a matrix, each column must be even length, and `haart` operates on each column of x .

level — Maximum level

positive integer

Maximum level to which to perform the Haar transform, specified as a positive integer. The default value depends on the length of the input signal, x .

- If the length of x is a power of two, the Haar transform is obtained down to level $\log_2(\text{length}(x))$.
- If the length of x is even, but not a power of two, the Haar transform is obtained down to level $\text{floor}(\log_2(\text{length}(x)/2))$.

If `level` is 1, the detail coefficients, d , are returned as a vector or matrix, depending on whether the input is a vector or matrix, respectively.

integerflag — Integer-valued data handling

'noninteger' (default) | 'integer'

Integer-valued data handling, specified as either 'noninteger' or 'integer'. 'noninteger' does not preserve integer-valued data in the Haar transform, and 'integer' preserves it. The 'integer' option applies only if all elements of the input, x , are integers. For integer-valued input, `haart` returns integer-valued wavelet coefficients. For both 'noninteger' and 'integer', however, the Haar transform algorithm uses floating-point arithmetic. The data type of outputs a and d , is always `double`.

Output Arguments

a — Approximation coefficients

scalar | vector | matrix

Approximation coefficients, returned as a scalar, vector, or matrix of coefficients, depending on the level to which the transform is calculated. Approximation, or scaling, coefficients are a lowpass representation of the input. At each level, the approximation coefficients are divided into coarser approximation and detail coefficients.

Data Types: `double`

d — Detail coefficients

scalar | vector | matrix | cell array

Detail coefficients, returned as a scalar, vector, matrix, or cell array. Detail coefficients are generally referred to as the wavelet coefficients and are a highpass representation of the input. The number of detail coefficients depends on the selected level and the length of

the input. The order of the elements of `d` is from fine to coarse resolution levels. The coarsest resolution level element of the `d` cell array is a scalar value. If you specify only two levels, the detail coefficient is a scalar.

Data Types: `double`

See Also

`haart2` | `ihaart` | `ihaart2`

Topics

“Haar Transforms for Time Series Data and Images”

Introduced in R2016b

haart2

2-D Haar wavelet transform

Syntax

```
[a,h,v,d] = haart2(x)
[a,h,v,d] = haart2(x,level)
[a,h,v,d] = haart2( ____,integerflag)
```

Description

`[a,h,v,d] = haart2(x)` performs the 2-D Haar discrete wavelet transform (DWT) of the matrix, `x`. `haart2` returns the approximation coefficients, `a`, at the coarsest level. `haart2` also returns cell arrays of matrices containing the horizontal, vertical, and diagonal detail coefficients by level. If the 2-D Haar transform is computed only at one level coarser in resolution, then `h`, `v`, and `d` are matrices. The default `level` depends on the number of rows of `x`.

`[a,h,v,d] = haart2(x,level)` performs the 2-D Haar transform down to the specified level.

`[a,h,v,d] = haart2(____,integerflag)` specifies how the 2-D Haar transform handles integer-valued data, using any of the previous syntaxes.

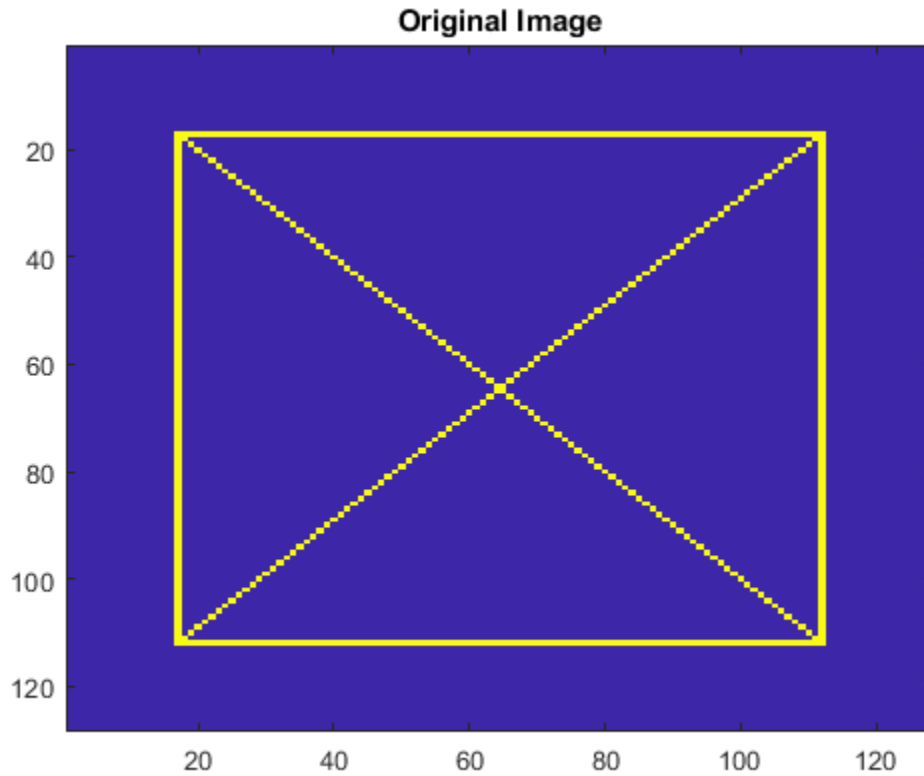
Examples

Haar Transform and First Level Details of 2-D Data

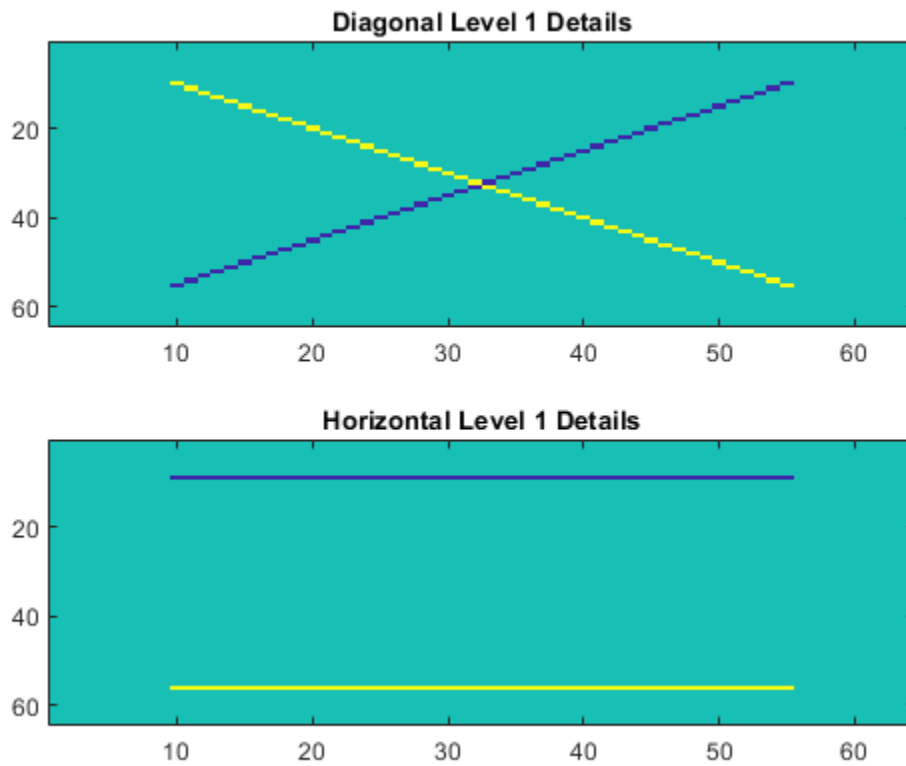
Obtain the 2-D Haar transform of 2-D data and plot its diagonal and horizontal level 1 details.

```
load xbox;
[a,h,v,d] = haart2(xbox);
```

```
imagesc(xbox)
title('Original Image')
```



```
figure
subplot(2,1,1)
imagesc(d{1})
title('Diagonal Level 1 Details')
subplot(2,1,2)
imagesc(h{1})
title('Horizontal Level 1 Details')
```

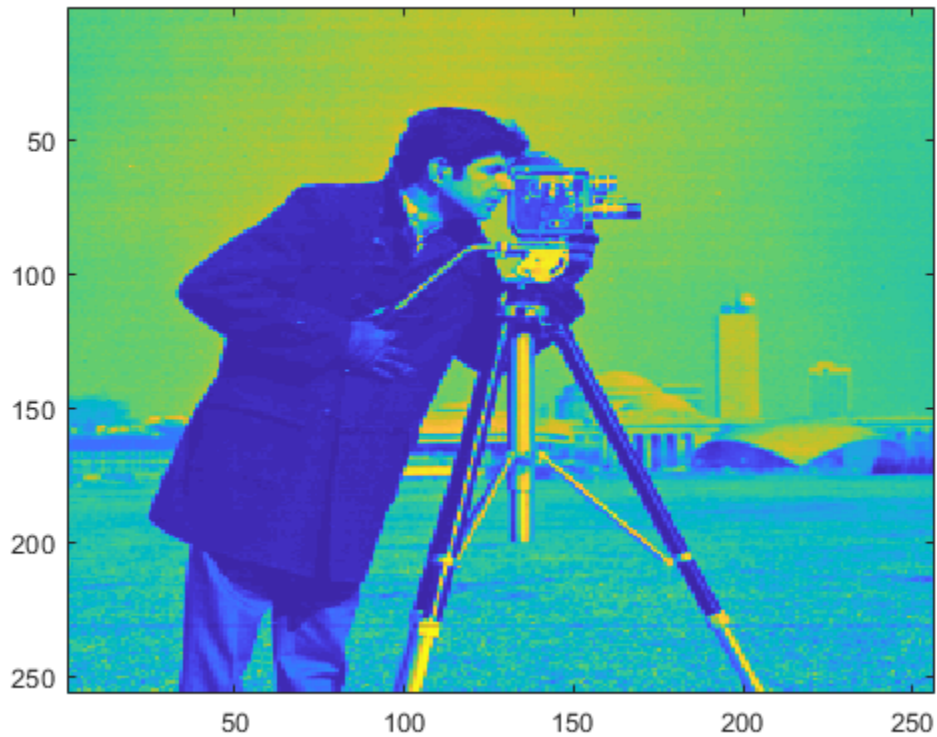



Haar Transform of Image Down to a Specified Level

Show the effect of limiting the maximum level of the 2-D Haar transform on an image.

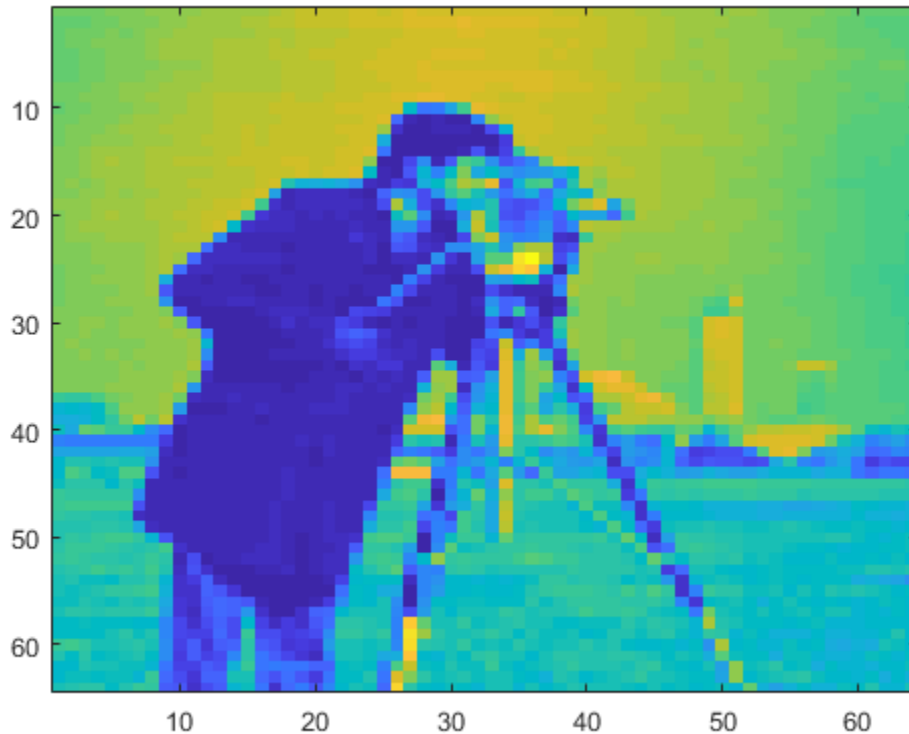
Load and display the image of a cameraman.

```
im = imread('cameraman.tif');  
imagesc(im)
```



Obtain the 2-D Haar transform to level 2 and view the level 2 approximation.

```
[a2,h2,v2,d2] = haart2(im,2);  
imagesc(a2)
```



Haar Transform Using Integer Image Data

Compare 2-D Haar transform results using the default 'noninteger' flag and the 'integer' flag. The cameraman image is uint8 data, so its maximum value is 255.

Obtain the default Haar transform. The approximation detail coefficient is outside the range 0 to 255.

```
im = imread('cameraman.tif');  
[a,h,v,d] = haart2(im);  
a
```

```
a = 3.0393e+04
```

Obtain the Haar transform, limiting it to integer values. The approximation detail is an integer and is within the range of the original image data.

```
[a,h,v,d] = haart2(im, 'integer');
```

```
a
```

```
a = 119
```

Input Arguments

x — Input signal

matrix of real values

Input signal, specified as a 2-D or 3-D matrix of real values. If **x** is 3-D, the third dimension of **x** must equal 3. The row and column sizes of **x** must be even length.

level — Maximum level

positive integer

Maximum level to which to perform the 2-D Haar transform, specified as a positive integer. The default value depends on the length of the input signal, **x**.

- If both the row and column sizes of **x** are powers of two, the 2-D Haar transform is obtained down to level $\log_2(\min(\text{size}(\mathbf{x})))$.
- If both the row and column sizes of **x** are even, but at least one is not a power of two, level is equal to $\text{floor}(\log_2(\min(\text{size}(\mathbf{x})/2))$.

If **level** is greater than 1, then **h**, **v**, and **d** are cell arrays. If **level** is equal to 1, then **h**, **v**, and **d** are matrices.

integerflag — Integer-valued data handling

'noninteger' (default) | 'integer'

Integer-valued data handling, specified as either 'noninteger' or 'integer'. 'noninteger' does not preserve integer-valued data in the 2-D Haar transform, and 'integer' preserves it. The 'integer' option applies only if all elements of the input, **x**, are integers. For integer-valued input, **haart2** returns integer-valued wavelet coefficients. For both 'noninteger' and 'integer', however, the 2-D Haar transform

algorithm uses floating-point arithmetic. The data type of outputs **a**, **h**, **v**, and **d**, is always double.

Output Arguments

a — Approximation coefficients

scalar | matrix

Approximation coefficients, returned as a scalar or matrix of coefficients, depending on the level to which the transform is calculated. Approximation, or scaling, coefficients are a lowpass representation of the input. At each level, the approximation coefficients are divided into coarser approximation and detail coefficients.

Data Types: double

h — Horizontal detail coefficients

matrix | cell array

Horizontal detail coefficients by level, returned as a matrix or cell array of matrices. If `level` is greater than 1, **h** is a cell array. If `level` is equal to 1, the 2-D Haar transform is computed at only one level coarser in resolution and **h** is a matrix.

Data Types: double

v — Vertical detail coefficients

matrix | cell array

Vertical detail coefficients by level, returned as a matrix or cell array of matrices. If `level` is greater than 1, **v** is a cell array. If `level` is equal to 1, the 2-D Haar transform is computed at only one level coarser in resolution and **v** is a matrix.

Data Types: double

d — Diagonal detail coefficients

matrix | cell array

Diagonal detail coefficients by level, returned as a matrix or cell array of matrices. If `level` is greater than 1, **d** is a cell array. If `level` is equal to 1, the 2-D Haar transform is computed at only one level coarser in resolution and **d** is a matrix.

Data Types: double

See Also

haart | ihaart | ihaart2

Topics

“Haar Transforms for Time Series Data and Images”

Introduced in R2016b

head

Get top rows of labels table

Syntax

```
val = head(lss)
```

Description

`val = head(lss)` returns the top rows of the labels table of the labeled signal set `lss`.

Examples

Top Label Values

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
    Source: {2x1 cell}  
    NumMembers: 2  
    TimeInformation: "sampleRate"  
    SampleRate: 4000  
    Labels: [2x3 table]  
    Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Get the top rows of the labels table.

```
head(lss)
```

```
ans=2x3 table
```

	WhaleType	MoanRegions	TrillRegions
Member{1}	blue	[3x2 table]	[1x3 table]
Member{2}	blue	[3x2 table]	[1x3 table]

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1),randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

Output Arguments

val — Top rows of labels

table

Top rows of labels, returned as a table.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

hht

Hilbert-Huang transform

Syntax

```
hs = hht(imf,fs)
[hs,f,t] = hht(imf,fs)
[hs,f,t,imfinsf,imfinse] = hht( ___ )
[ ___ ] = hht( ___ ,Name,Value)

hht( ___ )
hht( ___ ,freqlocation)
```

Description

`hs = hht(imf,fs)` returns the Hilbert spectrum `hs` of the signal specified by intrinsic mode functions `imf`, and the optional sampling frequency `fs`. `hs` is useful for analyzing signals that comprise a mixture of signals whose spectral content changes in time. Use `hht` to perform Hilbert spectral analysis on signals to identify localized features.

`[hs,f,t] = hht(imf,fs)` returns frequency vector `f` and time vector `t` in addition to `hs`.

`[hs,f,t,imfinsf,imfinse] = hht(___)` also returns the instantaneous frequencies `imfinsf` and the instantaneous energies `imfinse` of the intrinsic mode functions for signal diagnostics.

`[___] = hht(___ ,Name,Value)` estimates Hilbert spectrum parameters with additional options specified by one or more `Name,Value` pair arguments.

`hht(___)` with no output arguments plots the Hilbert spectrum in the current figure window. You can use this syntax with any of the input arguments in previous syntaxes.

`hht(___ ,freqlocation)` plots the Hilbert spectrum with the optional `freqlocation` argument to specify the location of the frequency axis. Frequency is represented on the `y`-axis by default.

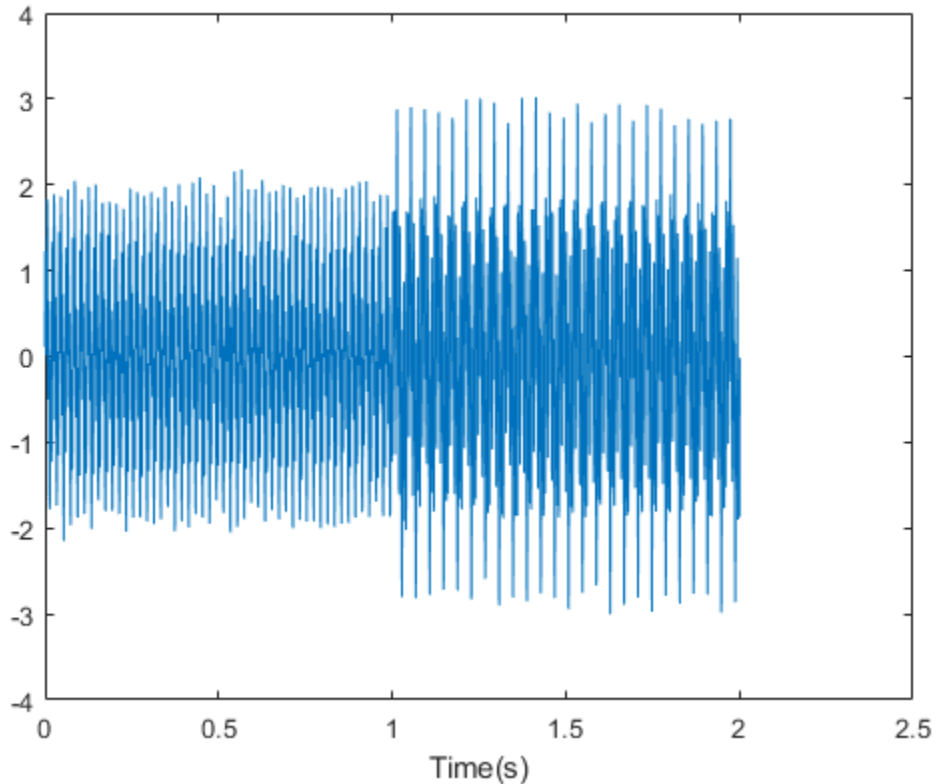
Examples

Perform Empirical Mode Decomposition and Visualize Hilbert Spectrum of Signal

For this example, consider a nonstationary continuous signal composed of sinusoidal waves with a distinct change in frequency. The vibration of a jackhammer and the sound of fireworks are examples of nonstationary continuous signals.

Load the nonstationary signal data with the sampling frequency `fs`, and visualize the mixed sinusoidal signal.

```
load('sinusoidalSignalExampleData.mat','X','fs')
t = (0:length(X)-1)/fs;
plot(t,X)
xlabel('Time(s)')
```



The mixed signal contains sinusoidal waves with different amplitude and frequency values.

To create the Hilbert spectrum plot, you need the intrinsic mode functions (IMFs) of the signal. Perform empirical mode decomposition to compute the IMFs and residuals of the signal. Since the signal is not smooth, specify 'pchip' as the Interpolation method.

```
[imf,residual,info] = emd(X,'Interpolation','pchip');
```

Current IMF	#Sift Iter	Relative Tol	Stop Criterion Hit
1	2	0.026352	SiftMaxRelativeTolerance
2	2	0.0039573	SiftMaxRelativeTolerance
3	1	0.024838	SiftMaxRelativeTolerance
4	2	0.05929	SiftMaxRelativeTolerance

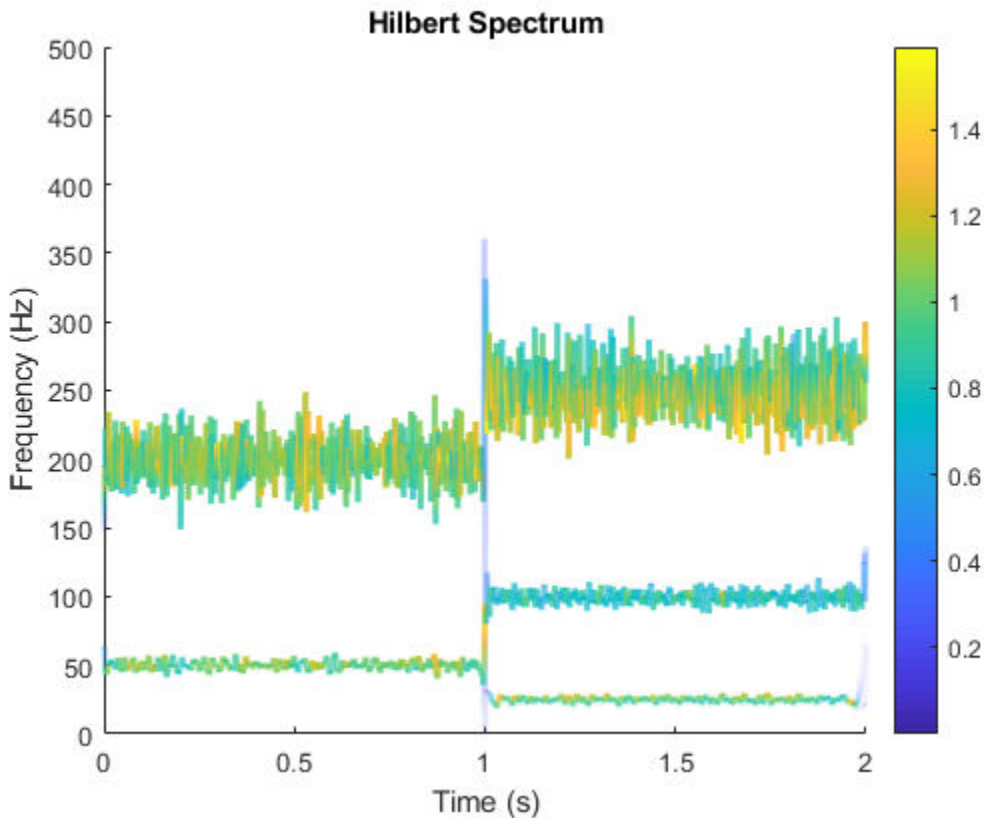
5		2		0.11317		SiftMaxRelativeTolerance
6		2		0.12599		SiftMaxRelativeTolerance
7		2		0.13802		SiftMaxRelativeTolerance
8		3		0.15937		SiftMaxRelativeTolerance
9		2		0.15923		SiftMaxRelativeTolerance

The decomposition stopped because the number of extrema of the residual signal is less

The table generated in the command window indicates the number of sift iterations, the relative tolerance, and the sift stop criterion for each generated IMF. This information is also contained in `info`. You can hide the table by adding the `'Display', 0` name value pair.

Create the Hilbert spectrum plot using the `imf` components obtained using empirical mode decomposition.

```
hht(imf, fs)
```



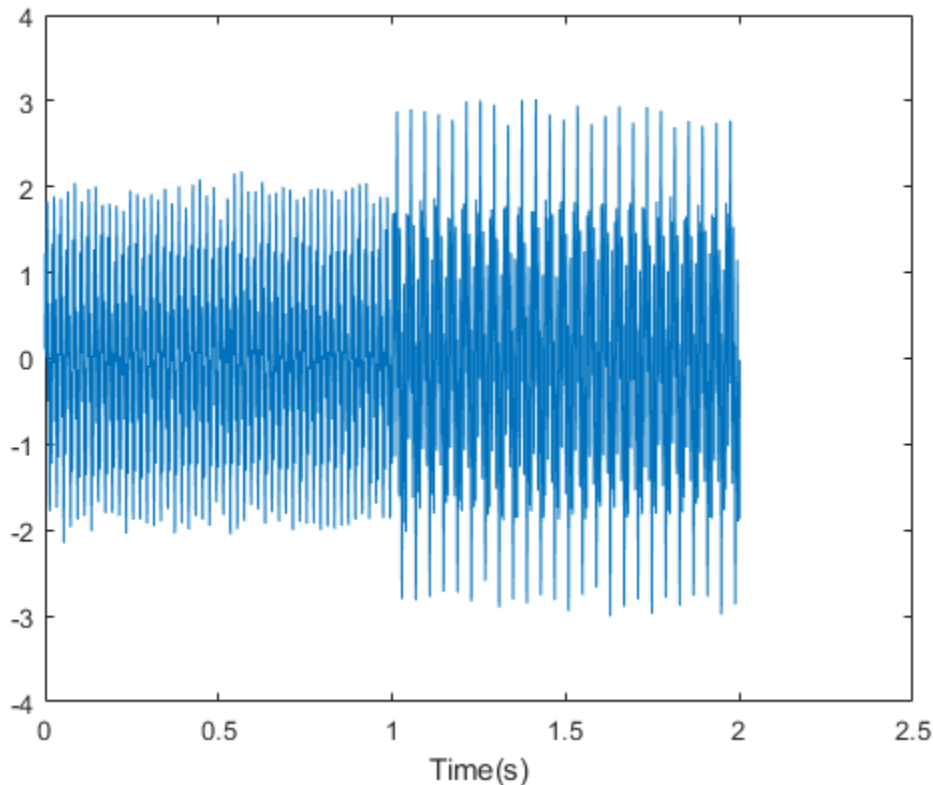
The frequency versus time plot is a sparse plot with a vertical color bar indicating the instantaneous energy at each point in the IMF. The plot represents the instantaneous frequency spectrum of each component decomposed from the original mixed signal. Three IMFs appear in the plot with a distinct change in frequency at 1 second.

Compute Hilbert Spectrum Parameters of Signal

For this example, consider a nonstationary continuous signal composed of sinusoidal waves with a distinct change in frequency. The vibration of a jackhammer and the sound of fireworks are examples of nonstationary continuous signals.

Load the nonstationary signal data with the sampling frequency fs , and visualize the mixed sinusoidal signal.

```
load('sinusoidalSignalExampleData.mat','X','fs')
t = (0:length(X)-1)/fs;
plot(t,X)
xlabel('Time(s)')
```



The mixed signal contains sinusoidal waves with different amplitude and frequency values.

To compute the Hilbert spectrum parameters, you need the IMFs of the signal. Perform empirical mode decomposition to compute the intrinsic mode functions and residuals of the signal. Since the signal is not smooth, specify 'pchip' as the Interpolation method.

```
[imf,residual,info] = emd(X,'Interpolation','pchip');
```

Current IMF	#Sift Iter	Relative Tol	Stop Criterion Hit
1	2	0.026352	SiftMaxRelativeTolerance
2	2	0.0039573	SiftMaxRelativeTolerance
3	1	0.024838	SiftMaxRelativeTolerance
4	2	0.05929	SiftMaxRelativeTolerance
5	2	0.11317	SiftMaxRelativeTolerance
6	2	0.12599	SiftMaxRelativeTolerance
7	2	0.13802	SiftMaxRelativeTolerance
8	3	0.15937	SiftMaxRelativeTolerance
9	2	0.15923	SiftMaxRelativeTolerance

The decomposition stopped because the number of extrema of the residual signal is less

The table generated in the command window indicates the number of sift iterations, the relative tolerance, and the sift stop criterion for each generated IMF. This information is also contained in `info`. You can hide the table by specifying `Display` as `0`.

Compute the Hilbert spectrum parameters: Hilbert spectrum `hs`, frequency vector `fs`, time vector `t`, instantaneous frequency `imfinsf`, and instantaneous energy `imfinse`.

```
[hs,f,t,imfinsf,imfinse] = hht(imf,fs);
```

Use the computed Hilbert spectrum parameters for time-frequency analysis and signal diagnostics.

Input Arguments

imf — Intrinsic mode function

matrix | timetable

Intrinsic mode function, specified as a matrix or timetable. `imf` is any signal whose envelope is symmetric with respect to zero and whose numbers of extrema and zero crossings differ by at most one. `emd` is used to decompose and simplify complicated signals into a finite number of intrinsic mode functions required to perform Hilbert spectral analysis.

`hht` treats each column in `imf` as an intrinsic mode function. For more information on computing `imf`, see `emd`.

fs — Sampling frequency 2π (default) | positive scalar

Sampling frequency, specified as a positive scalar. Sampling frequency or sampling rate is the number of samples per unit time.

If `fs` is not supplied, a normalized frequency of 2π is used to compute the Hilbert spectrum. If `imf` is specified as a timetable, the sampling time is inferred from it.

freqlocation — Location of frequency axis on plot

'yaxis' (default) | 'xaxis'

Location of frequency axis on the plot, specified as 'yaxis' or 'xaxis'. To display frequency data on the y-axis or x-axis of the plot, specify `freqlocation` as 'yaxis' or 'xaxis' respectively.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: 'FrequencyResolution', 1

FrequencyLimits — Frequency limits to compute Hilbert spectrum $[0, fs/2]$ (default) | 1-by-2 integer-valued vector

Frequency limits to compute Hilbert spectrum, specified as the comma-separated pair consisting of 'FrequencyLimits' and a 1-by-2 integer-valued vector. `FrequencyLimits` is specified in Hz.

FrequencyResolution — Frequency resolution to discretize frequency range $(f_high - f_low)/100$ (default) | positive scalar

Frequency resolution to discretize frequency limits, specified as the comma-separated pair consisting of 'FrequencyResolution' and a positive scalar.

Specify `FrequencyResolution` in Hz. If 'FrequencyResolution' is not specified, a value of $(f_high - f_low)/100$ is inferred from `FrequencyLimits`. Here, `f_high` is the upper limit of `FrequencyLimits`, while `f_low` is the lower limit.

MinThreshold — Minimum threshold value of Hilbert spectrum

-inf (default) | scalar

Minimum threshold value of Hilbert spectrum, specified as the comma-separated pair consisting of 'MinThreshold' and a scalar.

MinThreshold sets elements of `hs` to 0 when the corresponding elements of $10\log_{10}(hs)$ are less than MinThreshold.

Output Arguments

hs — Hilbert spectrum of signal

sparse array

Hilbert spectrum of the signal, returned as a sparse array. Use `hs` for time-frequency analysis and to identify localized features in the signal.

f — Frequency vector of signal

array

Frequency vector of the signal, returned as a numeric array. `hht` uses the frequency vector `f` and the time vector `t` to create the Hilbert spectrum plot.

Mathematically, `f` is denoted as:

$$f = f_low:FrequencyResolution:f_high$$
t — Time vector of signal

array | duration array

Time vector of the signal, returned as a numeric array or a duration array. `hht` uses the time vector `t` and the frequency vector `f` to create the Hilbert spectrum plot.

`t` is returned as:

- An array, if `imf` is specified as an array.
- A duration array, if `imf` is specified as a uniformly sampled timetable.

imfinsf — Instantaneous frequency of each IMF

array | timetable

Instantaneous frequency of each IMF, returned as an array or timetable.

`insf` has the same number of columns as `imf` and is returned as:

- An array, if `imf` is specified as an array.
- A timetable, if `imf` is specified as a uniformly sampled timetable.

imfinse — Instantaneous energy of each IMF

array | timetable

Instantaneous energy of each IMF, returned as an array or timetable.

`inse` has the same number of columns as `imf` and is returned as:

- An array, if `imf` is specified as an array.
- A timetable, if `imf` is specified as a uniformly sampled timetable.

Algorithms

The Hilbert-Huang transform is useful for performing time-frequency analysis of nonstationary and nonlinear data. The Hilbert-Huang procedure consists of the following steps:

- 1** `emd` decomposes the data set x into a finite number of intrinsic mode functions.
- 2** For each intrinsic mode function, x_i , the function `hht`:
 - a** Uses `hilbert` to compute the analytic signal, $z_i(t) = x_i(t) + jH\{x_i(t)\}$, where $H\{x_i\}$ is the Hilbert transform of x_i .
 - b** Expresses z_i as $z_i(t) = a_i(t) e^{j\theta_i(t)}$, where $a_i(t)$ is the instantaneous amplitude and $\theta_i(t)$ is the instantaneous phase.
 - c** Computes the instantaneous energy, $|a_i(t)|^2$, and the instantaneous frequency, $\omega_i(t) \equiv d\theta_i(t)/dt$. If given a sample rate, `hht` converts $\omega_i(t)$ to a frequency in Hz.
 - d** Outputs the instantaneous energy in `imfinse` and the instantaneous frequency in `imfinsf`.
- 3** When called with no output arguments, `hht` plots the energy of the signal as a function of time and frequency, with color proportional to amplitude.

References

- [1] Huang, Norden E., and Samuel S. P. Shen. *Hilbert-Huang Transform and Its Applications*. Singapore: World Scientific, 2014.
- [2] Huang, Norden E., Zhaohua Wu, Steven R. Long, Kenneth C. Arnold, Xianyao Chen, and Karin Blank. "On Instantaneous Frequency." *Advances in Adaptive Data Analysis*. Vol. 1, No. 2, 2009, pp. 177-229.

See Also

emd

Introduced in R2018a

icqt

Inverse constant-Q transform using nonstationary Gabor frames

Syntax

```
xrec = icqt(cfs,g,fshifts)
xrec = icqt( ____, 'SignalType', sigtype)
[xrec,gdual] = icqt( ____ )
```

Description

`xrec = icqt(cfs,g,fshifts)` returns the inverse constant-Q transform, `xrec`, of the coefficients `cfs`. `cfs` is a matrix, cell array, or structure array. `g` is the cell array of nonstationary Gabor constant-Q analysis filters used to obtain the coefficients `cfs`. `fshifts` is a vector of frequency bin shifts for the constant-Q bandpass filters in `g`. `icqt` assumes by default that the original signal was real-valued. To indicate the original input signal was complex-valued, use the 'SignalType' name-value pair. If the input to `cqt` was a single signal, then `xrec` is a vector. If the input to `cqt` was a multichannel signal, then `xrec` is a matrix. `cfs`, `g`, and `fshifts` must be outputs of `cqt`.

`xrec = icqt(____, 'SignalType', sigtype)` designates whether the signal was real-valued or complex-valued. Valid options for `sigtype` are 'real' or 'complex'. If unspecified, `sigtype` defaults to 'real'.

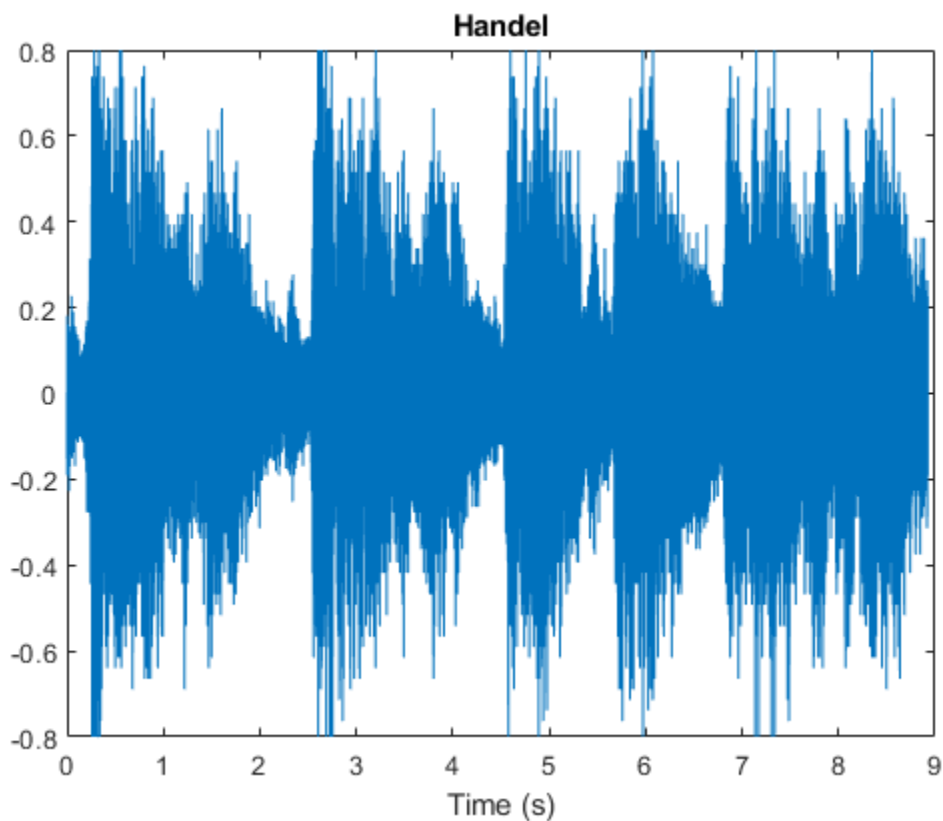
`[xrec,gdual] = icqt(____)` returns the dual frames of `xrec` as a cell array the same size as `g`. The dual frames are the canonical dual frames derived from the analysis filters.

Examples

Perfect Reconstruction of Constant-Q Transform

Load and plot the Handel signal.

```
load handel
t = (0:length(y)-1)/Fs;
plot(t,y)
title('Handel')
xlabel('Time (s)')
```



Obtain the constant-Q transform of the signal using the `sparse` transform option. Because the transform will be inverted, you must also return the Gabor frames and frequency shifts used in the analysis.

```
[cfs,~,g,fshifts] = cqt(y, 'SamplingFrequency',Fs, 'TransformType', 'sparse');
```

Invert the constant-Q transform and demonstrate perfect reconstruction by showing the maximum absolute reconstruction error and the relative energy error in dB.

```
xrec = icqt(cfs,g,fshifts);  
maxAbsError = max(abs(xrec-y))  
  
maxAbsError = 7.6328e-16  
  
relEnergyError = 20*log10(norm(xrec-y)/norm(y))  
  
relEnergyError = -301.4461
```

Input Arguments

cfs — Constant-Q coefficients

matrix | cell array | structure array

Constant-Q coefficients of a signal or multichannel signal, specified as a matrix, cell array, or structure array. `cfs` must be the output of `cqt`.

g — Nonstationary Gabor constant-Q analysis filters

cell array

Nonstationary Gabor constant-Q analysis filters used to obtain the coefficients `cfs`, specified as a cell array. `cfs` must be the output of `cqt`.

fshifts — Frequency bin shifts

real-valued vector

Frequency bin shifts for the constant-Q bandpass filters in `g`, specified as a real-valued vector. `fshifts` must be the output of `cqt`.

sigtype — Signal type

'real' (default) | 'complex'

Signal type of the original signal, specified as 'real' or 'complex'. Use `sigtype` to designate whether the original signal was real-valued or complex-valued. If unspecified, `sigtype` defaults to 'real'.

Output Arguments

xrec — Inverse constant-Q transform

vector | matrix

Inverse constant-Q transform, returned as a vector or matrix. If the input to `cqt` was a single signal, then `xrec` is a vector. If the input to `cqt` was a multichannel signal, then `xrec` is a matrix.

gdual — Dual frames

cell array

Dual frames used in the synthesis of `xrec`, returned as a cell array the same size as `g`. The dual frames are the canonical dual frames derived from the analysis filters.

Algorithms

The theory of nonstationary Gabor (NSG) frames for frequency-adaptive analysis and efficient algorithms for analysis and synthesis using NSG frames are due to Dörfler, Holighaus, Grill, and Velasco [1],[2]. The algorithms used in `cqt` and `icqt` were developed by Dörfler, Holighaus, Grill, and Velasco and are described in [1],[2]. In [3], Schörkhuber, Klapuri, Holighaus, and Dörfler develop and provide algorithms for a phase-corrected CQT transform which matches the CQT coefficients that would be obtained by naive convolution. The Large Time-Frequency Analysis Toolbox (<https://github.com/ltfat>) provides an extensive suite of algorithms for nonstationary Gabor frames [4].

References

- [1] Holighaus, N., M. Dörfler, G. A. Velasco, and T. Grill. "A framework for invertible real-time constant-Q transforms." *IEEE Transactions on Audio, Speech, and Language Processing*. Vol. 21, No. 4, 2013, pp. 775-785.
- [2] Velasco, G. A., N. Holighaus, M. Dörfler, and T. Grill. "Constructing an invertible constant-Q transform with nonstationary Gabor frames." In *Proceedings of the 14th International Conference on Digital Audio Effects (DAFx-11)*. Paris, France: 2011.
- [3] Schörkhuber, C., A. Klapuri, N. Holighaus, and M. Dörfler. "A Matlab Toolbox for Efficient Perfect Reconstruction Time-Frequency Transforms with Log-Frequency Resolution." Submitted to the *AES 53rd International Conference on Semantic Audio*. London, UK: 2014.
- [4] Průša, Z., P. L. Søndergaard, N. Holighaus, C. Wiesmeyer, and P. Balazs. *The Large Time-Frequency Analysis Toolbox 2.0*. Sound, Music, and Motion, Lecture Notes in Computer Science 2014, pp 419-442.

See Also

cqt

Topics

“Nonstationary Gabor Frames and the Constant-Q Transform”

Introduced in R2018a

icwt

Inverse continuous 1-D wavelet transform

Syntax

```
xrec = icwt(wt)
xrec = icwt(wt,wname)
xrec = icwt(wt,f,freqrage)
xrec = icwt(wt,period,periodrange)
xrec = icwt( ____,Name,Value)
```

Description

`xrec = icwt(wt)` inverts the continuous wavelet transform (CWT) coefficient matrix `wt` using default values. `icwt` assumes that you obtained the CWT using `cwt` with the default analytic Morse (3,60) wavelet. This wavelet has a symmetry of 3 and a time bandwidth of 60. `icwt` also assumes that the CWT uses default scales. If `wt` is a 2-D matrix, `icwt` assumes that the CWT was obtained from a real-valued signal. If `wt` is a 3-D matrix, `icwt` assumes that the CWT was obtained from a complex-valued signal. For a 3-D matrix, the first page of the `wt` is the CWT of the positive (counterclockwise) component and the second page of `wt` is the negative (clockwise) component. The pages represent the analytic and anti-analytic parts of the CWT, respectively.

`xrec = icwt(wt,wname)` uses the analytic wavelet `wname` to invert the CWT. The specified wavelet must be the same as the wavelet used in `cwt`. Valid options for `wname` are 'morse', 'amor', and 'bump', which specify the Morse, Morlet, and bump wavelet, respectively.

`xrec = icwt(wt,f,freqrage)` inverts the CWT over the frequency range specified in `freqrage`. `f` is the scale-to-frequency conversion obtained from `cwt`.

`xrec = icwt(wt,period,periodrange)` inverts the CWT over the range of periods specified in `periodrange`. `period` is an array of durations obtained from `cwt` with a duration input. The `period` is the `cwt` output obtained using a duration input. The period range must be increasing and contained in `period`.

`xrec = icwt(____, Name, Value)` returns the inverse CWT with additional options specified by one or more `Name, Value` pair arguments.

Examples

Inverse Continuous Wavelet Transform of Speech Signal

Obtain the CWT of a speech sample and invert the CWT using the default analytic Morse wavelet.

```
load mtlb;  
wt = cwt(mtlb);  
xrec = icwt(wt);
```

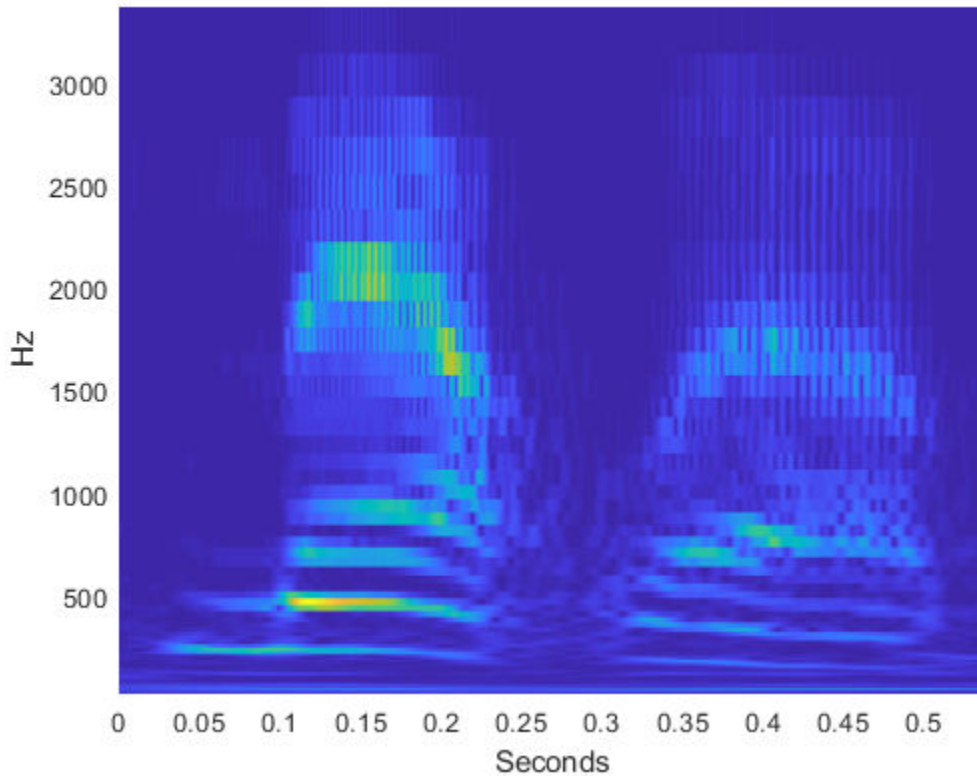
Inverse Continuous Wavelet Transform Using Specified Wavelet

Obtain the continuous wavelet transform of a speech sample and reconstruct the sample using the bump wavelet instead of the default Morse wavelet.

```
load mtlb;  
dt = 1/Fs;  
t = 0:dt:numel(mtlb)*dt-dt;
```

Obtain and plot the CWT.

```
[bumpmtlb, f] = cwt(mtlb, Fs, 'bump');  
p1 = pcolor(t, f, abs(bumpmtlb));  
p1.EdgeColor = 'none';  
xlabel('Seconds');  
ylabel('Hz');
```

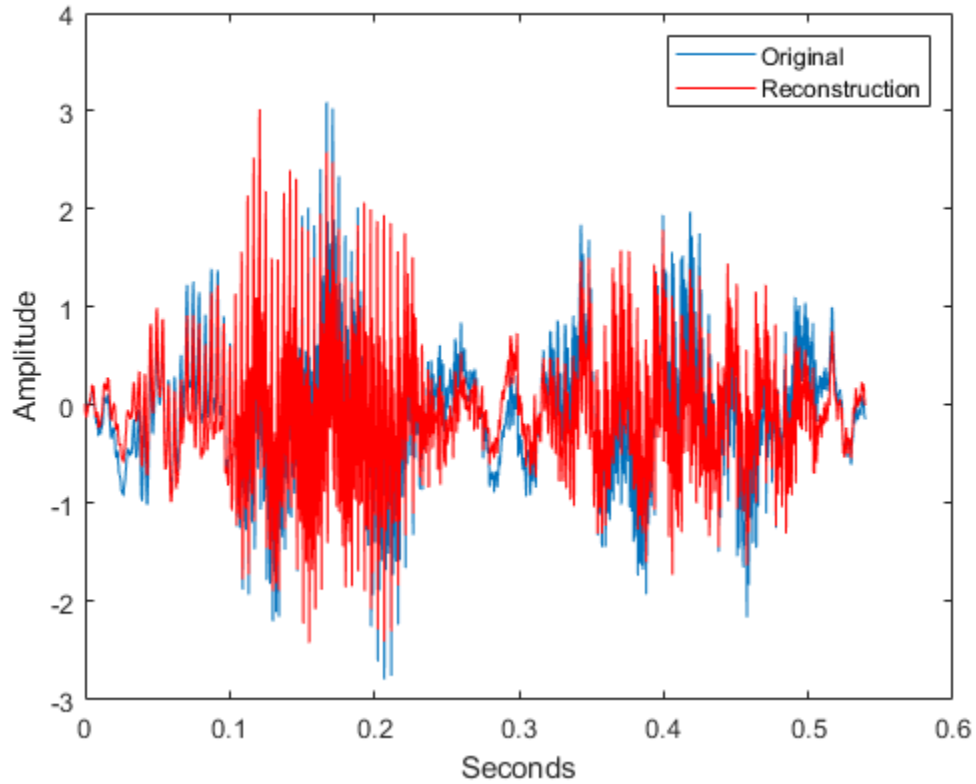


Obtain the inverse CWT.

```
xrec = icwt(bumpmtlb,'bump','SignalMean',mean(mtlb));
```

Plot the original and reconstructed signals.

```
plot(t,mtlb);  
xlabel('Seconds');  
ylabel('Amplitude');  
hold on;  
plot(t,xrec,'r');  
legend('Original','Reconstruction');
```



Play and compare the original and reconstructed signals.

```
p = audioplayer(mtlb,Fs);  
play(p);  
pause(2);  
px = audioplayer(xrec,Fs);  
play(px);
```

Reconstruct Frequency-Localized Data

Reconstruct a frequency-localized approximation to the Kobe earthquake data by extracting information from the CWT. The extracted information corresponds to frequencies in the range [0.030 0.070] Hz.

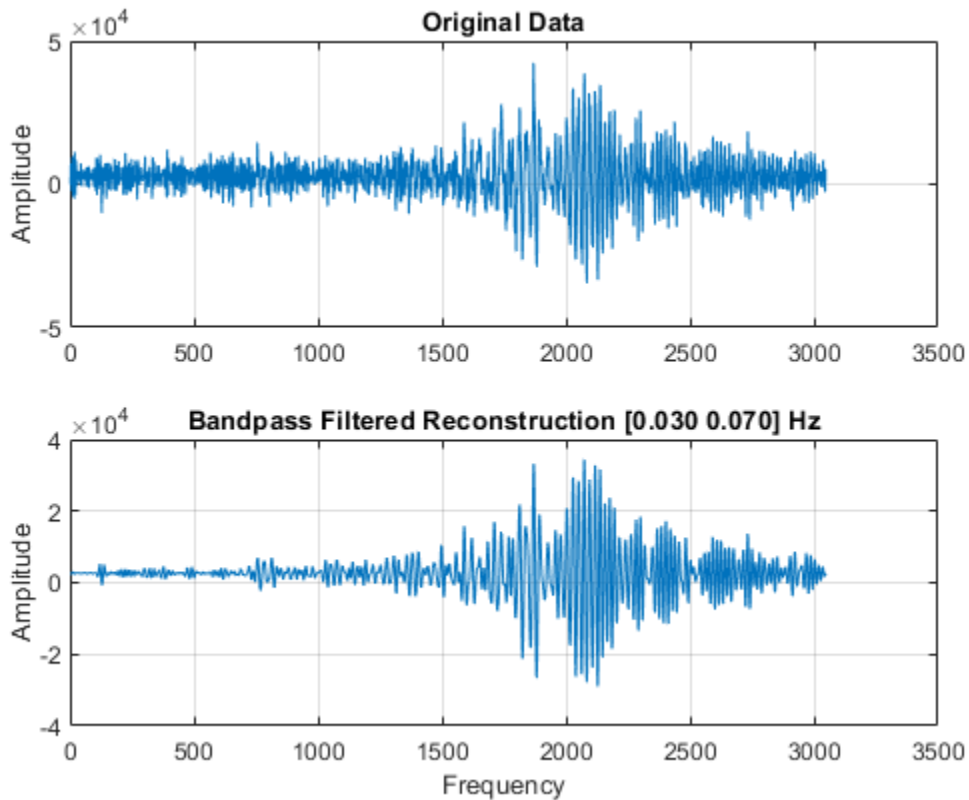
```
load kobe;
```

Obtain the CWT. Then, obtain the inverse CWT and add the signal mean back into the reconstructed data. The CWT does not preserve the signal mean.

```
[wt,f] = cwt(kobe,1);  
xrec = icwt(wt,f,[0.030 0.070], 'SignalMean', mean(kobe));
```

Plot the original and reconstructed data.

```
subplot(211)  
plot(kobe);  
grid on  
title('Original Data');  
ylabel('Amplitude')  
  
subplot(212)  
plot(xrec);  
grid on  
title('Bandpass Filtered Reconstruction [0.030 0.070] Hz');  
xlabel('Frequency');  
ylabel('Amplitude');
```



Reconstruct Data from Specific Time Period

Use the inverse continuous wavelet transform to reconstruct an approximation to El Nino data based on 2 to 8 year periods.

Load the El Nino data and obtain its CWT. The data is sampled monthly. To obtain the periods in years, specify the sampling interval as 1/12 of a year.

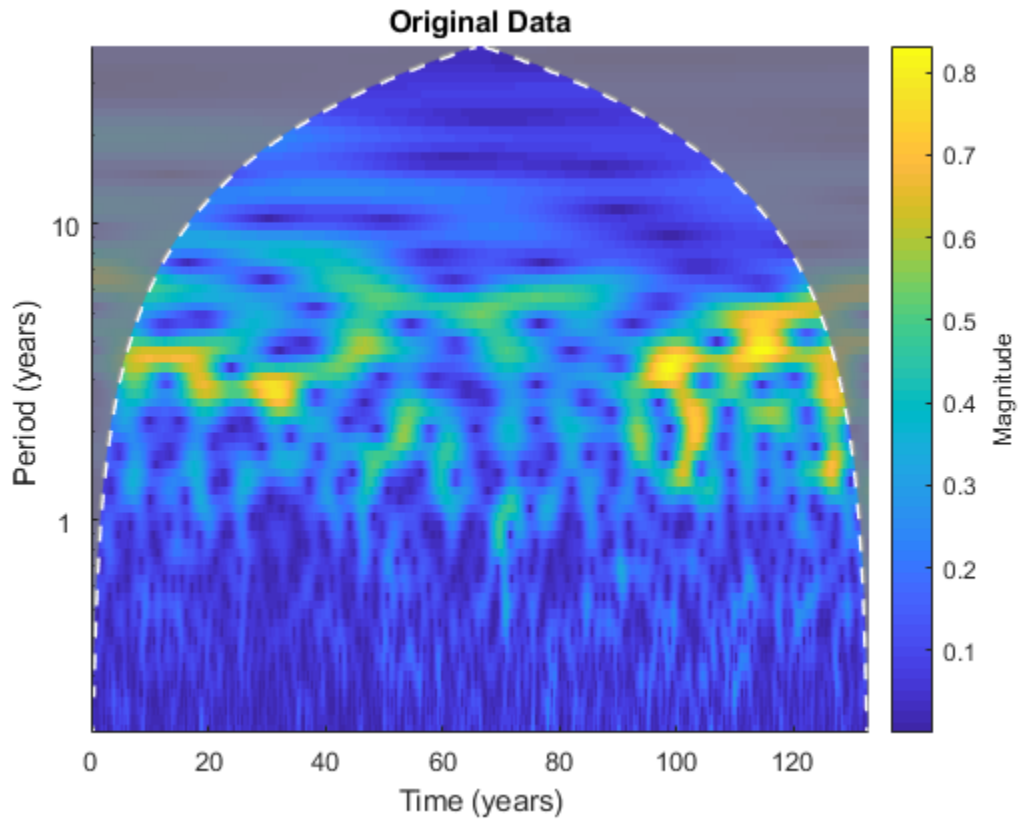
```
load ninoairdata;  
[cfs,period] = cwt(nino,years(1/12));
```

Obtain the inverse CWT for periods of 2 to 8 years.

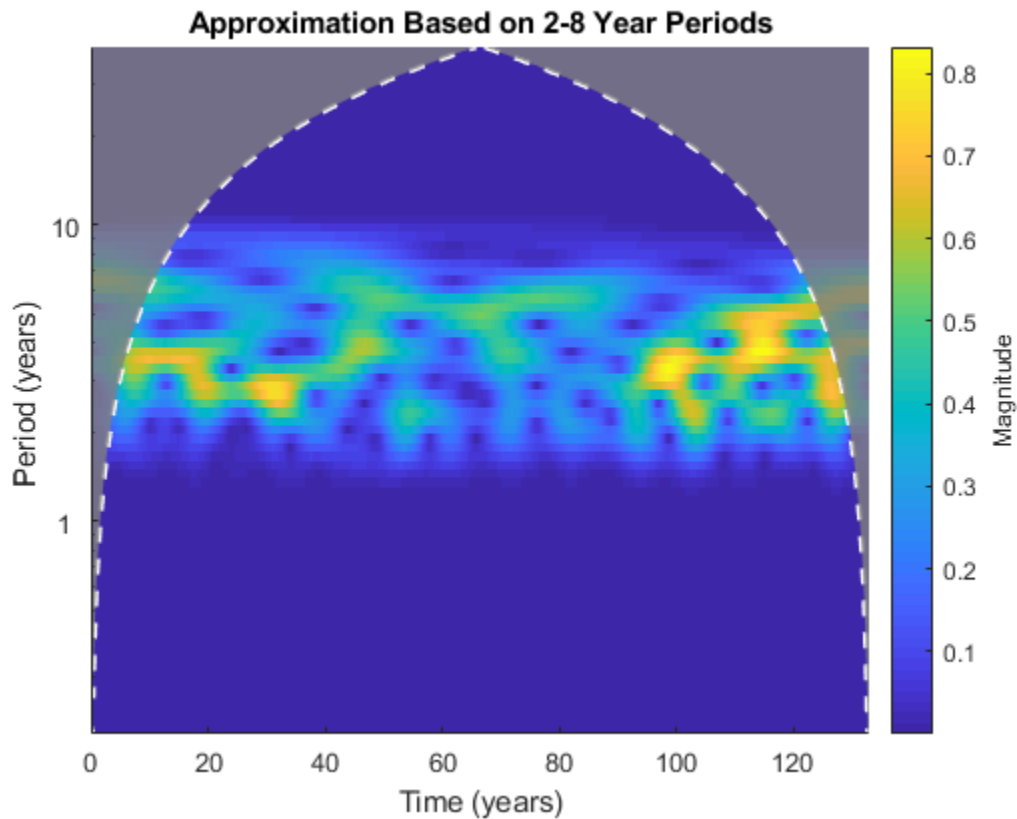
```
xrec = icwt(cfs,period,[years(2) years(8)]);
```

Plot the CWT of the reconstructed data and compare it to the CWT of the original data.

```
cwt(nino,years(1/12)); title('Original Data');
```



```
figure;  
cwt(xrec,years(1/12)); title('Approximation Based on 2-8 Year Periods');
```

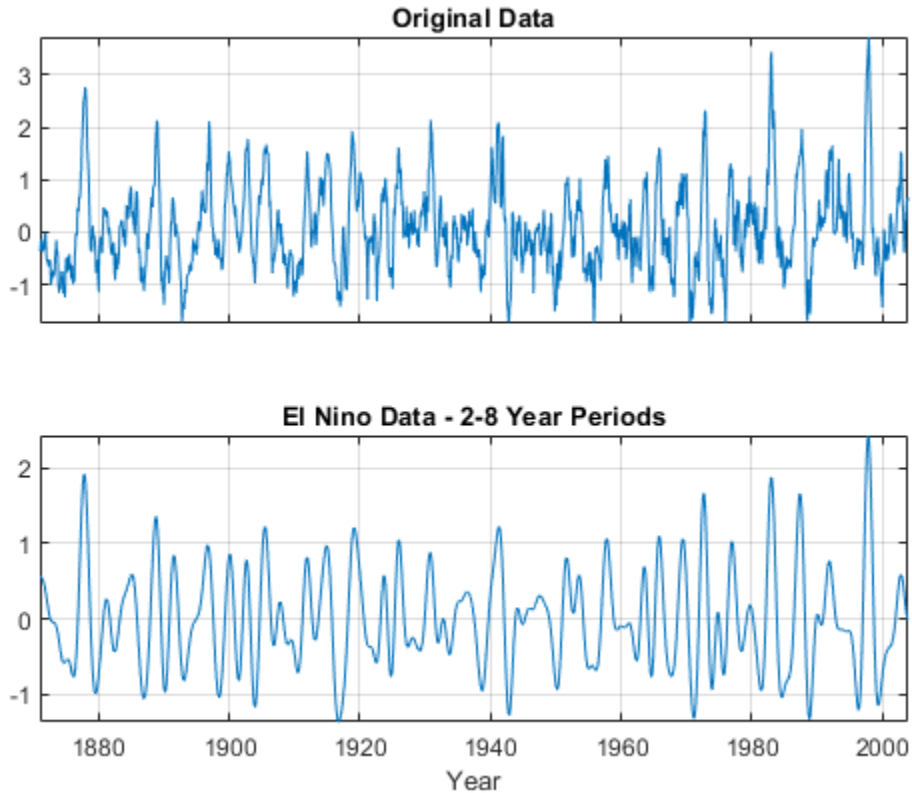


Compare the original data with the reconstructed data in time.

```
figure;  
subplot(211)  
plot(datayear,nino);  
grid on;  
ax = gca;  
ax.XTickLabel = '';  
axis tight;  
title('Original Data');  
  
subplot(212)  
plot(datayear,xrec);  
grid on;
```



```
axis tight;  
xlabel('Year');  
title('El Nino Data - 2-8 Year Periods');
```



Reconstruct Complex Data with Time-varying Trend

Add a trend to the continuous wavelet transform of a complex-valued dataset and reconstruct.

Obtain the CWT of the NPG2006 dataset.

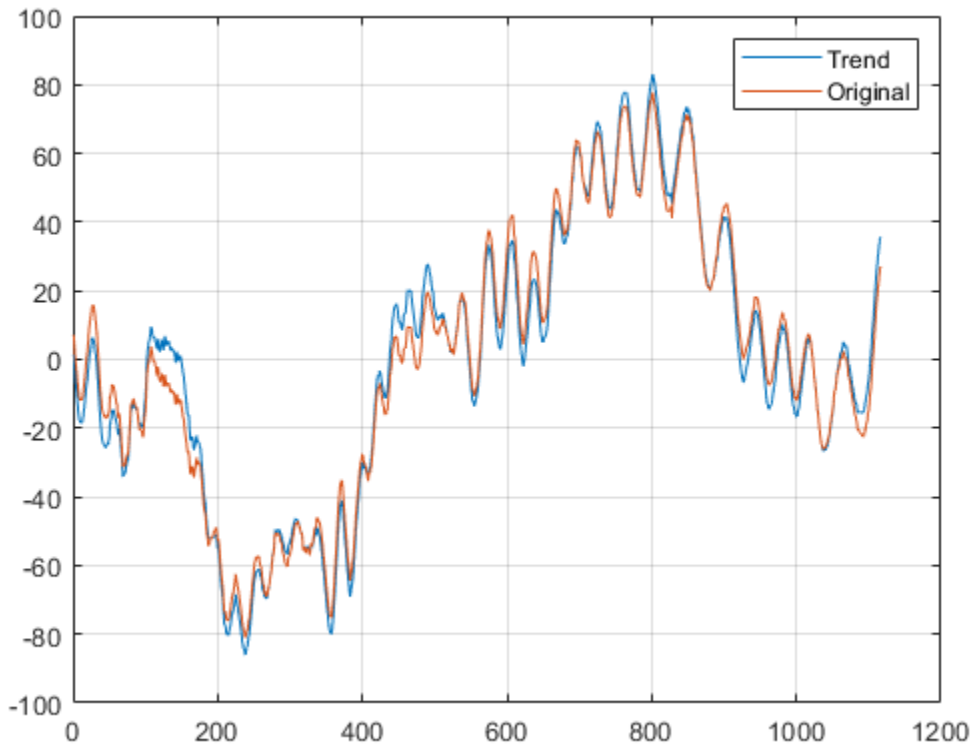
```
load npg2006.mat  
wt = cwt(npg2006.cx);
```

Create a time-varying trend derived from the data.

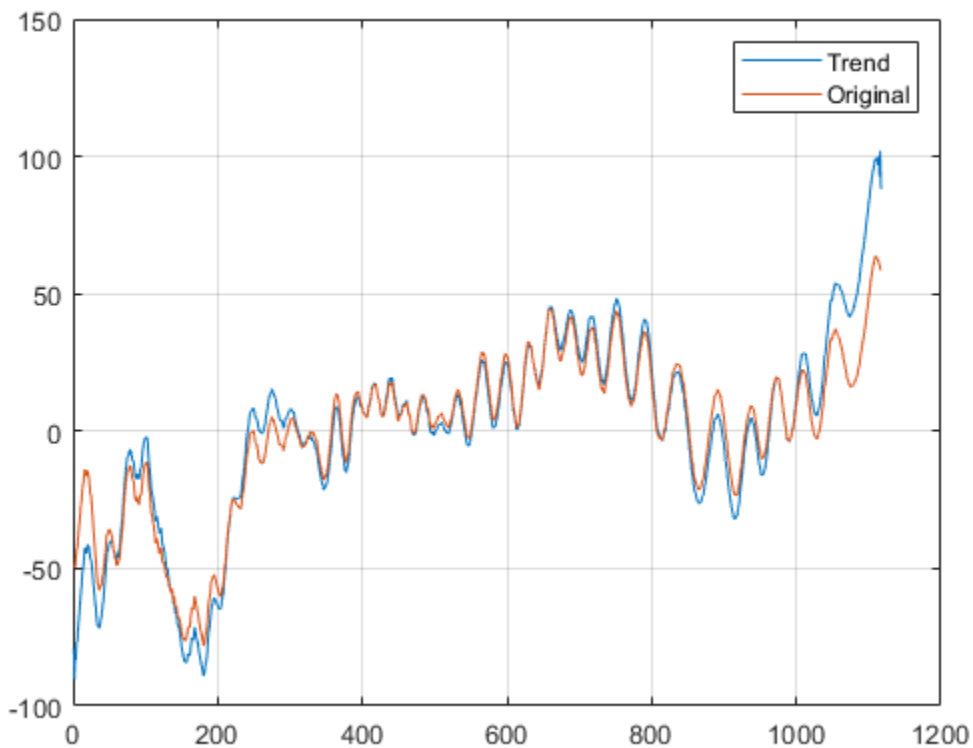
```
trend = smoothdata(npg2006.cx, 'movmean', 100);
```

Obtain the inverse CWT and add the trend. Plot the original data and the reconstructed data.

```
xrec = icwt(wt, 'SignalMean', trend);  
plot([real(xrec)' real(npg2006.cx)])  
grid on  
legend('Trend', 'Original')
```



```
figure
plot([imag(xrec)' imag(npg2006.cx)])
grid on
legend('Trend','Original')
```



Input Arguments

wt — Continuous wavelet transform coefficients

matrix

Continuous wavelet transform coefficients, specified as a matrix of complex values. **wt** is the output from the `cwt` function.

Data Types: double

Complex Number Support: Yes

wname — Analytic wavelet

'morse' (default) | 'amor' | 'bump'

Analytic wavelet used to invert the CWT, specified as 'morse', 'amor', or 'bump'. These character vectors specify the analytic Morse, Morlet, and bump wavelet, respectively. The specified wavelet must be the same type of wavelet used to obtain the cwt.

The default Morse wavelet uses a default symmetry parameter, γ , that is 3 and has a default time bandwidth of 60.

f — CWT frequencies

vector

CWT frequencies, specified as a vector. The number of elements in the frequency vector must equal to the number of rows in the input CWT coefficient matrix, wt. If you specify f, you must also specify freqrange.

freqrange — Frequency range

two-element vector | 2-by-2 matrix

Frequency range for which to return inverse continuous wavelet transform values, specified as a two-element vector or 2-by-2 matrix. If wt is a 2-D matrix, freqrange must be a two-element vector. If wt is a 3-D matrix, freqrange can be a two-element vector or a 2-by-2 matrix. If wt is a 3-D matrix and freqrange is a vector, inversion is performed over the same frequency range in both the positive (analytic) and negative (anti-analytic) components of wt. If freqrange is a 2-by-2 matrix, the first row contains the frequency range for the positive part of wt (first page) and the second row contains the frequency range for the negative part of wt (second page). For a vector, the elements of freqrange must be strictly increasing and contained in the range of the frequency vector f. For a matrix, each row of freqrange must be strictly increasing and contained in the range of f. f is the scale-to-frequency conversion obtained in CWT. For the inversion of a complex-valued signal, you can specify one row of freqrange as a vector of zeros. If the first row of freqrange is a vector of zeros, only the negative (anti-analytic part) is used in the inversion. If you specify freqrange, you must also specify f.

For example [0 0; 1/10 1/4] inverts the negative (clockwise) component over the frequency range [1/10 1/4]. The positive (counterclockwise) component is first set to all zeros before performing the inversion. Similarly, [1/10 1/4; 0 0] inverts the CWT

by selecting the frequency range `[1/10 1/4]` from the positive (counterclockwise) component and setting the negative component to all zeros.

period — Time periods

vector

Time periods corresponding to the rows of CWT coefficient matrix `wt`, specified as a vector. `period` is the output of `cwt`, when the CWT is obtained using a duration input.

periodrange — Period range

two-element vector | 2-by-2 matrix

Period range for which to return inverse continuous wavelet transform values, specified as a two-element vector or 2-by-2 matrix. If `wt` is a 2-D matrix, `periodrange` must be a two-element vector of durations. If `wt` is a 3-D matrix, `periodrange` can be a two-element vector of durations or 2-by-2 matrix of durations. If `periodrange` is a vector of durations and `wt` is a 3-D matrix, inversion is performed over the same frequency range in both the positive (analytic) and negative (anti-analytic) components of `wt`. If `periodrange` is a 2-by-2 matrix of durations, the first row contains the period range for the positive part of `wt` (first page) and the second row contains the period range for the negative part of `wt` (second page). For a vector, the elements of `periodrange` must be strictly increasing and contained in the range of the period vector `period`. The elements of `periodrange` and `period` must have the same units. For a matrix, each row of `periodrange` must be strictly increasing and contained in the range of the period vector `P`. For the inversion of a complex-valued signal, you can specify one row of `periodrange` as a vector of zero durations. If the first row of `periodrange` is a vector of zero durations, only the negative (anti-analytic part) is used in the inversion. If you specify `periodrange`, you must also specify `period`.

For example `[seconds(0) seconds(0); seconds(1/10) seconds(1/4)]` inverts the negative(clockwise) component over the period range `[seconds(1/10) seconds(1/4)]`. The positive (counterclockwise) component is first set to all zeros before performing the inversion. Similarly, `[seconds(1/10) seconds(1/4); seconds(0) seconds(0)]` inverts the CWT by selecting the period range `[1/10 1/4]` from the positive (counterclockwise) component and setting the negative component to all zeros.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes.

You can specify several name and value pair arguments in any order as `Name1,Value1,...,NameN,ValueN`.

Example: `'TimeBandwidth',45` sets the time bandwidth to 45.

TimeBandwidth — Time-bandwidth of Morse wavelet

60 (default) | scalar greater than 3 and less than or equal to 120

Time bandwidth of the Morse wavelet, specified as a comma-separated pair consisting of `'TimeBandwidth'` and a scalar greater than 3 and less than or equal to 120. The specified time bandwidth must be the same as the time bandwidth used in the `cwt`. The symmetry of the Morse wavelet is assumed to be 3.

You cannot specify both the `'TimeBandwidth'` and `'WaveletParameters'`. If you specify `'TimeBandwidth'`, you cannot specify `'WaveletParameters'`. To specify both the symmetry and time bandwidth, use `'WaveletParameters'` instead.

WaveletParameters — Symmetry and time bandwidth of Morse wavelet

(3,60) (default) | two-element vector of scalars

Symmetry and time bandwidth of Morse wavelet, specified as the comma-separated pair consisting of `'WaveletParameters'` and a two-element vector of scalars. The first element of the vector is the symmetry, γ , and the second element is the time-bandwidth. The specified wavelet parameters must be the same as the parameters used in `cwt`.

You cannot specify both `'WaveletParameters'` and `'TimeBandwidth'`. If you specify `'WaveletParameters'`, you cannot specify `'TimeBandwidth'`. To specify the time bandwidth and use the default symmetry value of 3, use `'TimeBandwidth'` instead.

SignalMean — Signal mean

scalar | vector

Signal mean to add to the `icwt` output, specified as the comma-separated pair consisting of `'SignalMean'` and a scalar or vector. If signal mean is a vector, it must be the same length as the column size of the wavelet coefficient matrix. If `wt` is a 2-D matrix, the signal mean must be a real-valued scalar or vector. If `wt` is a 3-D matrix, the signal mean must be a complex-valued scalar or vector. Because `cwt` does not preserve the signal mean, `icwt` does not contain the signal mean by default. Adding a non-zero signal mean to a frequency- or period-limited reconstruction adds a zero-frequency component to the reconstruction.

ScalingCoefficients — Scaling coefficients

real- or complex-valued vector

Scaling coefficients to use in the inverse CWT, specified as a real- or complex-valued vector, obtained as an optional output of `cwt`.

The scaling coefficient output is only supported for Morse wavelets and the analytic Morlet wavelet. The length of `ScalingCoefficients` is equal to the column size of `wt`.

You cannot specify both the `'SignalMean'` and `'ScalingCoefficients'` name-value pairs.

VoicesPerOctave — Number of voices per octave

10 (default) | even integer from 4 to 48

Number of voices per octave, specified as the comma-separated pair consisting of `'VoicesPerOctave'` and an even integer from 4 to 48. The CWT scales are discretized using the specified number of voices per octave. The number of voices per octave must be the same as the number of voices per octave used to obtain the CWT.

You cannot specify the number of voices per octave if you specify either the frequency, `f`, or duration, `period`.

Output Arguments

xrec — Inverse 1-D continuous wavelet transform

real- or complex-valued row vector

Inverse 1-D continuous wavelet transform, returned as a real- or complex-valued row vector.

Data Types: `double`

References

- [1] Lilly, J. M., and S. C. Olhede. "Generalized Morse Wavelets as a Superfamily of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 60, No. 11, 2012, pp. 6036-6041.
- [2] Lilly, J. M., and S. C. Olhede. "Higher-Order Properties of Analytic Wavelets." *IEEE Transactions on Signal Processing*. Vol. 57, No. 1, 2009, pp. 146-160.
- [3] Lilly, J. M. *jLab: A data analysis package for Matlab*, version 1.6.2. 2016. <http://www.jmlilly.net/jmlsoft.html>.

See Also

cwt | cwtfilterbank | cwtfreqbounds | duration | dwt | wavedec | wavefun |
waveinfo | wcodemat | wcoherence | wsst

Topics

“Continuous and Discrete Wavelet Transforms”

“Time-Frequency Analysis with the Continuous Wavelet Transform”

“Morse Wavelets”

Introduced in R2016b

icwtft

Inverse CWT

Note This function is no longer recommended. Use `icwt` instead.

Syntax

```
xrec = icwtft(cwtstruct)
xrec = icwtft(cwtstruct,'plot')
xrec = icwtft(cwtstruct,'signal',SIG,'plot')
```

Description

`xrec = icwtft(cwtstruct)` returns the inverse continuous wavelet transform of the CWT coefficients contained in the `cfs` field of the structure array `cwtstruct`. Obtain the structure array `cwtstruct` as the output of `cwtft`.

`xrec = icwtft(cwtstruct,'plot')` plots the reconstructed signal.

`xrec = icwtft(cwtstruct,'signal',SIG,'plot')` places a radio button in the bottom left corner of the plot. Enabling the radio button superimposes the plot of the input signal `SIG` on the plot of the reconstructed signal. By default the radio button is not enabled and only the reconstructed signal is plotted.

Input Arguments

`cwtstruct`

Structure array containing six fields.

- `cfs` — CWT coefficient matrix
- `scales` — Vector of scales
- `frequencies` — frequencies in cycles per unit time (or space) corresponding to the scales. If the sampling period units are seconds, the frequencies are in hertz. The

elements of frequencies are in decreasing order to correspond to the elements in the scales vector.

- `omega` — Angular frequencies used in the Fourier transform
- `meanSig` — Mean of the analyzed signal
- `dt` — The sampling period
- `wav` — Analyzing wavelet used in the CWT with parameters specified

`cwtstruct` is the output of `cwtft`.

Output Arguments

`xrec`

Reconstructed signal

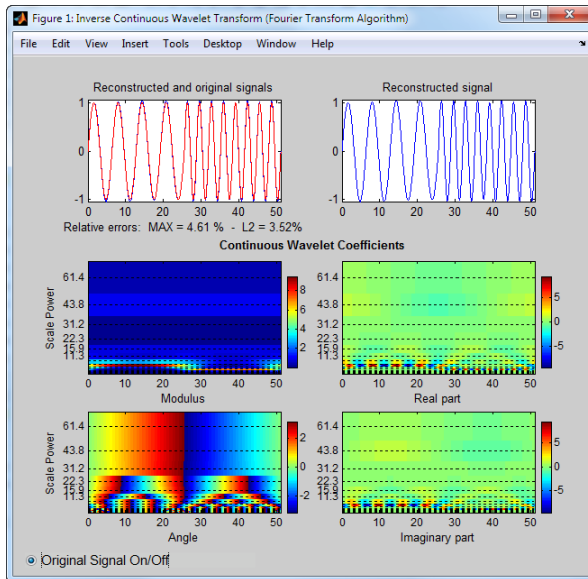
Examples

Compute the CWT and inverse CWT of two sinusoids with disjoint support.

```
N = 1024;
t = linspace(0,1,N);
y = sin(2*pi*8*t).*(t<=0.5)+sin(2*pi*16*t).*(t>0.5);
dt = 0.05;
s0 = 2*dt;
ds = 0.4875;
NbSc = 20;
wname = 'morl';
sig = {y,dt};
sca = {s0,ds,NbSc};
wave = {wname,[]};
cwtsig = cwtft(sig,'scales',sca,'wavelet',wave);

% Compute inverse CWT and plot reconstructed signal with original
sigrec = icwtft(cwtsig,'signal',sig,'plot');
```

Select the radio button in the bottom left corner of the plot.



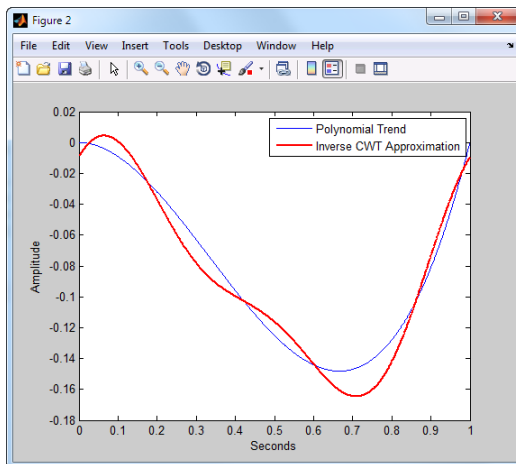
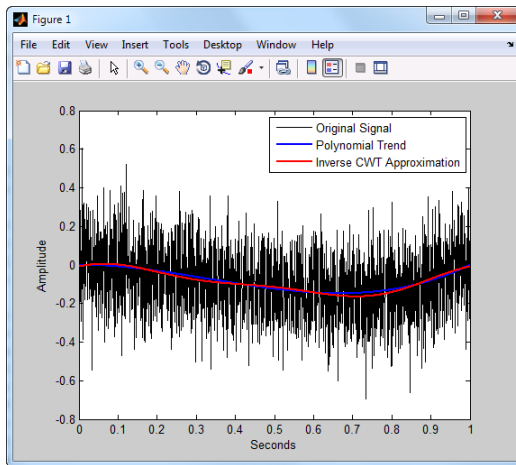
Use the inverse CWT to approximate a trend in a time series. Construct a time series consisting of a polynomial trend, a sinewave (oscillatory component), and additive white Gaussian noise. Obtain the CWT of the input signal and use the inverse CWT based on only the coarsest scales to reconstruct an approximation to the trend. To obtain an accurate approximation based on select scales use the default power of two spacing for the scales in the continuous wavelet transform. See `cwtft` for details.

```
t = linspace(0,1,1e3);
% Polynomial trend
x = t.^3-t.^2;
% Periodic term
x1 = 0.25*cos(2*pi*250*t);
% Reset random number generator for reproducible results
rng default
y = x+x1+0.1*randn(size(t));
% Obtain CWT of input time series
cwt = cwtft({y,0.001},'wavelet','morl');
% Zero out all but the coarsest scale CWT coefficients
cwt.cfs(1:16,:) = 0;
% Reconstruct a signal approximation based on the coarsest scales
xrec = icwtft(cwt);
plot(t,y,'k'); hold on;
xlabel('Seconds'); ylabel('Amplitude');
```

```

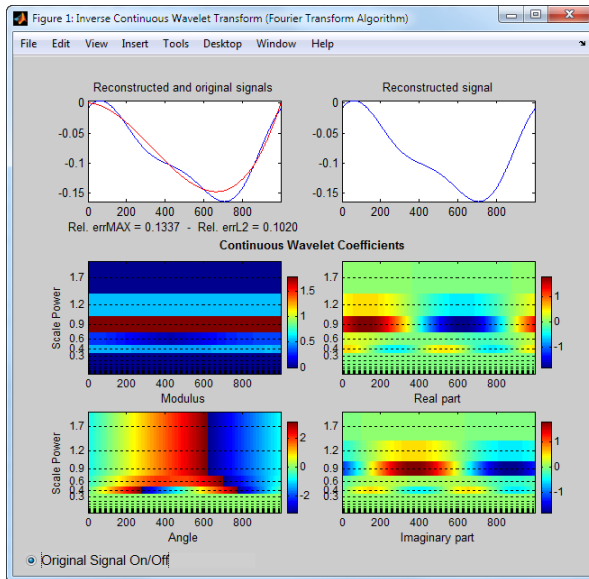
plot(t,x,'b','linewidth',2);
plot(t,xrec,'r','linewidth',2);
legend('Original Signal','Polynomial Trend',...
      'Inverse CWT Approximation');
figure
plot(t,x,'b'); hold on;
xlabel('Seconds'); ylabel('Amplitude');
plot(t,xrec,'r','linewidth',2);
legend('Polynomial Trend','Inverse CWT Approximation');

```



You can also use the following syntax to plot the approximation. Select the radio button to view the original polynomial trend superimposed on the wavelet approximation.

```
% Input the polynomial trend as the value of 'signal'
xrec = icwtft(cwty, 'signal', x, 'plot');
```



Definitions

Inverse CWT

`icwtft` computes the inverse CWT based on a discretized version of the single integral formula due to Morlet. The Wavelet Toolbox Getting Started Guide contains a brief description of the theoretical foundation for the single integral formula in “Inverse Continuous Wavelet Transform”. The discretized version of this integral is presented in [5]

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, Philadelphia, PA: Society for Industrial and Applied Mathematics (SIAM), 1992.
- [2] Farge, M. "Wavelet Transforms and Their Application to Turbulence", *Ann. Rev. Fluid. Mech.*, 1992, 24, 395-457.
- [3] Mallat, S. *A Wavelet Tour of Signal Processing*, San Diego, CA: Academic Press, 1998.
- [4] Sun, W. "Convergence of Morlet's Reconstruction Formula", *preprint*, 2010.
- [5] Torrence, C. and G.P. Compo "A Practical Guide to Wavelet Analysis", *Bull. Am. Meteorol. Soc.*, 79, 61-78, 1998.

See Also

cwt | cwtft | icwtlin

Topics

"Continuous and Discrete Wavelet Transforms"
"Continuous Wavelet Transform and Scale-Based Analysis"
"Inverse Continuous Wavelet Transform"

Introduced in R2011a

icwtlin

Inverse continuous wavelet transform (CWT) for linearly spaced scales

Note This function is no longer recommended. Use `icwt` instead.

Syntax

```
xrec = icwtlin(cwtstruct)
xrec = icwtlin(wav,meanSIG,cfs,scales,dt)
xrec = icwtin(...,'plot')
xrec = icwtlin(...,'signal',SIG,'plot')
xrec = icwtlin(...,Name,Value)
```

Description

`xrec = icwtlin(cwtstruct)` returns the inverse continuous wavelet transform (CWT) of the CWT coefficients obtained at linearly spaced scales.

Note To use `icwtlin` you must:

- Use linearly-spaced scales in the CWT. `icwtlin` does not verify that the scales are linearly-spaced.
 - Use one of the supported wavelets. See “Input Arguments” on page 1-554 for a list of supported wavelets.
-

`xrec = icwtlin(wav,meanSIG,cfs,scales,dt)` returns the inverse CWT of the coefficients in `cfs`. The inverse CWT is obtained using the wavelet `wav`, the linearly spaced scales `scales`, the sampling period `dt`, and the mean signal value `meanSig`.

`xrec = icwtin(...,'plot')` plots the reconstructed signal `xrec` along with the CWT coefficients and CWT moduli. If the analyzing wavelet is complex-valued, the plot includes the real and imaginary parts of the CWT coefficients.

`xrec = icwtlin(..., 'signal', SIG, 'plot')` places a radio button in the bottom-left corner of the plot. Enabling the radio button superimposes the plot of the input signal `SIG` on the plot of the reconstructed signal. `SIG` can be a structure array, a cell array, or a vector. If `SIG` is a structure array, there must be two fields: `val` and `period`. The `val` field contains the signal and the `period` field contains the sampling period. If `SIG` is a cell array, `SIG{1}` contains the signal and `SIG{2}` is the sampling period.

`xrec = icwtlin(..., Name, Value)` returns the inverse CWT transform with additional options specified by one or more `Name, Value` pair arguments.

Input Arguments

cwtstruct

A structure array that is the output of `cwtft` or constructed from the output of `cwt`. If you obtain `cwtstruct` from `cwtft`, the structure array contains seven fields:

- `cfs` — CWT coefficient matrix
- `scales` — Vector of linearly spaced scales. The scale vector must be linearly-spaced to ensure accurate reconstruction. `icwtlin` does not check that the spacing of your scale vector is linear.
- `frequencies` — frequencies in cycles per unit time (or space) corresponding to the scales. If the sampling period units are seconds, the frequencies are in hertz. The elements of `frequencies` are in decreasing order to correspond to the elements in the `scales` vector.
- `omega` — Angular frequencies used in the Fourier transform in radians/sample
- `MeanSIG` — Signal mean
- `dt` — Sampling period in seconds
- `wav` — Analyzing wavelet. `icwtlin` uses this wavelet as the reconstruction wavelet. The supported wavelets are:
 - `'dog'` — An m -th order derivative of Gaussian wavelet where m is a positive even integer. $m = 2$ is the Mexican-hat or Ricker wavelet.
 - `'morl'` — Analytic Morlet wavelet
 - `'morlex'` — Nonanalytic Morlet wavelet
 - `'morl0'` — Nonanalytic Morlet wavelet with exact zero mean

- 'mexh' — Mexican-hat wavelet. This argument represents a special case of the derivative of Gaussian wavelet with $m = 2$. This wavelet is also known as the Ricker wavelet.
- 'paul' — Paul wavelet
- 'bump' — Bump wavelet

If you create `cwtstruct` from the output of `cwt`, `cwtstruct` contains all of the preceding fields except `omega`.

Using `cwt` to obtain the CWT coefficients, the valid analyzing wavelets are:

- Coiflets — 'coif1', 'coif2', 'coif3', 'coif4', 'coif5'
- Biorthogonal wavelets — 'bior2.2', 'bior2.4', 'bior2.6', 'bior2.8', 'bior4.4', 'bior5.5', 'bior6.8'
- Reverse biorthogonal wavelets — 'rbio2.2', 'rbio2.4', 'rbio2.6', 'rbio2.8', 'rbio4.4', 'rbio5.5', 'rbio6.8'
- Complex Gaussian wavelets — 'cgau2', 'cgau4', 'cgau6', 'cgau8'

Name-Value Pair Arguments

IdxSc

Vector of scales to use in the signal reconstruction. Specifying a subset of scales results in a scale-localized approximation of the analyzed signal.

Output Arguments

xrec

Reconstructed signal. Signal approximation based on the input CWT coefficient matrix, analyzing wavelet, selected scales, and sampling period.

The purpose of the CWT inversion algorithm is not to produce a perfect reconstruction of the input signal. The inversion preserves time and scale-localized features in the reconstructed signal. The amplitude scaling in the reconstructed signal, however, can be significantly different. This difference in scaling can occur whether or not you use all the CWT coefficients in the inversion.

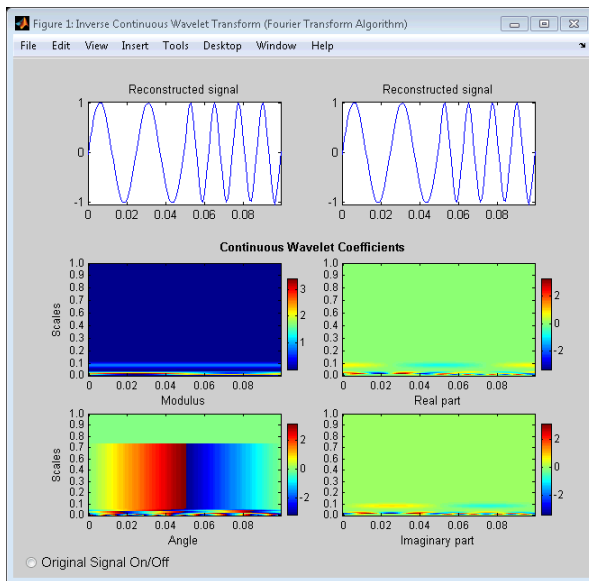
Examples

Compute the inverse CWT of a sum of sine waves with disjoint support.

```
% Define the signal
N = 100;
t = linspace(0,1,N);
Y = sin(8*pi*t).*(t<=0.5) + sin(16*pi*t).*(t>0.5) ;

% Define parameters before analysis
dt = 0.001;
maxsca = 1; s0 = 2*dt; ds = 2*dt;
scales = s0:ds:maxsca;
wname = 'morl';
SIG = {Y,dt};
WAV = {wname,[]};

% Compute the CWT using cwtft with linear scales
cwtS = cwtft(SIG,'scales',scales,'wavelet',WAV);
% Compute inverse CWT using linear scales
Yrec = icwtlin(cwtS,'Signal',Y,'plot');
```



Reconstruct an approximation to a noisy Doppler signal based on thresholded coefficients. Use the universal threshold. Assume the sampling period is 0.05 seconds.

```
load noisdopp;
Y = noisdopp;
N = length(Y);

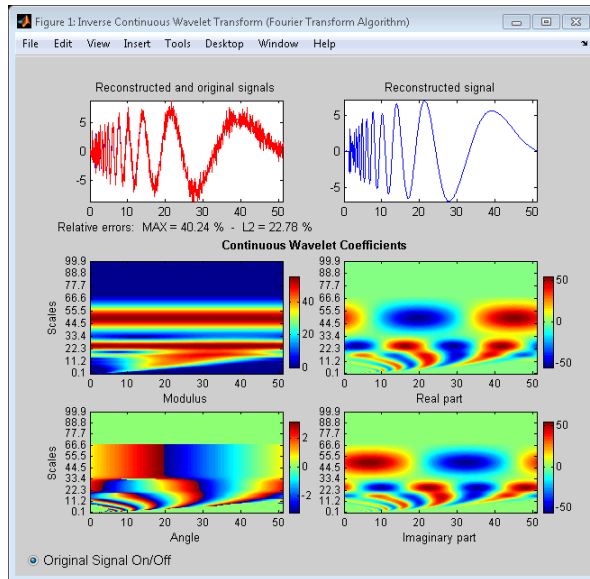
% Define parameters before analysis
% Assume sampling period is 0.05
dt = 0.05;
maxsca = 100; s0 = 2*dt; ds = 4*dt;
scales = s0:ds:maxsca;
wname = 'morl';
SIG = {Y,dt};
WAV = {wname,[]};

% Compute CWT
cwtS = cwtft(SIG,'scales',scales,'wavelet',WAV,'plot');

% Select subset of coefficients
cwtS1 = cwtS;
Hfreq = cwtS.cfs(1:10,:);
% Set threshold
thr = sqrt(2*log(N))*median(abs(Hfreq(:)))/0.6745;
newCFS = cwtS.cfs;
% Set coefficients smaller than threshold in absolute value to 0
newCFS(abs(newCFS)<thr) = 0;
cwtS1.cfs = newCFS;

% Reconstruction from the modified structure
YRDen = icwtlin(cwtS1,'signal',Y,'plot');
```

Enable the **Reconstructed Signal On/Off** radio button in the bottom-left corner.



Algorithms

See [4] for a description of the inverse CWT algorithm for linearly spaced scales. The `icwtlin` function uses heuristic scaling factors for the analyzing wavelets. These scaling factors can result in significant differences in the amplitude scaling of the reconstructed signal.

Alternatives

- `icwtft` — Computes the inverse for the CWT obtained using `cwtft` with logarithmically spaced scales. If you use linearly spaced scales in `cwtft`, or you obtain the CWT with `cwt`, use `icwtlin` to compute the inverse.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, Philadelphia, PA: Society for Industrial and Applied Mathematics (SIAM), 1992.

- [2] Farge, M. "Wavelet Transforms and Their Application to Turbulence", *Ann. Rev. Fluid. Mech.*, 1992, 24, 395-457.
- [3] Mallat, S. *A Wavelet Tour of Signal Processing*, San Diego, CA: Academic Press, 1998.
- [4] Sun, W. "Convergence of Morlet's Reconstruction Formula", *preprint*, 2010.
- [5] Torrence, C. and G.P. Compo. "A Practical Guide to Wavelet Analysis", *Bull. Am. Meteorol. Soc.*, 79, 61-78, 1998.

See Also

cwt | cwtft | icwtft

Topics

"Continuous and Discrete Wavelet Transforms"

"Continuous Wavelet Transform and Scale-Based Analysis"

"Inverse Continuous Wavelet Transform"

Introduced in R2011b

iddtree

Inverse dual-tree and double-density 1-D wavelet transform

Syntax

```
xrec = iddtree(wt)
```

Description

`xrec = iddtree(wt)` returns the inverse wavelet transform of the wavelet decomposition (analysis filter bank), `wt`. `wt` is the output of `dddtree`.

Examples

Perfect Reconstruction Using Dual-Tree Double-Density Wavelet Filter Bank

Demonstrate perfect reconstruction of a signal using a dual-tree double-density wavelet transform.

Load the noisy Doppler signal. Obtain the dual-tree double-density wavelet transform down to level 5. Invert the transform and demonstrate perfect reconstruction.

```
load noisdopp;
wt = dddtree('cplxddd',noisdopp,5,'FSdoubledualfilt',...
    'doubledualfilt');
xrec = iddtree(wt);
max(abs(noisdopp-xrec))

ans = 1.9291e-12
```

Input Arguments

wt — Wavelet transform

structure

Wavelet transform, returned as a structure from `dddtree` with these fields:

type — Type of wavelet decomposition (filter bank)

'dwt' | 'ddt' | 'cplxdt' | 'cplxdddtt'

Type of wavelet decomposition (filter bank), specified as one of 'dwt', 'ddt', 'cplxdt', or 'cplxdddtt'. The type, 'dwt', gives a critically sampled discrete wavelet transform. The other types are oversampled wavelet transforms. 'ddt' is a double-density wavelet transform, 'cplxdt' is a dual-tree complex wavelet transform, and 'cplxdddtt' is a double-density dual-tree complex wavelet transform.

level — Level of wavelet decomposition

positive integer

Level of wavelet decomposition, specified as a positive integer.

filters — Decomposition (analysis) and reconstruction (synthesis) filters

structure

Decomposition (analysis) and reconstruction (synthesis) filters, specified as a structure with these fields:

Fdf — First-stage analysis filters

matrix | cell array

First-stage analysis filters, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

Df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1 , specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the analysis filters for the corresponding tree.

FrF — First-level reconstruction filters

matrix | cell array

First-level reconstruction filters, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

Rf — Reconstruction filters for levels > 1

matrix | cell array

Reconstruction filters for levels > 1 , specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the synthesis filters for the corresponding tree.

cfs — Wavelet transform coefficients

cell array of matrices

Wavelet transform coefficients, specified as a 1-by-(level+1) cell array of matrices. The size and structure of the matrix elements of the cell array depend on the type of wavelet transform as follows:

- 'dwt' — cfs{j}

- $j = 1, 2, \dots$ level is the level.
- $cfs\{\text{level}+1\}$ are the lowpass, or scaling, coefficients.
- 'ddt' — $cfs\{j\}(:, :, k)$
 - $j = 1, 2, \dots$ level is the level.
 - $k = 1, 2$ is the wavelet filter.
 - $cfs\{\text{level}+1\}(:, :,)$ are the lowpass, or scaling, coefficients.
- 'cplxdt' — $cfs\{j\}(:, :, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :,)$ are the lowpass, or scaling, coefficients.
- 'cplxddd' — $cfs\{j\}(:, :, k, m)$
 - $j = 1, 2$ level is the level.
 - $k = 1, 2$ is the wavelet filter.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :,)$ are the lowpass, or scaling, coefficients.

Output Arguments

xrec — Synthesized 1-D signal

vector

Synthesized 1-D signal, returned as a vector. The row or column orientation of `xrec` depends on the row or column orientation of the 1-D signal input to `dddtree`.

Data Types: double

See Also

`dddtree` | `dddtreecfs` | `plotted`

Topics

“Analytic Wavelets Using the Dual-Tree Wavelet Transform”

“Critically Sampled and Oversampled Wavelet Filter Banks”

Introduced in R2013b

idddtree2

Inverse dual-tree and double-density 2-D wavelet transform

Syntax

```
xrec = idddtree2(wt)
```

Description

`xrec = idddtree2(wt)` returns the inverse wavelet transform of the 2-D decomposition (analysis filter bank), `wt`. `wt` is the output of `dddtree2`.

Examples

Perfect Reconstruction Using Complex Oriented Dual-Tree Wavelet Filter Bank

Demonstrate perfect reconstruction of an image using a complex oriented dual-tree wavelet transform.

Load the image and obtain the complex oriented dual-tree wavelet transform down to level 5 using `dddtree2`. Reconstruct the image using `idddtree2` and demonstrate perfect reconstruction.

```
load woman;  
wt = dddtree2('cplxdt',X,5,'dtf2');  
xrec = idddtree2(wt);  
max(max(abs(X-xrec)))  
  
ans = 7.3612e-12
```

Input Arguments

wt — Wavelet transform

structure

Wavelet transform, returned as a structure from `ddtree2` with these fields:

type — Type of wavelet decomposition (filter bank)

'dwt' | 'ddt' | 'realdt' | 'cplxdt' | 'realdtdt' | 'cplxdttdt'

Type of wavelet decomposition (filter bank), specified as one of 'dwt', 'ddt', 'realdt', 'cplxdt', 'realdtdt', or 'cplxdttdt'. 'dwt' is the critically sampled DWT. 'ddt' produces a double-density wavelet transform with one scaling and two wavelet filters for both row and column filtering. 'realdt' and 'cplxdt' produce oriented dual-tree wavelet transforms consisting of two and four separable wavelet transforms. 'realdtdt' and 'cplxdttdt' produce double-density dual-tree wavelet transforms consisting of two and four separable wavelet transforms.

level — Level of the wavelet decomposition

positive integer

Level of the wavelet decomposition, specified as a positive integer.

filters — Decomposition (analysis) and reconstruction (synthesis) filters

structure

Decomposition (analysis) and reconstruction (synthesis) filters, specified as a structure with these fields:

Fdf — First-stage analysis filters

matrix | cell array

First-stage analysis filters, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

Df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the analysis filters for the corresponding tree.

FrF — First-level reconstruction filters

matrix | cell array

First-level reconstruction filters, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

Rf — Reconstruction filters for levels > 1

matrix | cell array

Reconstruction filters for levels > 1, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

cfs — Wavelet transform coefficients

cell array of matrices

Wavelet transform coefficients, specified as a 1-by-(level+1) cell array of matrices. The size and structure of the matrix elements of the cell array depend on the type of wavelet transform as follows:

- 'dwt' — $cfs\{j\}(:, :, d)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $cfs\{level+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'ddt' — $cfs\{j\}(:, :, d)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3, 4, 5, 6, 7, 8$ is the orientation.
 - $cfs\{level+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'realddt' — $cfs\{j\}(:, :, d, k)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $cfs\{level+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $m = 1, 2$ are the real and imaginary parts.
 - $cfs\{level+1\}(:, :, :)$ are the lowpass, or scaling, coefficients..
- 'realdddt' — $cfs\{j\}(:, :, d, k)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.
 - $cfs\{level+1\}(:, :, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdddt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1, 2, \dots$ level is the level.
 - $d = 1, 2, 3$ is the orientation.
 - $k = 1, 2$ is the wavelet transform tree.

- $m = 1, 2$ are the real and imaginary parts.
- `cfs{level+1}(:, :)` are the lowpass, or scaling, coefficients.

Output Arguments

xrec — Synthesized 2-D image

matrix

Synthesized image, returned as a matrix.

Data Types: `double`

See Also

`dddtree2` | `dddtreecfs`

Topics

“Analytic Wavelets Using the Dual-Tree Wavelet Transform”

“Critically Sampled and Oversampled Wavelet Filter Banks”

Introduced in R2013b

idualtree3

3-D dual-tree complex wavelet reconstruction

Syntax

```
xrec = idualtree3(a,d)  
xrec = idualtree3(a,d,Name,Value)
```

Description

`xrec = idualtree3(a,d)` returns the inverse 3-D dual-tree complex wavelet transform of the final-level approximation coefficients, `a`, and cell array of wavelet coefficients, `d`.

`xrec = idualtree3(a,d,Name,Value)` specifies options using name-value pair arguments.

Examples

Wavelet Coefficients

Generate all-zero sets of scaling and wavelet coefficients by computing the 3-D dual-tree complex wavelet transform of an array of zeros.

```
zr = zeros(64,64,64);
```

```
[a,d] = dualtree3(zr,4);
```

Find the real (4,5) wavelet coefficient of the 19th subband of the third level by assigning 1 to the corresponding array element and inverting the transform.

```
d{3}(4,5,19) = 1;
```

```
xr = idualtree3(a,d);
```


Find the corresponding imaginary coefficient assigning the imaginary unit to the array element and then inverting the transform.

```
[a,d] = dualtree3(zr,4);
```

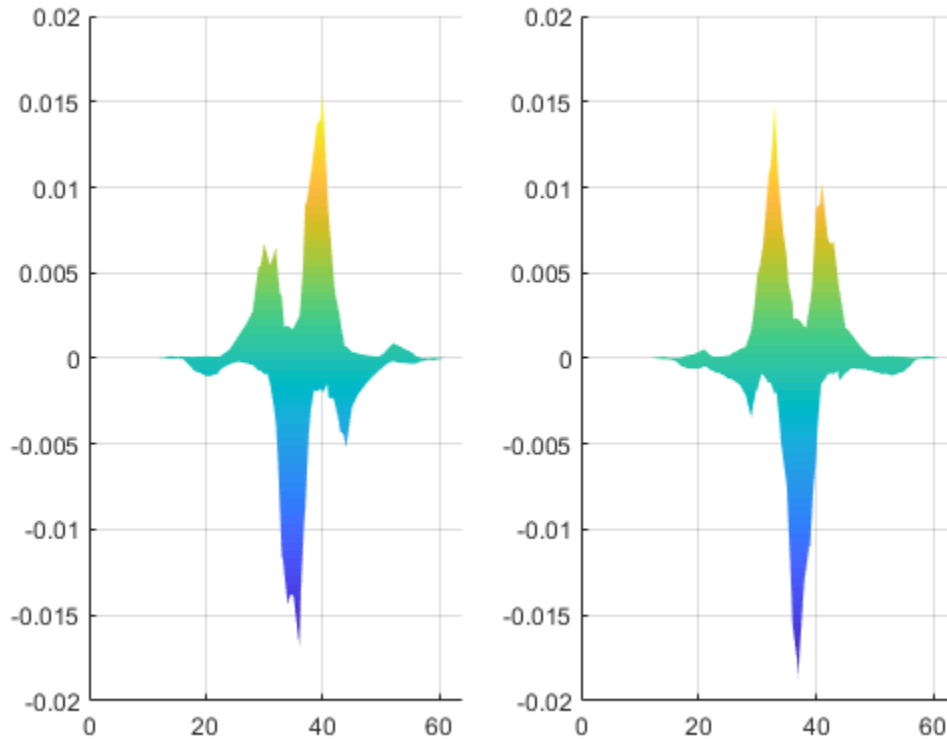
```
d{3}(4,5,19) = 1j;
```

```
xi = idualtree3(a,d);
```

Display the 18th page of the real and imaginary reconstructions.

```
subplot(1,2,1)  
surf(xr(:,:,18))  
view(0,0)  
zlim([-0.02 0.02])  
shading interp
```

```
subplot(1,2,2)  
surf(xi(:,:,18))  
view(0,0)  
zlim([-0.02 0.02])  
shading interp
```



Input Arguments

a — Final-level scaling coefficients

real-valued matrix

Final-level scaling coefficients, specified as a real-valued matrix. **a** is an output of `dualtree3`.

Data Types: `single` | `double`

d — Wavelet coefficients

cell array

Wavelet coefficients, specified as a cell array. `d` is an output of `dualtree3`.

Data Types: `single` | `double`

Complex Number Support: Yes

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'LevelOneFilter', 'legall', 'FilterLength', 6` inverts a transform using LeGall synthesis filters with scaling length 3 and wavelet length 5 at level 1, and length-6 Q-shift filters at levels 2 and greater.

FilterLength — Hilbert Q-shift filter-pair length

10 (default) | 6 | 14 | 16 | 18

Hilbert Q-shift filter-pair length, specified as the comma-separated pair consisting of `'FilterLength'` and one of 6, 10, 14, 16, or 18. The synthesis filters used by `idualtree3` must match the analysis filters used by `dualtree3`.

Data Types: `double` | `single`

LevelOneFilter — First-level biorthogonal analysis filter

`'nearsym5_7'` (default) | `'nearsym13_19'` | `'antonini'` | `'legall'`

First-level biorthogonal analysis filter, specified as the comma-separated pair consisting of `'LevelOneFilter'` and a character vector or string. By default, `idualtree3` uses the near-symmetric biorthogonal wavelet filter with lengths 7 (scaling synthesis filter) and 5 (wavelet synthesis filter) in the reconstruction.

Data Types: `char` | `string`

OriginalDataSize — Size of the original data

three-element vector of even integers

Size of the original data, specified as the comma-separated pair consisting of `'OriginalDataSize'` and a three-element vector of even integers. This vector must match the size of the original input to the 3-D dual-tree wavelet transform. When the first-level wavelet coefficients are not available, the reconstructed data size can differ from the original input data size. If you call `dualtree3` with the `'excludeL1'` option, then

'OriginalDataSize' adjusts the size of `xrec` to match the size of the original input data. If you do not use the 'excludeL1' option, then this argument is ignored.

Data Types: `double` | `single`

Output Arguments

`xrec` — Inverse 3-D dual-tree complex wavelet transform

3-D array

Inverse 3-D dual-tree complex wavelet transform, returned as a 3-D array.

References

- [1] Chen, H., and N. G. Kingsbury. "Efficient Registration of Nonrigid 3-D Bodies." *IEEE Transactions on Image Processing*. Vol 21, January 2012, pp. 262-272.
- [2] Kingsbury, N. G. "Complex Wavelets for Shift Invariant Analysis and Filtering of Signals." *Journal of Applied and Computational Harmonic Analysis*. Vol. 10, May 2001, pp. 234-253.

See Also

`dddtree2` | `dualtree3` | `wavedec3` | `waverec3`

Topics

Dual-Tree Wavelet Transforms

Introduced in R2017a

idwt

Single-level inverse discrete 1-D wavelet transform

Syntax

```
X = idwt(cA,cD,'wname')
X = idwt(cA,cD,Lo_R,Hi_R)
X = idwt(cA,cD,'wname',L)
X = idwt(cA,cD,Lo_R,Hi_R,L)
idwt(cA,cD,'wname')
X = idwt(...,'mode',MODE)
X = idwt(cA,[],...)
X = idwt([],cD,...)
```

Description

The `idwt` command performs a single-level one-dimensional wavelet reconstruction with respect to either a particular wavelet (`'wname'`, see `wfilters` for more information) or particular wavelet reconstruction filters (`Lo_R` and `Hi_R`) that you specify.

`X = idwt(cA,cD,'wname')` returns the single-level reconstructed approximation coefficients vector `X` based on approximation and detail coefficients vectors `cA` and `cD`, and using the wavelet `'wname'`.

`X = idwt(cA,cD,Lo_R,Hi_R)` reconstructs as above using filters that you specify.

- `Lo_R` is the reconstruction low-pass filter.
- `Hi_R` is the reconstruction high-pass filter.

`Lo_R` and `Hi_R` must be the same length.

Let `la` be the length of `cA` (which also equals the length of `cD`) and `lf` the length of the filters `Lo_R` and `Hi_R`; then $\text{length}(X) = LX$ where $LX = 2*la$ if the DWT extension mode is set to periodization. For the other extension modes $LX = 2*la - lf + 2$.

For more information about the different Discrete Wavelet Transform extension modes, see `dwtmode`.

`X = idwt(cA,cD,'wname',L)` or `X = idwt(cA,cD,Lo_R,Hi_R,L)` returns the length-`L` central portion of the result obtained using `idwt(cA,cD,'wname')`. `L` must be less than `LX`.

`X = idwt(...,'mode',MODE)` computes the wavelet reconstruction using the specified extension mode `MODE`.

`X = idwt(cA,[],...)` returns the single-level reconstructed approximation coefficients vector `X` based on approximation coefficients vector `cA`.

`X = idwt([],cD,...)` returns the single-level reconstructed detail coefficients vector `X` based on detail coefficients vector `cD`.

Examples

Inverse DWT Using Orthogonal Wavelet

Demonstrate perfect reconstruction using `dwt` and `idwt` with an orthonormal wavelet.

```
load noisdopp;
[A,D] = dwt(noisdopp,'sym4');
x = idwt(A,D,'sym4');
max(abs(noisdopp-x))

ans = 3.2156e-12
```

Inverse DWT Using Biorthogonal Wavelet

Demonstrate perfect reconstruction using `dwt` and `idwt` with a biorthogonal wavelet.

```
load noisdopp;
[Lo_D,Hi_D,Lo_R,Hi_R] = wfilters('bior3.5');
[A,D] = dwt(noisdopp,Lo_D,Hi_D);
x = idwt(A,D,Lo_R,Hi_R);
max(abs(noisdopp-x))
```

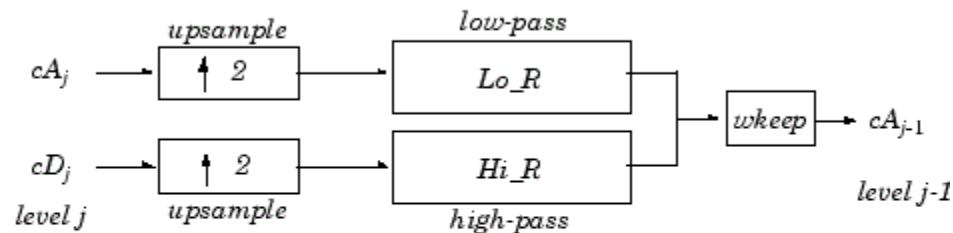
ans = 3.5527e-15

Algorithms

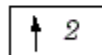
Starting from the approximation and detail coefficients at level j , cA_j and cD_j , the inverse discrete wavelet transform reconstructs cA_{j-1} , inverting the decomposition step by inserting zeros and convolving the results with the reconstruction filters.

One-Dimensional IDWT

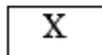
Reconstruction step



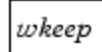
Where



Insert zeros at even-indexed elements



Convolve with filter X



Take the central part of U with the convenient length

References

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed.

Mallat, S. (1989), "A theory for multiresolution signal decomposition: the wavelet representation," *IEEE Pattern Anal. and Machine Intell.*, vol. 11, no. 7, pp. 674-693.

Meyer, Y. (1990), *Ondelettes et opérateurs*, Tome 1, Hermann Ed. (English translation: *Wavelets and operators*, Cambridge Univ. Press. 1993.)

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`dwt` | `dwtmode` | `upwlev`

Introduced before R2006a

idwt2

Single-level inverse discrete 2-D wavelet transform

Syntax

```
X = idwt2(cA,cH,cV,cD,'wname')
X = idwt2(cA,cH,cV,cD,Lo_R,Hi_R)
X = idwt2(cA,cH,cV,cD,'wname',S)
X = idwt2(cA,cH,cV,cD,Lo_R,Hi_R,S)
idwt2(cA,cH,cV,cD,'wname')
X = idwt2(...,'mode',MODE)
X = idwt2(cA,[],[],[],...)
X = idwt2([],cH,[],[],...)
```

Description

The `idwt2` command performs a single-level two-dimensional wavelet reconstruction with respect to either a particular wavelet (`'wname'`, see `wfilters` for more information) or particular wavelet reconstruction filters (`Lo_R` and `Hi_R`) that you specify.

`X = idwt2(cA,cH,cV,cD,'wname')` uses the wavelet `'wname'` to compute the single-level reconstructed approximation coefficients matrix `X`, based on approximation matrix `cA` and details matrices `cH`, `cV`, and `cD` (horizontal, vertical, and diagonal, respectively).

`X = idwt2(cA,cH,cV,cD,Lo_R,Hi_R)` reconstructs as above, using filters that you specify.

- `Lo_R` is the reconstruction low-pass filter.
- `Hi_R` is the reconstruction high-pass filter.

`Lo_R` and `Hi_R` must be the same length.

Let `sa = size(cA) = size(cH) = size(cV) = size(cD)` and `lf` the length of the filters; then `size(X) = SX`, where `SX = 2* SA`, if the DWT extension mode is set to periodization. For the other extension modes, `SX = 2*size(cA) - lf + 2`.

For more information about the different Discrete Wavelet Transform extension modes, see `dwtmode`.

`X = idwt2(cA,cH,cV,cD,'wname',S)` and `X = idwt2(cA,cH,cV,cD,Lo_R,Hi_R,S)` return the size-`S` central portion of the result obtained using the syntax `idwt2(cA,cH,cV,cD,'wname')`. `S` must be less than `SX`.

`X = idwt2(...,'mode',MODE)` computes the wavelet reconstruction using the extension mode `MODE` that you specify.

`X = idwt2(cA,[],[],[],...)` returns the single-level reconstructed approximation coefficients matrix `X` based on approximation coefficients matrix `cA`.

`X = idwt2([],cH,[],[],...)` returns the single-level reconstructed detail coefficients matrix `X` based on horizontal detail coefficients matrix `cH`.

The same result holds for `X = idwt2([],[],cV,[],...)` and `X = idwt2([],[],[],cD,...)`, based on vertical and diagonal details.

More generally, `X = idwt2(AA,HH,VV,DD,...)` returns the single-level reconstructed matrix `X`, where `AA` can be `cA` or `[]`, and so on.

`idwt2` is the inverse function of `dwt2` in the sense that the abstract statement `idwt2(dwt2(X,'wname'),'wname')` would give back `X`.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load original image.  
load woman;
```

```
% X contains the loaded image.  
sX = size(X);
```

```
% Perform single-level decomposition  
% of X using db4.  
[cA1,cH1,cV1,cD1] = dwt2(X,'db4');
```

```
% Invert directly decomposition of X  
% using coefficients at level 1.  
A0 = idwt2(cA1,cH1,cV1,cD1,'db4',sX);
```

```
% Check for perfect reconstruction.
max(max(abs(X-A0)))
ans =
    3.4176e-10
```

Tips

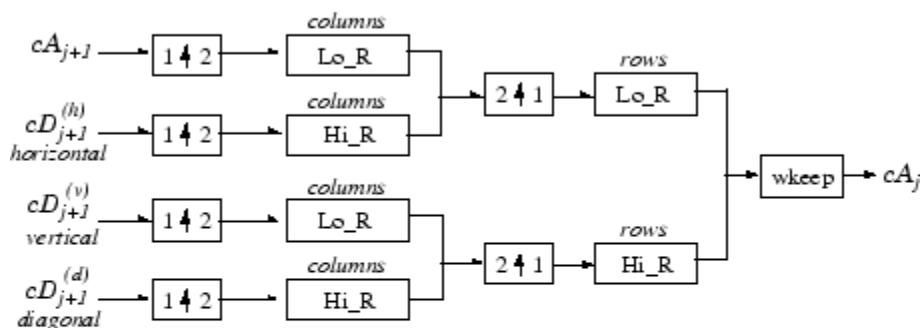
If cA , cH , cV , cD are obtained from an indexed image analysis or a truecolor image analysis, they are m -by- n matrices or m -by- n -by-3 arrays, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Algorithms

Two-Dimensional IDWT

Reconstruction step



Where

$2 \uparrow 1$	Upsample columns: insert zeros at odd-indexed columns.
$1 \uparrow 2$	Upsample rows: insert zeros at odd-indexed rows.
$\begin{matrix} rows \\ X \end{matrix}$	Convolve with filter X the rows of the entry.
$\begin{matrix} columns \\ X \end{matrix}$	Convolve with filter X the columns of the entry.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

`dwt2` | `dwtmode` | `upwlev2`

Introduced before R2006a

idwt3

Single-level inverse discrete 3-D wavelet transform

Syntax

```
X = idwt3(WT)
C = idwt3(WT,TYPE)
```

Description

The `idwt3` command performs a single-level three-dimensional wavelet reconstruction starting from a single-level three-dimensional wavelet decomposition.

`X = idwt3(WT)` computes the single-level reconstructed 3-D array `X`, based on the three-dimensional wavelet decomposition stored in the `WT` structure. This structure contains the following fields.

<code>sizeINI</code>	Size of the three-dimensional array <code>X</code> .
<code>mode</code>	Name of the wavelet transform extension mode.
<code>filters</code>	Structure with 4 fields, <code>LoD</code> , <code>HiD</code> , <code>LoR</code> , <code>HiR</code> , which contain the filters used for DWT.
<code>dec</code>	2 x 2 x 2 cell array containing the coefficients of the decomposition. <code>dec{i,j,k}</code> , <code>i,j,k = 1</code> or <code>2</code> contains the coefficients obtained by low-pass filtering (for <code>i</code> or <code>j</code> or <code>k = 1</code>) or high-pass filtering (for <code>i</code> or <code>j</code> or <code>k = 2</code>).

`C = idwt3(WT,TYPE)` computes the single-level reconstructed component based on the three-dimensional wavelet decomposition. Valid values for `TYPE` are:

- A group of three characters `'xyz'`, one per direction, with `'x'`, `'y'` and `'z'` selected in the set `{'a','d','l','h'}` or in the corresponding uppercase set `{'A','D','L','H'}`, where `'A'` (or `'L'`) specifies low-pass filter and `'D'` (or `'H'`) specifies highpass filter.

- The char 'd' (or 'h' or 'D' or 'H') which specifies the sum of all the components different from the lowpass component.

Examples

Single-Level Three-Dimensional Wavelet Reconstruction

Define the original 3-D data.

```
X = reshape(1:64,4,4,4)
```

```
X =
```

```
X(:,:,1) =
```

```
    1    5    9   13
    2    6   10   14
    3    7   11   15
    4    8   12   16
```

```
X(:,:,2) =
```

```
   17   21   25   29
   18   22   26   30
   19   23   27   31
   20   24   28   32
```

```
X(:,:,3) =
```

```
   33   37   41   45
   34   38   42   46
   35   39   43   47
   36   40   44   48
```

```
X(:,:,4) =
```

```
   49   53   57   61
   50   54   58   62
   51   55   59   63
```

52 56 60 64

Decompose X using 'db1'.

```
wt = dwt3(X, 'db1');
```

Reconstruct X from the coefficients. Verify that the reconstructed data agrees with the original data to machine precision.

```
XR = idwt3(wt);
```

```
dff = max(abs(X-XR))
```

```
dff =
```

```
dff(:, :, 1) =
```

```
1.0e-13 *
```

```
0.0266    0.0355    0.0888    0.1066
```

```
dff(:, :, 2) =
```

```
1.0e-13 *
```

```
0.1066    0.1066    0.2132    0.2132
```

```
dff(:, :, 3) =
```

```
1.0e-13 *
```

```
0.1421    0.1421    0.2132    0.2132
```

```
dff(:, :, 4) =
```

```
1.0e-13 *
```

```
0.3553    0.3553    0.2842    0.2842
```

Compute the reconstructed approximation, which consists of the lowpass component.

```
A = idwt3(wt, 'aaa');
```

Compute the sum of all the components different from the lowpass component.

```
D = idwt3(wt, 'd');
```

Reconstruct the component associated with lowpass in the x and z directions and highpass in the y direction.

```
ADA = idwt3(wt, 'ada');
```

See Also

[dwt3](#) | [wavedec3](#) | [waverec3](#)

Introduced in R2010a

ihaart

Inverse 1-D Haar wavelet transform

Syntax

```
xrec = ihaart(a,d)
xrec = ihaart(a,d,level)
xrec = ihaart( ____,integerflag)
```

Description

`xrec = ihaart(a,d)` returns the inverse 1-D Haar transform, `xrec`, for the approximation coefficients, `a`, and the wavelet coefficients, `d`. Both `a` and `d` are obtained from `haart`.

`xrec = ihaart(a,d,level)` returns the inverse 1-D Haar transform at the specified level.

`xrec = ihaart(____,integerflag)` specifies how the inverse 1-D Haar transform handles integer-valued data, using any of the previous syntaxes.

Examples

Inverse Haar Transform of Noisy Data

Obtain the Haar and inverse Haar transforms of noisy data.

Load the noisy data signal

```
load noisdopp;
```

Obtain the Haar transform of the noisy signal.

```
[a,d] = haart(noisdopp);
```

Reconstruct the data by inverting the Haar transform.

```
xrec = ihaart(a,d);
```

Compare the original and reconstructed data by determining the maximum difference between them. The difference is essentially zero, which indicates a near-perfect reconstruction.

```
max(abs(xrec-noisdopp'))
```

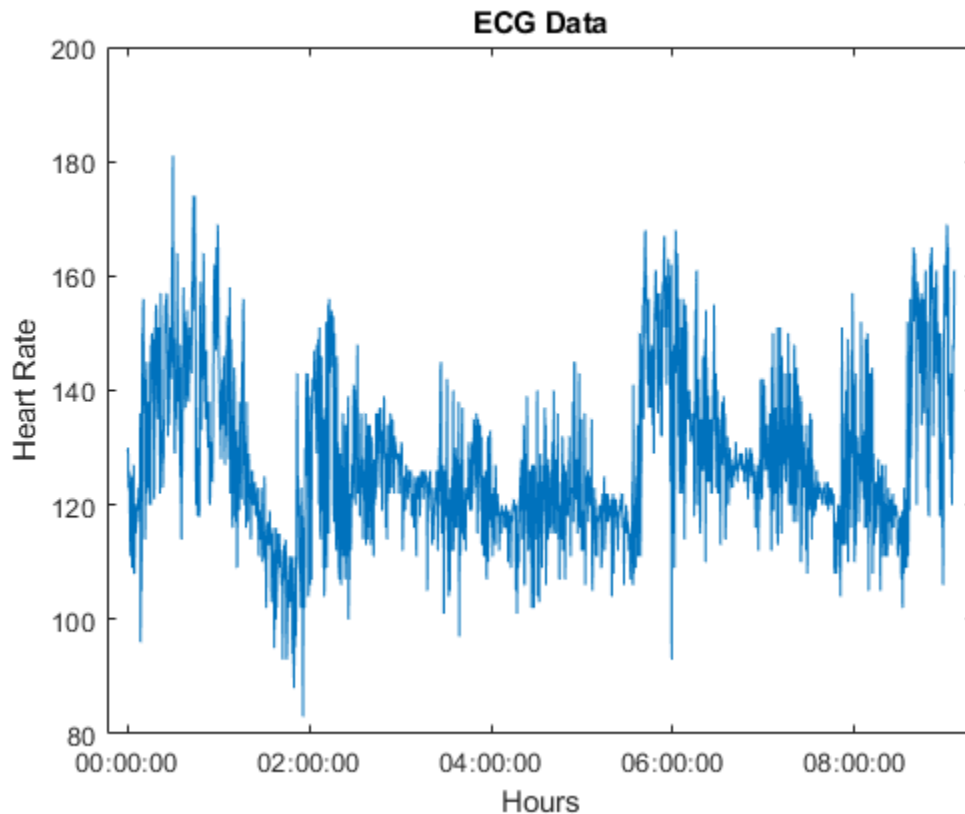
```
ans = 4.4409e-15
```

Inverse Haar Transform of ECG Data

Obtain the Haar transform and inverse Haar transform of ECG heart rate data.

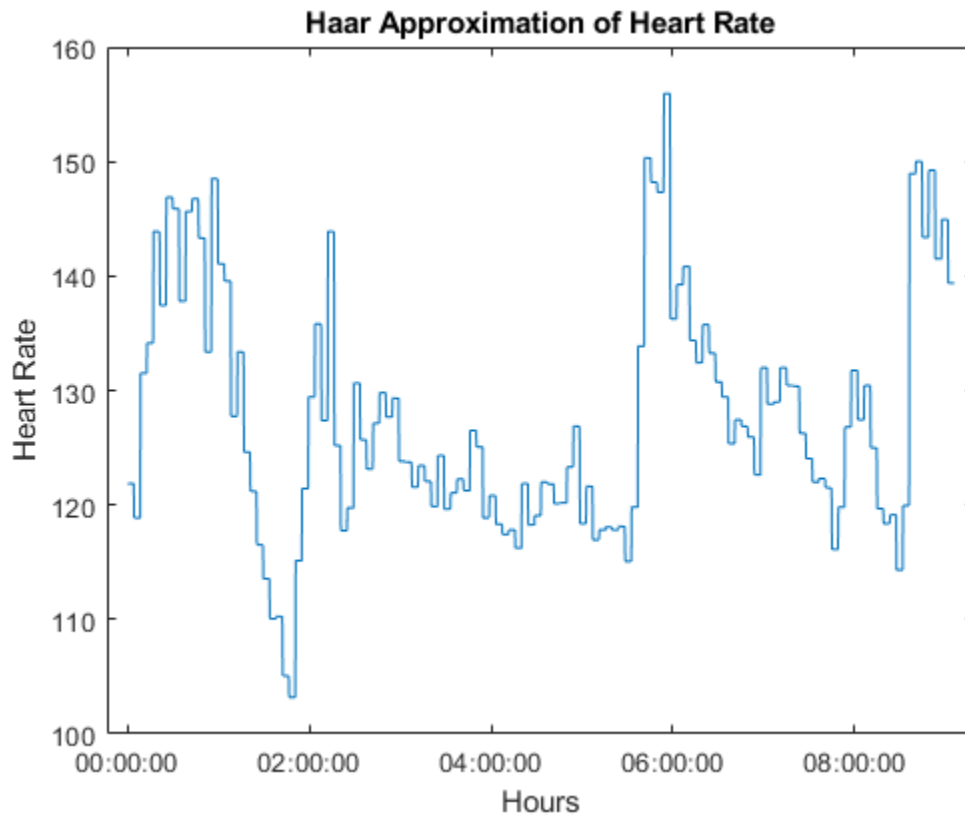
Load and plot the ECG data.

```
load BabyECGData;  
plot(times,HR)  
xlabel('Hours')  
ylabel('Heart Rate')  
title('ECG Data')
```



Obtain the Haar transform and inverse Haar transform. Compare the reconstructed data at level 4 to the original data.

```
[a,d] = haart(HR);  
HaarHR = ihaart(a,d,4);  
figure  
plot(times,HaarHR)  
xlabel('Hours')  
ylabel('Heart Rate')  
title('Haar Approximation of Heart Rate')
```



Inverse Haar Transform of Integer Data

Obtain the Haar and inverse Haar transforms for a series of random integers.

Create the series.

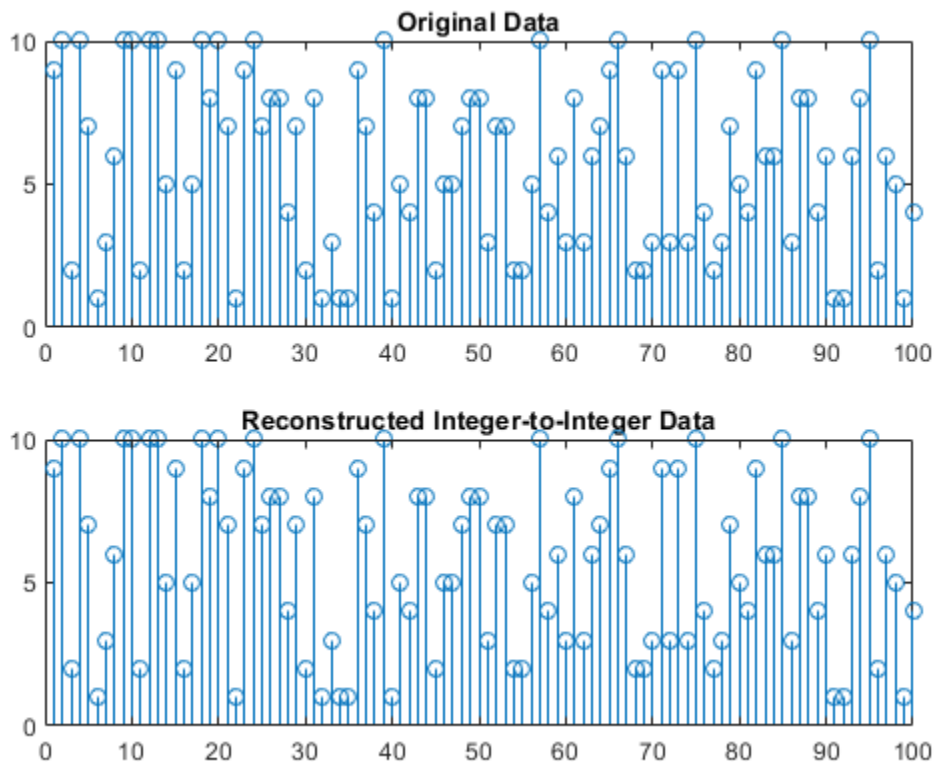
```
x = randi(10,100,1);
```

Obtain the Haar and inverse Haar transforms.

```
[a,d] = haart(x,'integer');  
xrec = ihaart(a,d,'integer');
```

Plot and compare the original and reconstructed data.

```
subplot(2,1,1)
stem(x); title('Original Data')
subplot(2,1,2)
stem(xrec)
title('Reconstructed Integer-to-Integer Data')
```



Determine the maximum difference between the original data values and the reconstructed values. The difference is zero, which indicates perfect reconstruction.

```
max(abs(x(:)-xrec(:)))
```

```
ans = 0
```

Input Arguments

a — Approximation coefficients

scalar | vector | matrix

Approximation coefficients, specified as a scalar, vector, or matrix of coefficients, depending on the level to which the Haar transform was calculated. **a** is an output from the `haart` function.

Approximation, or scaling, coefficients are a lowpass representation of the input. At each level the approximation coefficients are divided into coarser approximation and detail coefficients.

Data Types: `double`

d — Detail coefficients

scalar | vector | matrix | cell array

Detail coefficients, specified as a scalar, vector, matrix, or cell array of wavelet coefficients, which are a highpass representation of the input. **d** is an output from the `haart` function. The number of detail coefficients depends on the selected level and the length of the input. The order of the elements of **d** is from fine to coarse resolution levels.

If **d** is a cell array, it can contain scalars, vectors, or matrices. The level of the Haar transform equals the number of elements in **d**. The coarsest resolution level element of the **d** cell array is a scalar value.

If **d** is a vector or matrix, the Haar transform was computed only down to one level coarser in resolution. If you specify only two levels, the detail coefficient is a scalar.

If **a** and the elements of **d** are vectors, `xrec` is a vector. If **a** and the elements of **d** are matrices, `xrec` is a matrix, where each column is the inverse Haar transform of the corresponding columns in **a** and **d**.

Data Types: `double`

level — Maximum level

0 (default) | nonnegative integer

Maximum level to which to invert the Haar transform, specified as a nonnegative integer. If **d** is a cell array, `level` is less than or equal to `length(d) - 1`. If **d** is a vector or matrix, `level` must equal 0 or be unspecified. The level must be less than the level used to obtain **a** and **d** from `haart`.

integerflag — Integer-valued data handling`'noninteger' (default) | 'integer'`

Integer-valued data handling, specified as either `'noninteger'` or `'integer'`. `'noninteger'` does not preserve integer-valued data, and `'integer'` preserves it. The `'integer'` option applies only if all elements of `a` and `d` are integer-valued. You must have used `'integer'` with `haart` to obtain integer-valued `a` and `d` inputs. The inverse 1-D Haar transform algorithm, however, uses floating-point arithmetic.

Output Arguments

xrec — Inverse 1-D Haar wavelet transform`vector | matrix`

Inverse 1-D Haar wavelet transform, returned as a vector or matrix. If `a` and the elements of `d` are vectors, `xrec` is a vector. If `a` and the elements of `d` are matrices, `xrec` is a matrix, where each column is the inverse 1-D Haar transform of the corresponding columns in `a` and `d`.

Data Types: `double`

See Also

`haart | haart2 | ihaart2`

Topics

“Haar Transforms for Time Series Data and Images”

Introduced in R2016b

ihaart2

Inverse 2-D Haar wavelet transform

Syntax

```
xrec = ihaart2(a,h,v,d)
xrec = ihaart2(a,h,v,d,level)
xrec = ihaart2( ____,integerflag)
```

Description

`xrec = ihaart2(a,h,v,d)` returns the inverse 2-D Haar transform, `xrec`, for the approximation coefficients, `a`, and the horizontal, vertical, and diagonal detail coefficients, `h`, `v`, and `d`. All the inputs, `a`, `h`, `v`, and `d`, are outputs of `haart2`.

`xrec = ihaart2(a,h,v,d,level)` returns the inverse 2-D Haar transform at the specified level.

`xrec = ihaart2(____,integerflag)` specifies how the inverse 2-D Haar transform handles integer-valued data, using any of the previous syntaxes.

Examples

Inverse 2-D Haar Transform of an Image

Obtain the inverse 2-D Haar transform of image and view the reconstructed image.

Load the image and obtain its 2-D Haar transform.

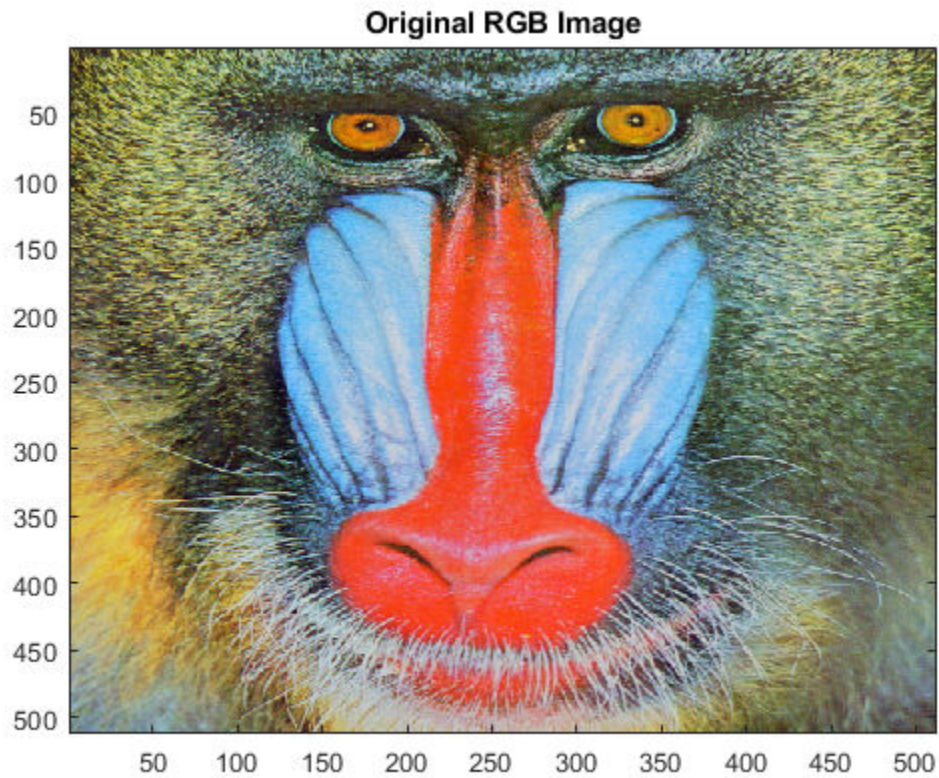
```
im = imread('mandrill.png');
[a,h,v,d] = haart2(im);
```

Use the inverse 2-D Haar transform to reconstruct the image.

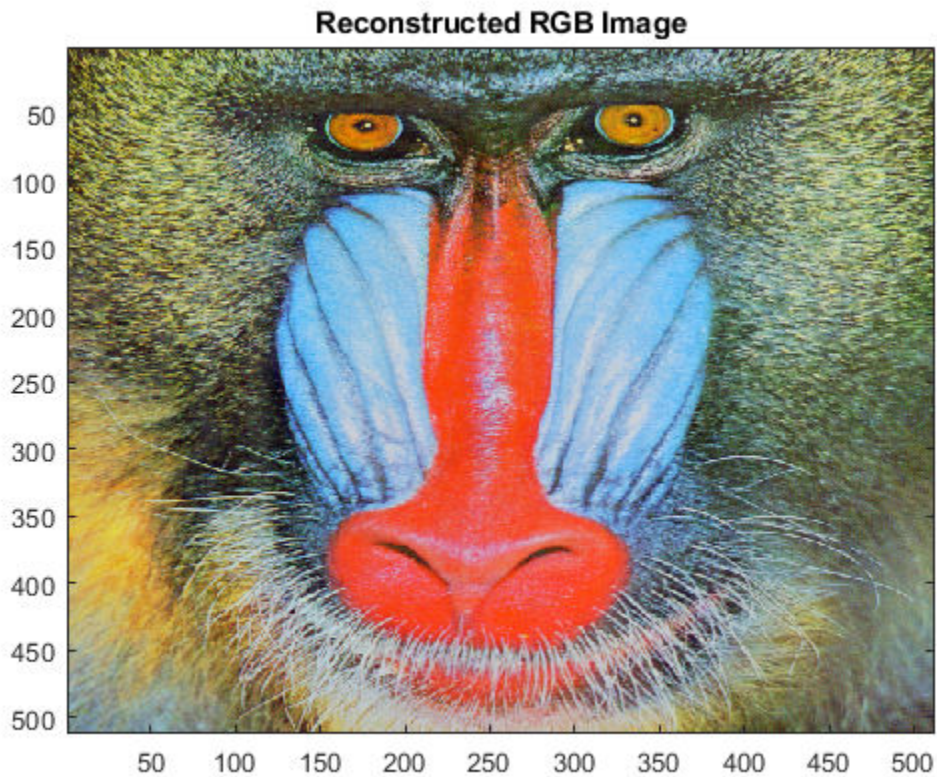
```
xrec = ihaart2(a,h,v,d);
```


Compare the original and reconstructed images.

```
imagesc(im)  
title('Original RGB Image')
```



```
figure  
imagesc(uint8(xrec))  
title('Reconstructed RGB Image')
```

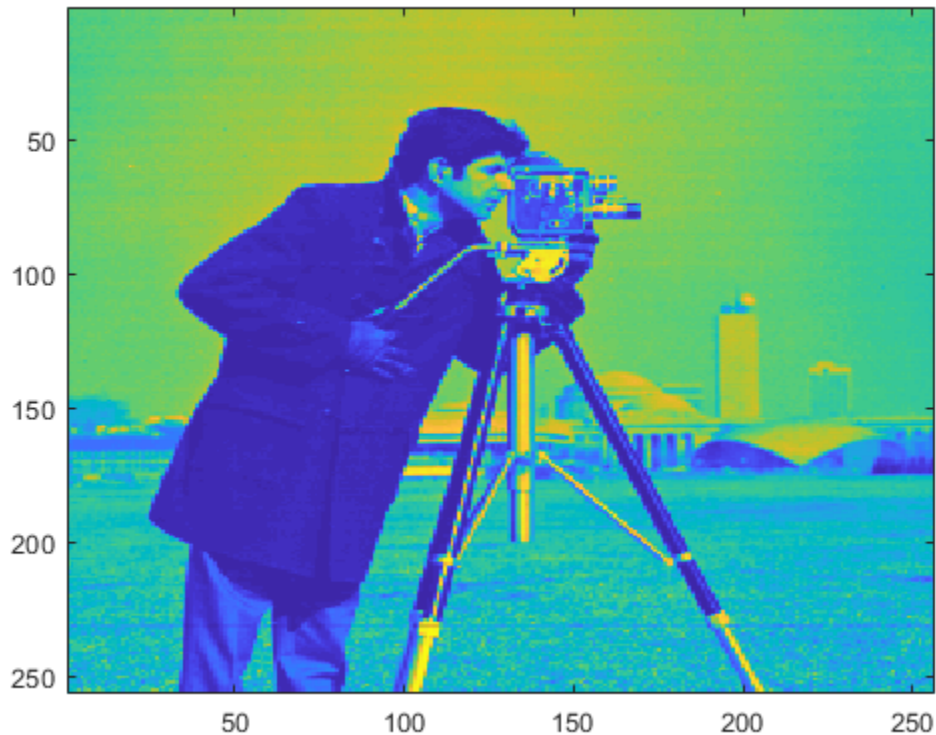


Inverse 2-D Haar Transform of Image Limited to Specified Level

Obtain the 2-D Haar transform of an image limiting the transform to 2 levels.

Load and view the image of a cameraman.

```
im = imread('cameraman.tif');  
imagesc(im)
```

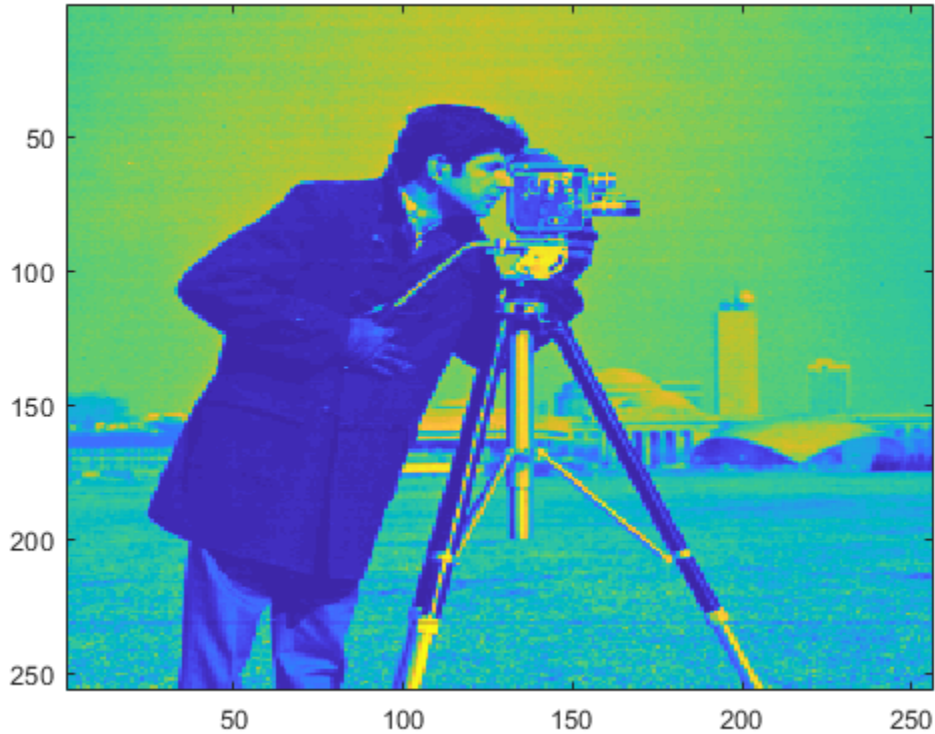


Obtain the 2-D Haar transform using the default maximum number of levels.

```
[a,h,v,d] = haart2(im);
```

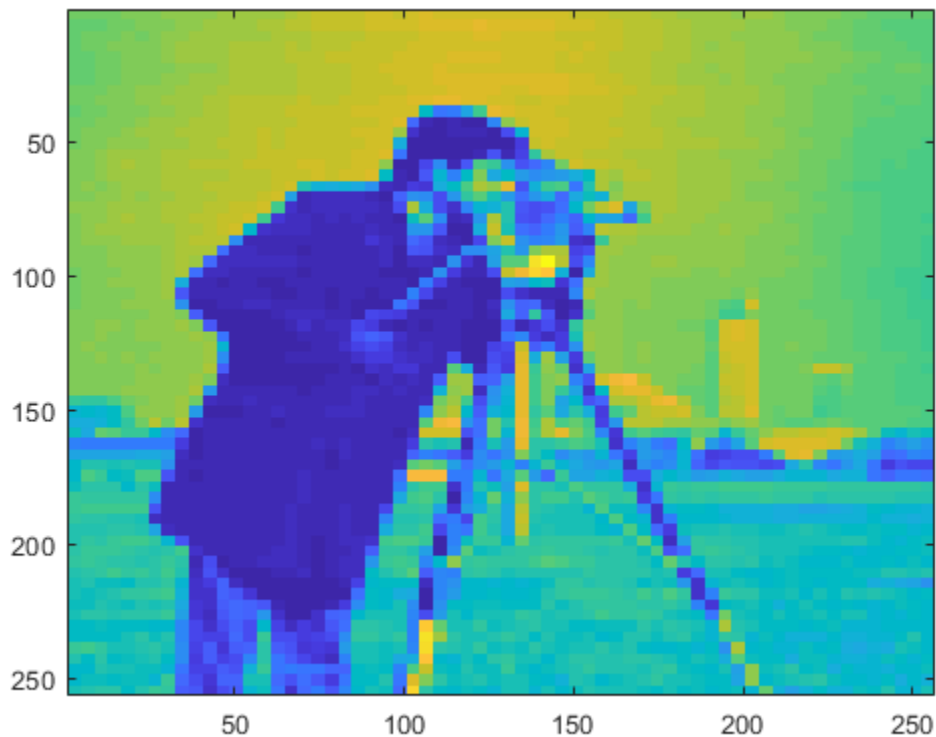
Reconstruct the image using the inverse 2-D Haar transform and view the image. Notice the near-perfect reconstruction.

```
xrec = ihaart2(a,h,v,d);  
imagesc(xrec)
```



Reconstruct and view the image using the inverse 2-D Haar transform, limited to level 2. Level 2 corresponds to the fourth scale because scale is defined as 2^j , where j is the level.

```
xrec1 = ihaart2(a,h,v,d,2);  
imagesc(xrec1)
```



Using fewer levels returns the average of the original image at level 2.

Inverse 2-D Haar Transform of Image Limited to Integer Data

Obtain the 2-D Haar transform of an image limiting the transform to integer data.

Load the image of a cameraman.

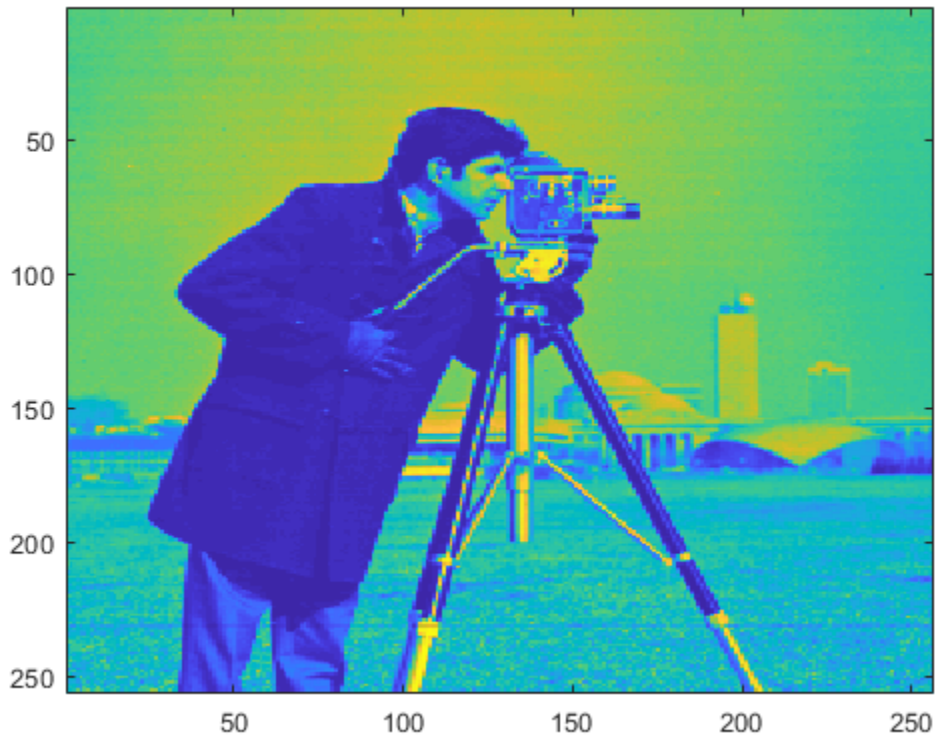
```
im = imread('cameraman.tif');
```

Obtain the 2-D Haar transform using the 'integer' flag.

```
[a,h,v,d]=haart2(im,'integer');
```

Reconstruct the image using the inverse 2-D Haar transform and view the image.

```
xrec = ihaart2(a,h,v,d,'integer');  
imagesc(xrec)
```



Use integer data when you need to reduce the amount of memory used compared to noninteger data.

Input Arguments

a — Approximation coefficients

scalar | matrix

Approximation coefficients, specified as a scalar or matrix of coefficients, depending on the level to which the 2-D Haar transform was calculated. **a** is an output from the `haart2` function. Approximation, or scaling, coefficients are a lowpass representation of the input. If **a** and the elements of **h**, **v**, and **d**, are vectors, **xrec** is a vector. If **a** and the elements of **h**, **v**, and **d** are matrices, **xrec** is a matrix, where each column is the inverse 2-D Haar transform of the corresponding columns in **a** and **h**, **v**, or **d**.

Data Types: double

h — Horizontal detail coefficients

matrix | cell array

Horizontal detail coefficients by level, specified as a matrix or cell array of matrices. **h** is an output from the `haart2` function. If **h** is a matrix, the 2-D Haar transform was computed only down to one level coarser in resolution.

Data Types: double

v — Vertical detail coefficients

matrix or | cell array

Vertical detail coefficients by level, specified as a matrix or cell array of matrices. **v** is an output from the `haart2` function. If **v** is a matrix, the 2-D Haar transform was computed only down to one level coarser in resolution.

Data Types: double

d — Diagonal detail coefficients

matrix or | cell array

Diagonal detail coefficients by level, specified as a matrix or cell array of matrices. **d** is an output from the `haart2` function. If **d** is a matrix, the 2-D Haar transform was computed only down to one level coarser in resolution.

Data Types: double

level — Maximum level

0 (default) | nonnegative integer

Maximum level to which to invert the Haar transform, specified as a nonnegative integer. If `h` is a cell array, `level` is less than or equal to `length(h) - 1`. If `h` is a vector or matrix, `level` must equal `0` or be unspecified.

integerflag — Integer-valued data handling

'noninteger' (default) | 'integer'

Integer-valued data handling, specified as either 'noninteger' or 'integer'. 'noninteger' does not preserve integer-valued data in the 2-D Haar transform, and 'integer' preserves it. The 'integer' option applies only if all elements of inputs, `a`, `h`, `v`, and `d`, are integer-valued. The inverse 2-D Haar transform algorithm, however, uses floating-point arithmetic.

Output Arguments

xrec — Inverse 2-D Haar wavelet transform

matrix

2-D Haar wavelet transform, returned as a matrix.

Data Types: double

See Also

haart | haart2 | ihaart

Topics

“Haar Transforms for Time Series Data and Images”

Introduced in R2016b

ilwt

Inverse 1-D lifting wavelet transform

Syntax

```
X = ilwt(AD_In_Place,W)
X = ilwt(CA,CD,W)
X = ilwt(AD_In_Place,W,LEVEL)
X = ILWT(CA,CD,W,LEVEL)
X = ilwt(AD_In_Place,W,LEVEL,'typeDEC',typeDEC)
X = ilwt(CA,CD,W,LEVEL,'typeDEC',typeDEC)
```

Description

`ilwt` performs a 1-D lifting wavelet reconstruction with respect to a particular lifted wavelet that you specify.

`X = ilwt(AD_In_Place,W)` computes the reconstructed vector `X` using the approximation and detail coefficients vector `AD_In_Place` obtained by a lifting wavelet reconstruction. `W` is a lifted wavelet name (see `liftwave`).

`X = ilwt(CA,CD,W)` computes the reconstructed vector `X` using the approximation coefficients vector `CA` and detail coefficients vector `CD` obtained by a lifting wavelet reconstruction.

`X = ilwt(AD_In_Place,W,LEVEL)` or `X = ILWT(CA,CD,W,LEVEL)` computes the lifting wavelet reconstruction, at level `LEVEL`.

`X = ilwt(AD_In_Place,W,LEVEL,'typeDEC',typeDEC)` or `X = ilwt(CA,CD,W,LEVEL,'typeDEC',typeDEC)` with `typeDEC = 'w'` or `'wp'` computes the wavelet or the wavelet packet decomposition using lifting, at level `LEVEL`.

Instead of a lifted wavelet name, you may use the associated lifting scheme `LS`: `X = ilwt(...,LS,...)` instead of `X = ILWT(...,W,...)`.

For more information about lifting schemes, see `lsinfo`.

Examples

```
% Start from the Haar wavelet and get the
% corresponding lifting scheme.
lshaar = liftwave('haar');

% Add a primal ELS to the lifting scheme.
els = {'p',[-0.125 0.125],0};
lsnew = addlift(lshaar,els);

% Perform LWT at level 1 of a simple signal.
x = 1:8;
[cA,cD] = lwt(x,lsnew);

% Perform integer LWT of the same signal.
lshaarInt = liftwave('haar','int2int');
lsnewInt = addlift(lshaarInt,els);
[cAint,cDint] = lwt(x,lsnewInt);

% Invert the two transforms.
xRec = ilwt(cA,cD,lsnew);
err = max(max(abs(x-xRec)))

err =

    4.4409e-016

xRecInt = ilwt(cAint,cDint,lsnewInt);
errInt = max(max(abs(x-xRecInt)))

errInt =

    0
```

See Also

lwt

Introduced before R2006a

ilwt2

Inverse 2-D lifting wavelet transform

Syntax

```
X = ilwt2(AD_In_Place,W)
X = ilwt2(CA,CH,CV,CD,W)
X = ilwt2(AD_In_Place,W,LEVEL)
X = ILWT2(CA,CH,CV,CD,W,LEVEL)
X = ilwt2(AD_In_Place,W,LEVEL,'typeDEC',typeDEC)
X = ilwt2(CA,CH,CV,CD,W,LEVEL,'typeDEC',typeDEC)
```

Description

`ilwt2` performs a 2-D lifting wavelet reconstruction with respect to a particular lifted wavelet that you specify.

`X = ilwt2(AD_In_Place,W)` computes the reconstructed matrix `X` using the approximation and detail coefficients matrix `AD_In_Place`, obtained by a lifting wavelet decomposition. `W` is a lifted wavelet name (see `liftwave`).

`X = ilwt2(CA,CH,CV,CD,W)` computes the reconstructed matrix `X` using the approximation coefficients vector `CA` and detail coefficients vectors `CH`, `CV`, and `CD` obtained by a lifting wavelet decomposition.

`X = ilwt2(AD_In_Place,W,LEVEL)` or `X = ILWT2(CA,CH,CV,CD,W,LEVEL)` computes the lifting wavelet reconstruction, at level `LEVEL`.

`X = ilwt2(AD_In_Place,W,LEVEL,'typeDEC',typeDEC)` or `X = ilwt2(CA,CH,CV,CD,W,LEVEL,'typeDEC',typeDEC)` with `typeDEC = 'w'` or `'wp'` computes the wavelet or the wavelet packet decomposition using lifting, at level `LEVEL`.

Instead of a lifted wavelet name, you may use the associated lifting scheme `LS`: `X = ilwt2(...,LS,...)` instead of `X = ilwt2(...,W,...)`.

For more information about lifting schemes, see `lsinfo`.

Examples

```
% Start from the Haar wavelet and get the
% corresponding lifting scheme.
lshaar = liftwave('haar');

% Add a primal ELS to the lifting scheme.
els = {'p',[-0.125 0.125],0};
lsnew = addlift(lshaar,els);

% Perform LWT at level 1 of a simple image.
x = reshape(1:16,4,4);
[cA,cH,cV,cD] = lwt2(x,lsnew);

% Perform integer LWT of the same image.
lshaarInt = liftwave('haar','int2int');
lsnewInt = addlift(lshaarInt,els);
[cAint,cHint,cVint,cDint] = lwt2(x,lsnewInt);

% Invert the two transforms.
xRec = ilwt2(cA,cH,cV,cD,lsnew);
err = max(max(abs(x-xRec)))

err =

    0

xRecInt = ilwt2(cAint,cHint,cVint,cDint,lsnewInt);
errInt = max(max(abs(x-xRecInt)))

errInt =

    0
```

Tips

If `AD_In_Place` or `cA,cH,cV,cD` are obtained from an indexed image analysis or a truecolor image analysis, they are `m-by-n` matrices or `m-by-n-by-3` arrays, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

See Also

lwt2

Introduced before R2006a

imlpt

Inverse multiscale local 1-D polynomial transform

Syntax

```
y = imlpt(coefs,T,coefsPerLevel,scalingMoments)
y = imlpt( ____,Name,Value)
```

Description

`y = imlpt(coefs,T,coefsPerLevel,scalingMoments)` returns the inverse multiscale local polynomial 1-D transform (MLPT) of `coefs`. The inputs to `imlpt` must be the outputs of `mlpt`.

`y = imlpt(____,Name,Value)` specifies `mlpt` properties using one or more `Name,Value` pair arguments and the input arguments from the previous syntax.

Examples

Multiscale Local 1-D Polynomial Transform and Inverse Transform

Create a signal with nonuniform sampling and verify good reconstruction when performing the `mlpt` and `imlpt`.

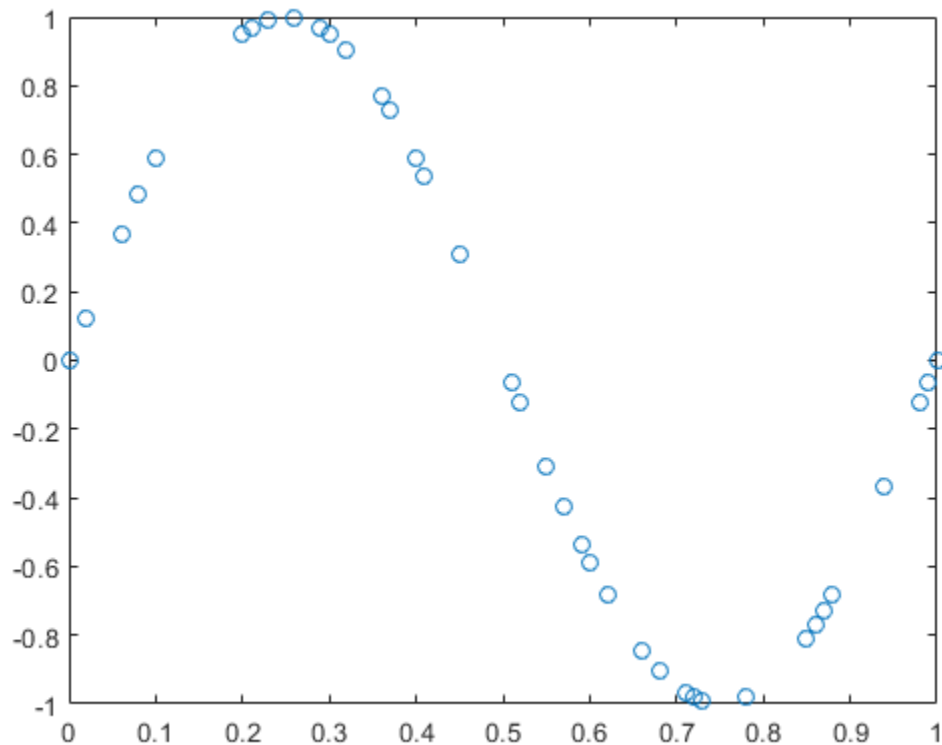
Create and plot a sine wave with non-uniform sampling.

```
timeVector = 0:0.01:1;
sineWave = sin(2*pi*timeVector)';

samplesToErase = randi(100,100,1);
sineWave(samplesToErase) = [];
timeVector(samplesToErase) = [];

figure(1)
```

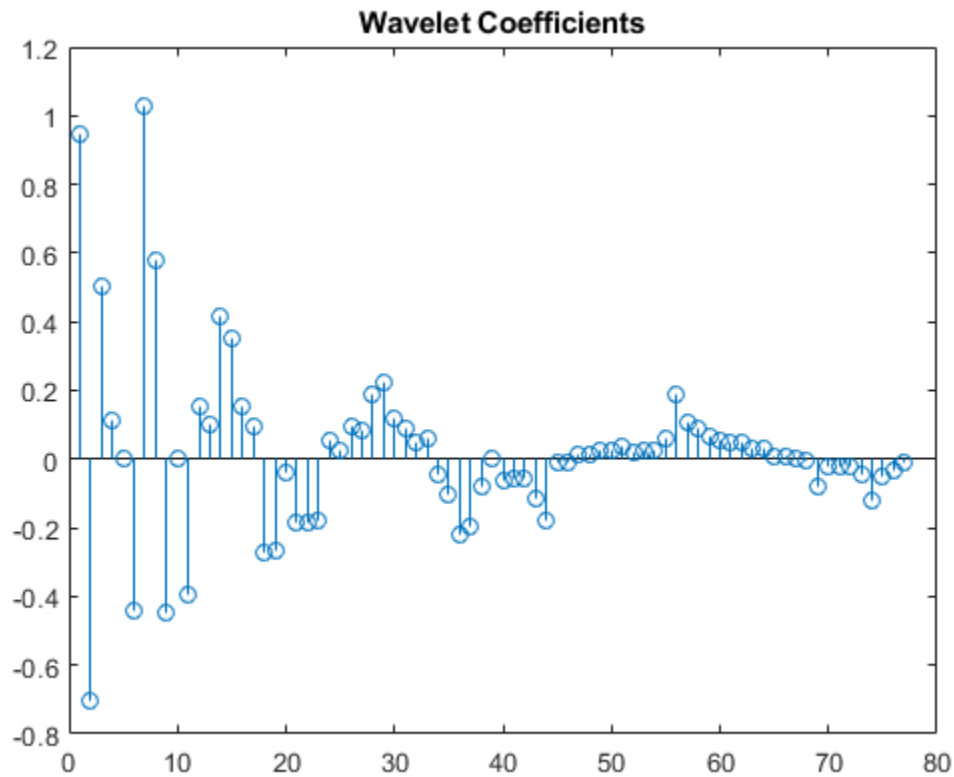
```
plot(timeVector,sineWave,'o')  
hold on
```



Perform the multiscale local 1-D polynomial transform (mlpt) on the signal. Visualize the coefficients.

```
[coefs,T,coefsPerLevel,scalingMoments] = mlpt(sineWave,timeVector);
```

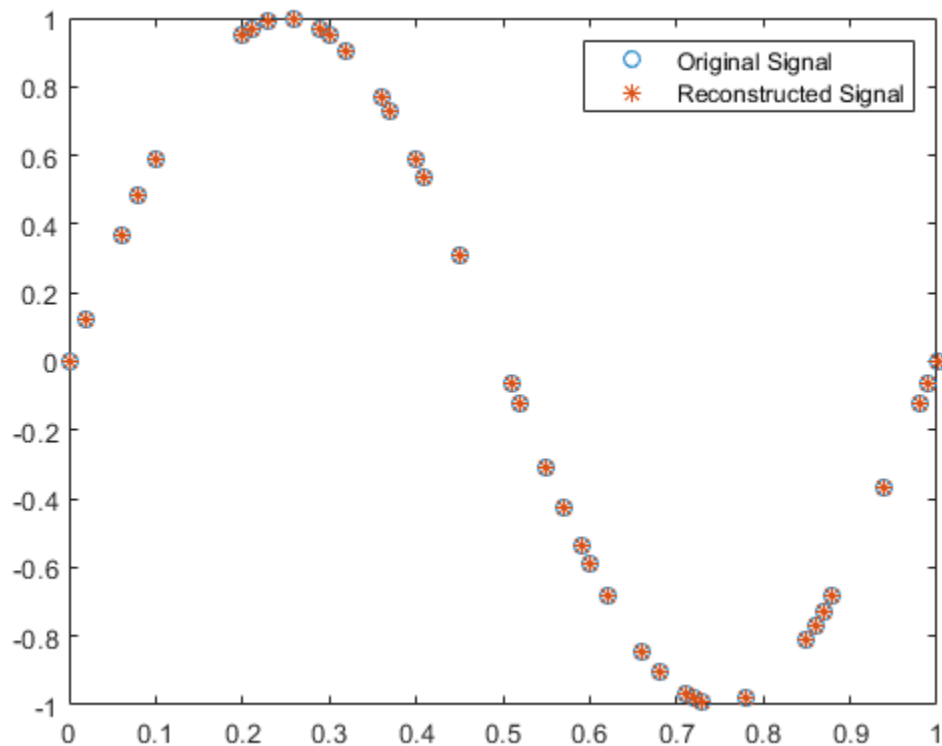
```
figure(2)  
stem(coefs)  
title('Wavelet Coefficients')
```



Perform the inverse multiscale local 1-D polynomial transform (`imlpt`) on the coefficients. Visualize the reconstructed signal.

```
y = imlpt(coefs,T,coefsPerLevel,scalingMoments);
```

```
figure(1)  
plot(T,y,'*')  
legend('Original Signal','Reconstructed Signal')  
hold off
```

Look at the total error to verify good reconstruction.

```
reconstructionError = sum(abs(y-sineWave))
```

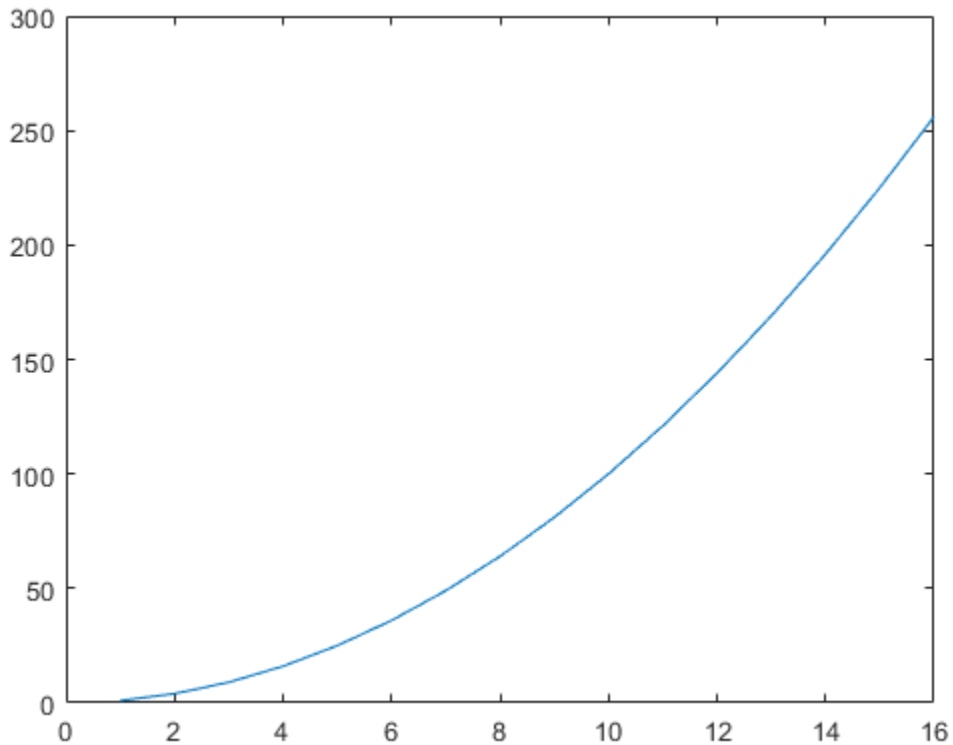
```
reconstructionError = 1.7552e-15
```

Specify Nondefault Dual Moments

Specify nondefault dual moments by using the `m_lpt` function. Compare the results of analysis and synthesis using the default and nondefault dual moments.

Create an input signal and visualize it.

```
T = (1:16)';  
x = T.^2;  
plot(x)  
hold on
```



Perform the forward and inverse transform for the input signal using the default and nondefault dual moments.

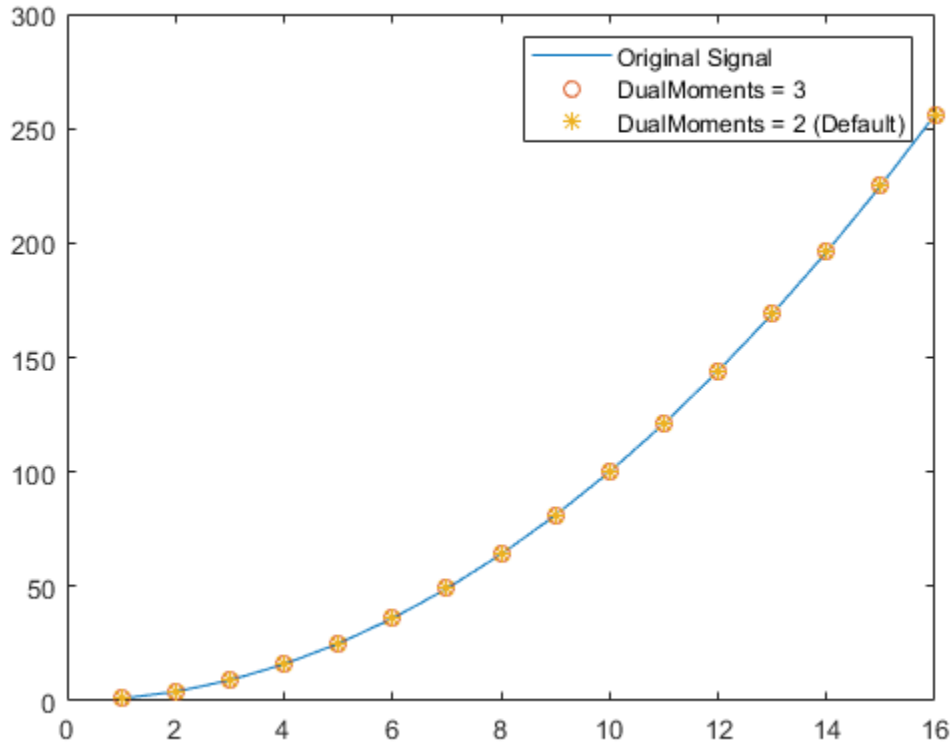
```
[w2,t2,nj2,scalingmoments2] = mlpt(x,T);  
y2 = imlpt(w2,t2,nj2,scalingmoments2);  
  
[w3,t3,nj3,scalingmoments3] = mlpt(x,T,'dualmoments',3);  
y3 = imlpt(w3,t3,nj3,scalingmoments3,'dualmoments',3);
```

Plot the reconstructed signal and verify perfect reconstruction using both the default and nondefault dual moments.

```
plot(y2,'o')
plot(y3,'*')
legend('Original Signal', ...
       'DualMoments = 3', ...
       'DualMoments = 2 (Default)');

fprintf('\nMean Reconstruction Error:\n');
Mean Reconstruction Error:
fprintf(' - Nondefault dual moments: %0.2f\n',mean(abs(y3-x)));
- Nondefault dual moments: 0.00
fprintf(' - Default dual moments: %0.2f\n\n',mean(abs(y2-x)));
- Default dual moments: 0.00

hold off
```



Input Arguments

coefs — MLPT coefficients

vector | matrix

MLPT coefficients, specified as a vector or matrix of MLPT coefficients returned by the `mlpt` function.

Data Types: double

T — Sampling instants corresponding to output

vector | duration array

Sampling instants corresponding to y , specified as a vector or `duration` array of increasing values returned by the `mLPT` function.

Data Types: `double` | `duration`

coefsPerLevel — Coefficients per resolution level

vector

Coefficients per resolution level, specified as a vector containing the number of coefficients at each resolution level in `coefs`. `coefsPerLevel` is an output argument of the `mLPT` function.

The elements of `coefsPerLevel` are organized as follows:

- `coefsPerLevel(1)` — Number of approximation coefficients at the coarsest resolution level.
- `coefsPerLevel(i)` — Number of detail coefficients at resolution level i , where $i = \text{numLevel} - i + 2$ for $i = 2, \dots, \text{numLevel} + 1$. `numLevel` is the number of resolution levels used to calculate the MLPT. `numLevel` is inferred from `coefsPerLevel`: `numLevel = length(coefsPerLevel)-1`.

The smaller the index i , the lower the resolution. The MLPT is two times redundant in the number of detail coefficients, but not in the number of approximation coefficients.

Data Types: `double`

scalingMoments — Scaling function moments

matrix

Scaling function moments, specified as a `length(coefs)`-by- P matrix, where P is the number of primal moments specified by the MLPT.

Data Types: `double`

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'DualMoments', 3` computes a transform using three dual vanishing moments.

DualMoments — Number of dual vanishing moments

2 (default) | 3 | 4

Number of dual vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of 'DualMoments' and 2, 3 or 4. The number of dual moments must match the number used by `mlpt`.

Data Types: `double`

Output Arguments

y — Reconstructed signal

vector | matrix

Reconstructed signal, returned as a vector or matrix, depending on the inputs to the `mlpt` function.

Data Types: `double`

Algorithms

Maarten Jansen developed the theoretical foundation of the multiscale local polynomial transform (MLPT) and algorithms for its efficient computation [1][2][3]. The MLPT uses a lifting scheme, wherein a kernel function smooths fine-scale coefficients with a given bandwidth to obtain the coarser resolution coefficients. The `mlpt` function uses only local polynomial interpolation, but the technique developed by Jansen is more general and admits many other kernel types with adjustable bandwidths [2].

References

- [1] Jansen, M. "Multiscale Local Polynomial Smoothing in a Lifted Pyramid for Non-Equispaced Data". *IEEE Transactions on Signal Processing*. Vol. 61, Number 3, 2013, pp.545-555.
- [2] Jansen, M., and M. Amghar. "Multiscale local polynomial decompositions using bandwidths as scales". *Statistics and Computing* (forthcoming). 2016.
- [3] Jansen, M., and Patrick Oonincx. *Second Generation Wavelets and Applications*. London: Springer, 2005.

See Also

`mlpt` | `mlptdenoise` | `mlptrecon`

Topics

Smoothing Nonuniformly Sampled Data

Introduced in R2017a

imodwpt

Inverse maximal overlap discrete wavelet packet transform

Syntax

```
xrec = imodwpt(coefs)
xrec = imodwpt(coefs,wname)
xrec = imodwpt(coefs,lo,hi)
```

Description

`xrec = imodwpt(coefs)` returns the inverse maximal overlap discrete wavelet packet transform (inverse MODWPT), in `xrec`. The inverse transform is for the terminal node coefficient matrix (`coefs`) obtained using `modwpt` with the default length 18 Fejér-Korovkin ('fk18') wavelet.

`xrec = imodwpt(coefs,wname)` returns the inverse MODWPT using the orthogonal filter specified by `wname`. This filter must be the same filter used in `modwpt`.

`xrec = imodwpt(coefs,lo,hi)` returns the inverse MODWPT using the orthogonal scaling filter, `lo`, and wavelet filter, `hi`.

Examples

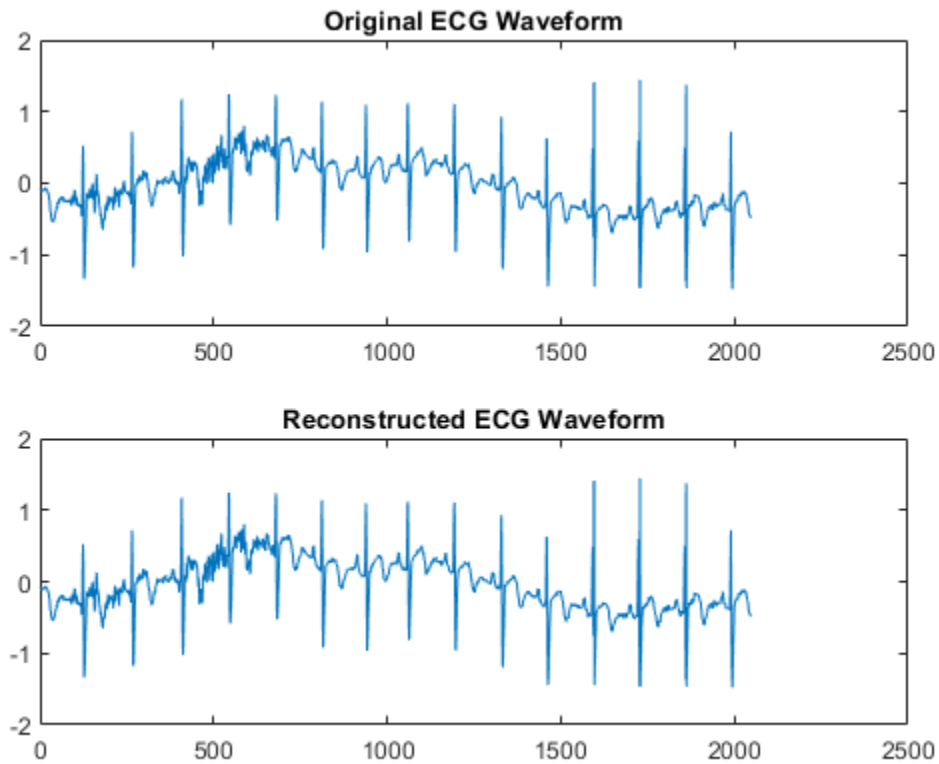
Perfect Reconstruction with the Inverse MODWPT

Obtain the MODWPT of an ECG waveform and demonstrate perfect reconstruction using the inverse MODWPT.

```
load wecg;
wpt = modwpt(wecg);
xrec = imodwpt(wpt);
subplot(2,1,1)
plot(wecg);
```



```
title('Original ECG Waveform');  
subplot(2,1,2)  
plot(xrec);  
title('Reconstructed ECG Waveform');
```



Find the largest absolute difference between the original signal and the reconstruction. The difference is on the order of 10^{-11} , which demonstrates perfect reconstruction.

```
max(abs(wecg-xrec'))
```

```
ans = 1.7902e-11
```

Inverse MODWPT Using Daubechies Extremal Phase Wavelet

Obtain the MODWPT of Southern Oscillation Index data using the Daubechies extremal phase wavelet with two vanishing moments ('db2'). Reconstruct the signal using the inverse MODWPT.

```
load soi;
wsoi = modwpt(soi,'db2');
xrec = imodwpt(wsoi,'db2');
```

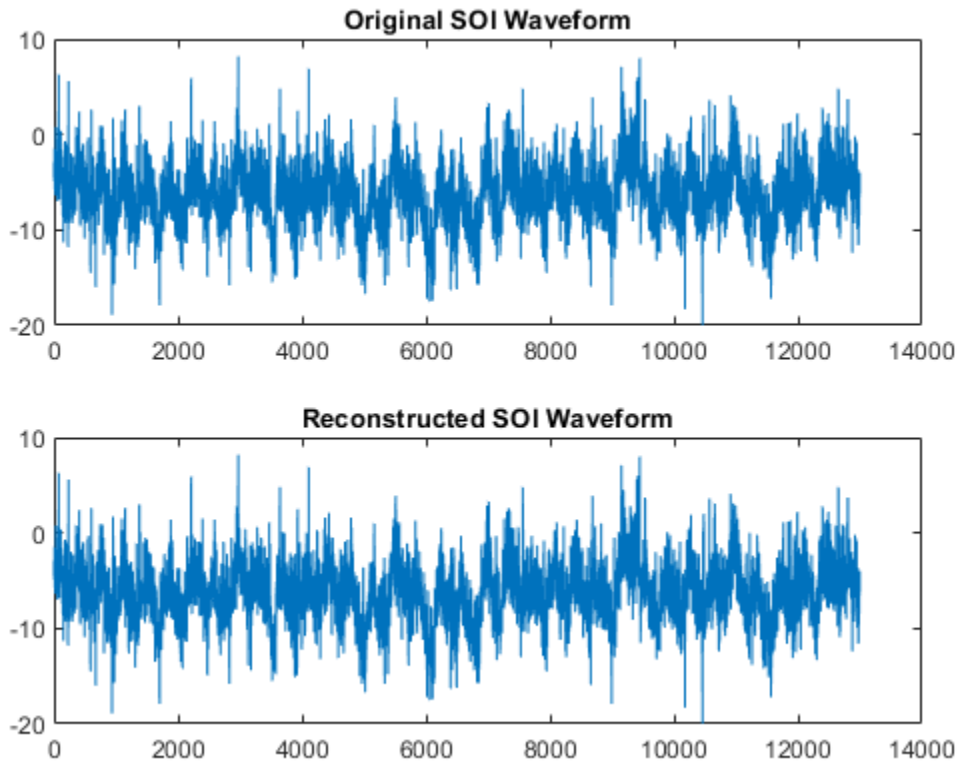
Inverse MODWPT Using Scaling and Wavelet Filters

Obtain the MODWPT of Southern Oscillation Index data using specified scaling and wavelets filters with the Daubechies extremal phase wavelet with two vanishing moments ('db2').

```
load soi;
[lo,hi] = wfilters('db2');
wpt = modwpt(soi,lo,hi);
xrec = imodwpt(wpt,lo,hi);
```

Plot the original SOI waveform and the reconstructed waveform.

```
subplot(2,1,1)
plot(soi)
title('Original SOI Waveform');
subplot(2,1,2)
plot(xrec)
title('Reconstructed SOI Waveform')
```



Input Arguments

coefs — Terminal node coefficients

matrix

Terminal node coefficients of a wavelet packet tree, specified as a matrix. You must obtain the coefficient matrix from `modwpt` using the `'FullTree', false` option. `'FullTree', false` is the default value of `modwpt`.

Data Types: double

wname — Synthesizing wavelet filter

fk18 (default) | character vector | string scalar

Synthesizing wavelet filter used to invert the MODWPT, specified as a character vector or string scalar. The specified wavelet must be the same wavelet as used in the analysis with `modwpt`.

lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. `lo` must be the same scaling filter as used in the analysis with `modwpt`. You cannot specify both a scaling-wavelet filter pair and a `wname` filter.

hi — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. `hi` must be the same wavelet filter used in the analysis with `modwpt`. You cannot specify both a scaling-wavelet filter pair and a `wname` filter.

Output Arguments

xrec — Inverse maximal overlap discrete wavelet packet transform

row vector

Inverse maximal overlap discrete wavelet packet transform, returned as a row vector. The inverse transform is the reconstructed version of the original signal based on the MODWPT terminal node coefficients. `xrec` has the same number of columns as the input `coefs` matrix.

References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.
- [2] Walden, A.T., and A. Contreras Cristan. "The phase-corrected undecimated discrete wavelet packet transform and its application to interpreting the timing of events." *Proceedings of the Royal Society of London A*. Vol. 454, Issue 1976, 1998, pp. 2243-2266.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

[modwpt](#) | [modwptdetails](#)

Introduced in R2016a

imodwt

Inverse maximal overlap discrete wavelet transform

Syntax

```
xrec = imodwt(w)
xrec = imodwt(w,wname)
xrec = imodwt(w,Lo,Hi)
xrec = imodwt( ____,lev)
xrec = imodwt( ____, 'reflection')
```

Description

`xrec = imodwt(w)` returns in `xrec` a reconstructed version of the signal. The reconstructed signal is based on `w`, the maximal overlap discrete wavelet transform (MODWT) coefficients and on the level of reconstruction, which defaults to zero.

`xrec = imodwt(w,wname)` reconstructs the signal using `wname`, the orthogonal wavelet. `wname` must be the same wavelet used to analyze the signal input to `modwt`. The reconstruction is up to level 0, which is a perfect reconstruction of the original signal.

`xrec = imodwt(w,Lo,Hi)` reconstructs the signal using the orthogonal scaling filter `Lo` and the wavelet filter `Hi`. The `Lo` and `Hi` filters must be the same filters used to analyze the signal input to `modwt`. The reconstruction is up to level 0, which is a perfect reconstruction of the original signal.

`xrec = imodwt(____,lev)` reconstructs the signal up to level `lev`. `xrec` is a projection onto the scaling space at level `lev`.

`xrec = imodwt(____, 'reflection')` uses the reflection boundary condition in the reconstruction. If you specify `'reflection'`, `imodwt` assumes that the length of the original signal length is one half the number of columns in the input coefficient matrix. By default, `imodwt` assumes periodic signal extension at the boundary.

Examples

Perfect Reconstruction with the Inverse MODWT

Obtain the MODWT of an ECG signal and demonstrate perfect reconstruction.

Load the ECG signal data and obtain the MODWT.

```
load wecg;
```

Obtain the MODWT and the Inverse MODWT.

```
w = modwt(wecg);  
xrec = imodwt(w);
```

Use the L-infinity norm to show that the difference between the original signal and the reconstruction is extremely small. The largest absolute difference between the original signal and the reconstruction is on the order of 10^{-12} , which demonstrates perfect reconstruction.

```
norm(abs(xrec' - wecg), Inf)  
ans = 2.3253e-12
```

Inverse MODWT with Specified Wavelet

Obtain the MODWT of Deutsche Mark-U.S. Dollar exchange rate data and demonstrate perfect reconstruction.

Load the Deutsche Mark-U.S. Dollar exchange rate data.

```
load DM_USD;
```

Obtain the MODWT and the Inverse MODWT using the 'db2' wavelet.

```
wdm = modwt(DM_USD, 'db2');  
xrec = imodwt(wdm, 'db2');
```

Use the L-infinity norm to show that the difference between the original signal and the reconstruction is extremely small. The largest absolute difference between the original

signal and the reconstruction is on the order of 10^{-13} , which demonstrates perfect reconstruction.

```
norm(abs(xrec' - DM_USD), Inf)
ans = 1.6362e-13
```

Inverse MODWT with Specified Filters

Obtain the MODWT of an ECG signal using the Fejer-Korovkin filters.

Load the ECG data.

```
load wecg;
```

Create the 8-coefficient Fejer-Korovkin filters.

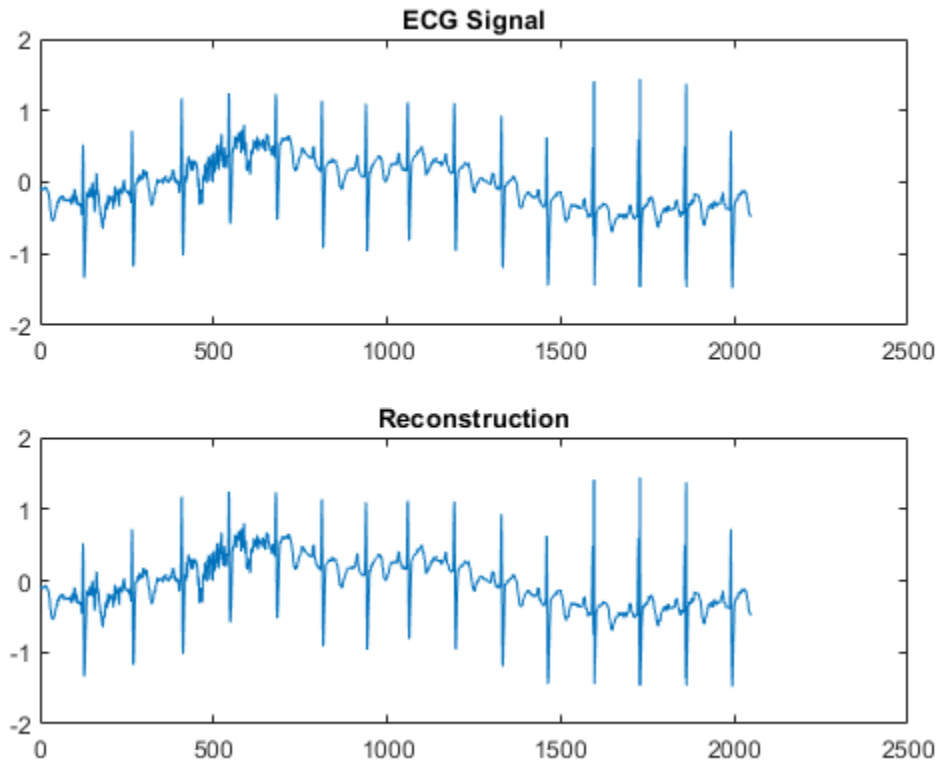
```
[Lo,Hi] = wfilters('fk8');
```

Obtain the MODWT and Inverse MODWT.

```
wtecg = modwt(wecg,Lo,Hi);
xrec = imodwt(wtecg,Lo,Hi);
```

Plot the original data and the reconstruction.

```
subplot(2,1,1)
plot(wecg)
title('ECG Signal');
subplot(2,1,2)
plot(xrec)
title('Reconstruction')
```

Obtain Projection onto Scaling Space

Obtain the MODWT of an ECG signal down to the maximum level and obtain the projection of the ECG signal onto the scaling space at level 3.

Load the ECG data.

```
load wecg;
```

Obtain the MODWT.

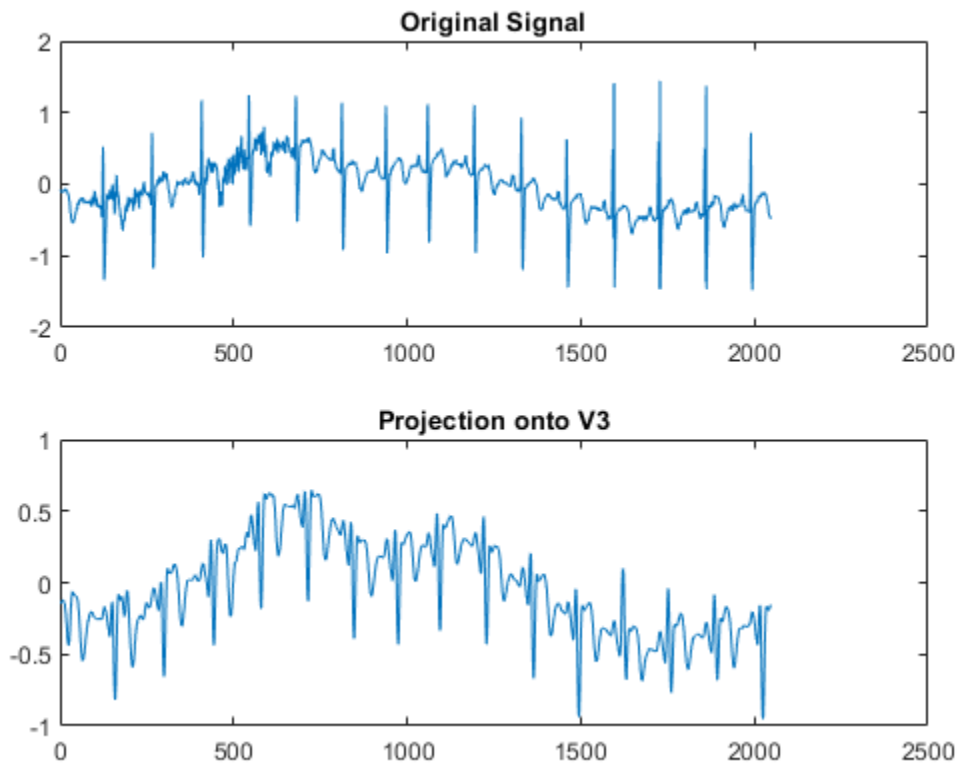
```
wtecg = modwt(wecg);
```

Obtain the projection of the ECG signal onto V_3 , the scaling space at level three by using the `imodwt` function.

```
v3proj = imodwt(wtecg,3);
```

Plot the original signal and the projection.

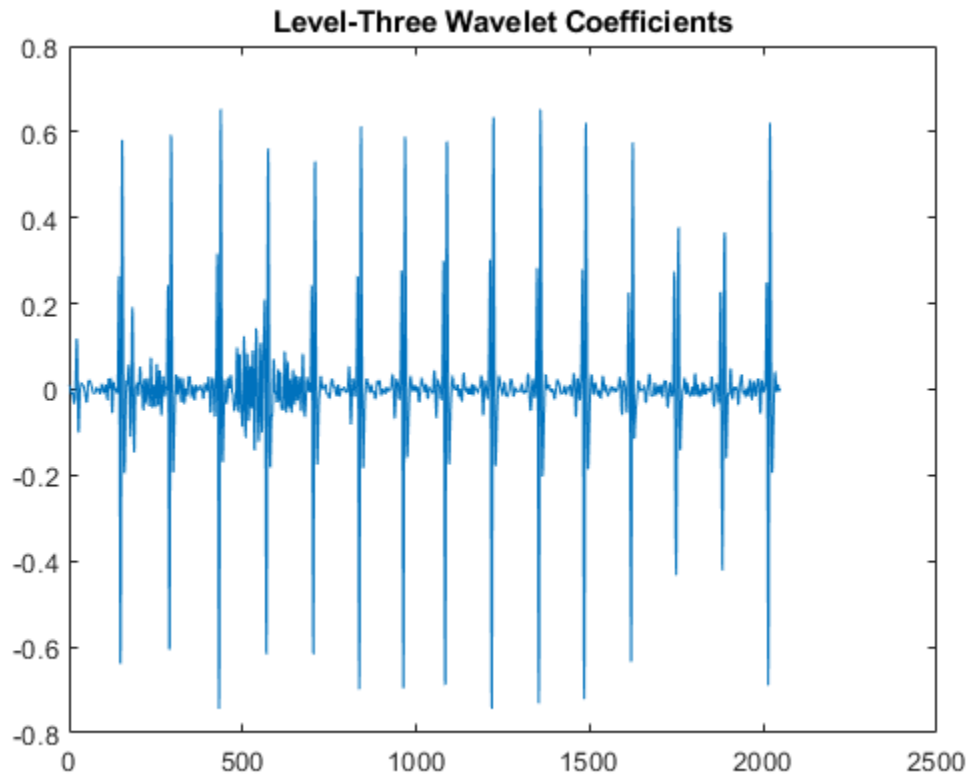
```
subplot(2,1,1)  
plot(wecg)  
title('Original Signal')  
subplot(2,1,2)  
plot(v3proj)  
title('Projection onto V3')
```



Note that the spikes characteristic of the R waves in the ECG are missing in the V_3 approximation. You can see the missing details by examining the wavelet coefficients at level three.

Plot the level-three wavelet coefficients.

```
figure
plot(wtecg(3,:))
title('Level-Three Wavelet Coefficients')
```



Inverse MODWT with Reflection Boundary

Obtain the inverse MODWT using reflection boundary handling for Southern Oscillation Index data. The sampling period is one day. `imodwt` with the `'reflection'` option assumes that the input matrix, which is the `modwt` output, is twice the length of the original signal length. `imodwt` reflection boundary handling reduces the number of wavelet and scaling coefficients at each scale by half.

```
load soi;
wsoi = modwt(soi,4,'reflection');
xrecsoi = imodwt(wsoi,'reflection');
```

Use the L-infinity norm to show that the difference between the original signal and the reconstruction is extremely small. The largest absolute difference between the original signal and the reconstruction is on the order of 10^{-11} , which demonstrates perfect reconstruction.

```
norm(abs(xrecsoi'-soi),Inf)
ans = 1.6433e-11
```

Input Arguments

w — MODWT transform

matrix

MODWT transform, specified as a matrix of size $L+1$ -by- N . `w` is the output of `modwt`, which is the MODWT of an N -point input signal down to level L . By default, `imodwt` assumes that you obtained the MODWT using the `'sym4'` wavelet with periodic boundary handling.

Data Types: `double`

wname — Synthesis wavelet

`'sym4'` (default) | `'dbN'` | `'coifN'` | `'haar'` | `'fkN'` | `'symN'`

Synthesis wavelet, specified as one of the following:

- `'haar'` — Haar wavelet
- `'dbN'` — Extremal phase Daubechies wavelet with N vanishing moments, where N is a positive integer from 1 to 45.

- 'symN' — Symlets wavelet with N vanishing moments, where N is a positive integer from 2 to 45.
- 'coifN' — Coiflets wavelet with N vanishing moments, where N is a positive integer from 1 to 5.
- 'fkN' — Fejér-Korovkin wavelet with N coefficients, where N = 4, 6, 8, 14, 18 and 22.

The synthesis wavelet must be the same wavelet used in the analysis with `modwt`.

Lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. You can specify `Lo` only if you do not specify `wname`. `Lo` must be the same scaling filter used in the analysis with `modwt`.

Hi — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. You can specify `Hi` only if you do not specify `wname`. `Hi` must be the same wavelet filter used in the analysis with `modwt`.

lev — Reconstruction level

0 (default) | nonnegative integer

Reconstruction level, specified as a nonnegative integer between 0 and `size(w,1)-2`. The level must be less than the level used to obtain `w` from `modwt`. If `lev` is 0 and you do not modify the coefficients, `imodwt` produces a perfect reconstruction of the signal.

Output Arguments

xrec — Reconstructed signal

row vector

Reconstructed version of the original signal based on the MODWT and the level of reconstruction, returned as a row vector.

References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`modwt` | `modwtmra`

Introduced in R2015b

ind2depo

Node index to node depth-position

Syntax

$[D, P] = \text{ind2depo}(\text{ORD}, [D \ P])$

Description

ind2depo is a tree-management utility.

For a tree of order ORD, $[D, P] = \text{ind2depo}(\text{ORD}, N)$ computes the depths D and the positions P (at these depths D) for the nodes with indices N.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

N must be a column vector of integers ($N \geq 0$).

Note that $[D, P] = \text{ind2depo}(\text{ORD}, [D \ P])$.

Examples

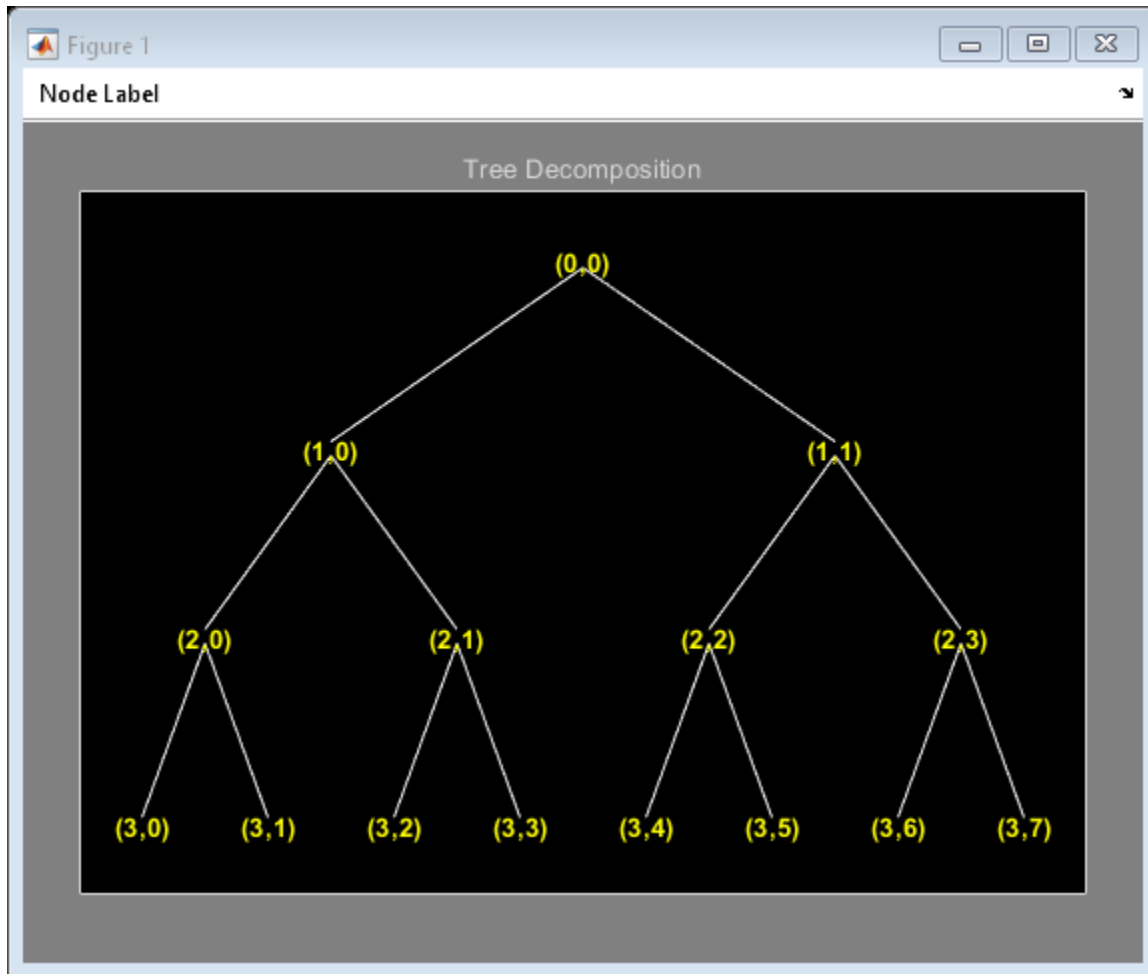
Depth and Position in Wavelet Packet Tree

Create a binary wavelet packet tree with three levels.

```
Ord = 2;  
Lev = 3;  
T = ntree(Ord, Lev);
```

Plot the binary wavelet packet tree.

```
plot(T)
```



Obtain the indices of the nodes in linear order.

```
idx = allnodes(T);
```

Convert the indices to depth-position format.

```
[depth,pos] = ind2depo(Ord,idx);  
table(depth,pos)
```



```
ans=15x2 table
  depth    pos
  -----
      0      0
      1      0
      1      1
      2      0
      2      1
      2      2
      2      3
      3      0
      3      1
      3      2
      3      3
      3      4
      3      5
      3      6
      3      7
```

See Also

depo2ind

Introduced before R2006a

intwave

Integrate wavelet function ψ (ψ)

Syntax

```
[INTEG,XVAL] = intwave('wname',PREC)
[INTDEC,XVAL,INTREC] = intwave('wname',PREC)
[INTEG,XVAL] = intwave('wname',PREC)
[INTEG,XVAL] = intwave('wname',PREC,0)
[INTEG,XVAL] = intwave('wname')
[INTEG,XVAL] = intwave('wname',8)
intwave('wname',IN2,IN3), PREC = max(IN2,IN3)
intwave('wname',0)
intwave('wname',8,IN3)
intwave('wname')
intwave('wname',8)
```

Description

[INTEG,XVAL] = intwave('wname',PREC) computes the integral, INTEG, of the wavelet function ψ (from $-\infty$ to XVAL values): $\int_{-\infty}^x \psi(y)dy$ for x in XVAL.

The function ψ is approximated on the 2^{PREC} points grid XVAL where PREC is a positive integer. 'wname' is a character vector containing the name of the wavelet ψ (see wfilters for more information).

Output argument INTEG is a real or complex vector depending on the wavelet type.

For biorthogonal wavelets,

[INTDEC,XVAL,INTREC] = intwave('wname',PREC) computes the integrals, INTDEC and INTREC, of the wavelet decomposition function ψ_{dec} and the wavelet reconstruction function ψ_{rec} .

`[INTEG,XVAL] = intwave('wname',PREC)` is equivalent to `[INTEG,XVAL] = intwave('wname',PREC,0)`.

`[INTEG,XVAL] = intwave('wname')` is equivalent to `[INTEG,XVAL] = intwave('wname',8)`.

When used with three arguments `intwave('wname',IN2,IN3)`, `PREC = max(IN2,IN3)` and plots are given.

When `IN2` is equal to the special value 0, `intwave('wname',0)` is equivalent to `intwave('wname',8,IN3)`.

`intwave('wname')` is equivalent to `intwave('wname',8)`.

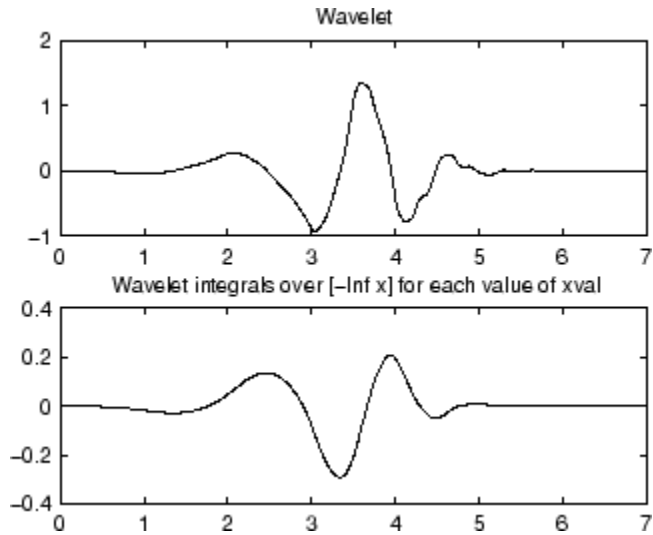
`intwave` is used only for continuous analysis (see `cwt` for more information).

Examples

```
% Set wavelet name.
wname = 'db4';

% Plot wavelet function.
[phi,psi,xval] = wavefun(wname,7);
subplot(211); plot(xval,psi); title('Wavelet');

% Compute and plot wavelet integrals approximations
% on a dyadic grid.
[integ,xval] = intwave(wname,7);
subplot(212); plot(xval,integ);
title(['Wavelet integrals over [-Inf x] ' ...
      'for each value of xval']);
```



Algorithms

First, the wavelet function is approximated on a grid of 2^{PREC} points using `wavefun`. A piecewise constant interpolation is used to compute the integrals using `cumsum`.

See Also

`wavefun`

Introduced before R2006a

isBiorthogonal

Determine if DWT filter bank is biorthogonal

Syntax

```
tf = isBiorthogonal(fb)
tf = isBiorthogonal(fb,tol)
```

Description

`tf = isBiorthogonal(fb)` returns `true` if the DWT filter bank `fb` is a biorthogonal filter bank and `false` otherwise.

Use `isOrthogonal` to determine whether a filter bank is orthogonal.

`tf = isBiorthogonal(fb,tol)` uses the positive real-valued tolerance, `tol`, to determine the biorthogonality of the filter bank `fb`. `tol` is a small positive number in the interval $(0, 10^{-2}]$. If unspecified, `tol` defaults to 10^{-5} .

Examples

Biorthogonality Test of DWT Filter Bank

Check whether a filter bank is biorthogonal.

```
fb = dwtfilterbank('Wavelet','bior4.4');
isBiorthogonal(fb)
```

```
ans = logical
      1
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

tol — Tolerance

10^{-2} (default) | positive scalar

Tolerance to use to determine biorthogonality of the filter bank, specified as a positive scalar in the interval $(0, 10^{-2}]$.

See Also

`dwtfilterbank`

Introduced in R2018a

isnode

Existing node test

Syntax

```
R = isnode(T,N)
```

Description

`isnode` is a tree-management utility.

`R = isnode(T,N)` returns 1's for nodes N , which exist in the tree T , and 0's for others.

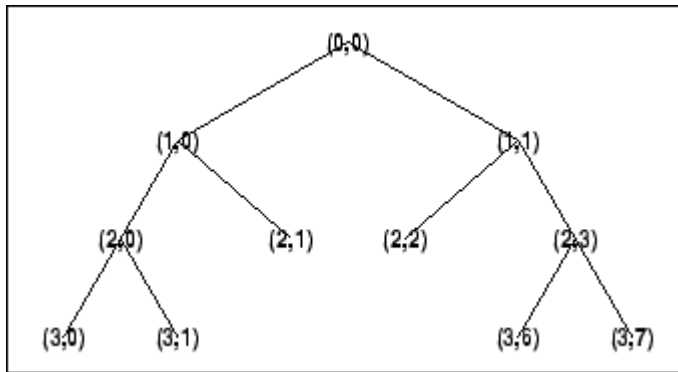
N can be a column vector containing the indices of nodes or a matrix, that contains the depths and positions of nodes.

In the last case, $N(i,1)$ is the depth of the i -th node and $N(i,2)$ is the position of the i -th node.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

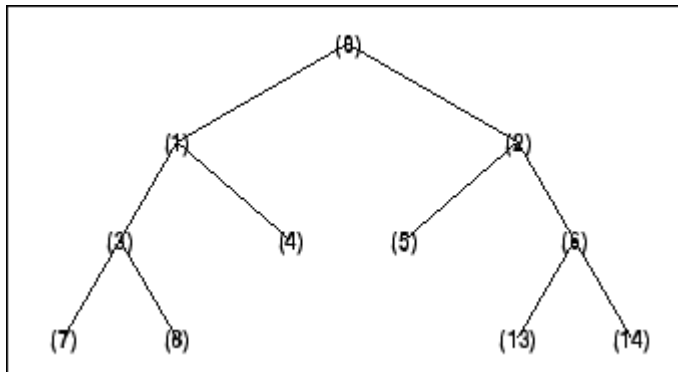
```
% Create initial tree.  
ord = 2;  
t = ntree(ord,3);    % binary tree of depth 3.  
t = nodejoin(t,5);  
t = nodejoin(t,4);  
plot(t)
```



```

% Change Node Label from Depth_Position to Index
% (see the plot function).

```



```

% Check node index.
isnode(t,[1;3;25])

```

```

ans =
     1
     1
     0

```

```

% Check node Depth_Position.
isnode(t,[1 0;3 1;4 5])

```

```

ans =
     1

```


1
0

See Also

istnode | wtreemgr

Introduced before R2006a

isOrthogonal

Determine if DWT filter bank is orthogonal

Syntax

```
tf = isOrthogonal(fb)
tf = isOrthogonal(fb,tol)
```

Description

`tf = isOrthogonal(fb)` returns `true` if the DWT filter bank `fb` is an orthogonal filter bank and `false` otherwise.

Use `isBiorthogonal` to determine whether a filter bank is biorthogonal.

`tf = isOrthogonal(fb,tol)` uses the positive real-valued tolerance, `tol`, to determine the orthogonality of the filter bank `fb`. `tol` is a small positive number in the interval $(0, 10^{-2}]$. If unspecified, `tol` defaults to 10^{-5} .

Examples

Orthogonality Test of DWT Filter Bank

Create a DWT filter bank using the Daubechies `db6` wavelet. Confirm the filter bank is orthogonal.

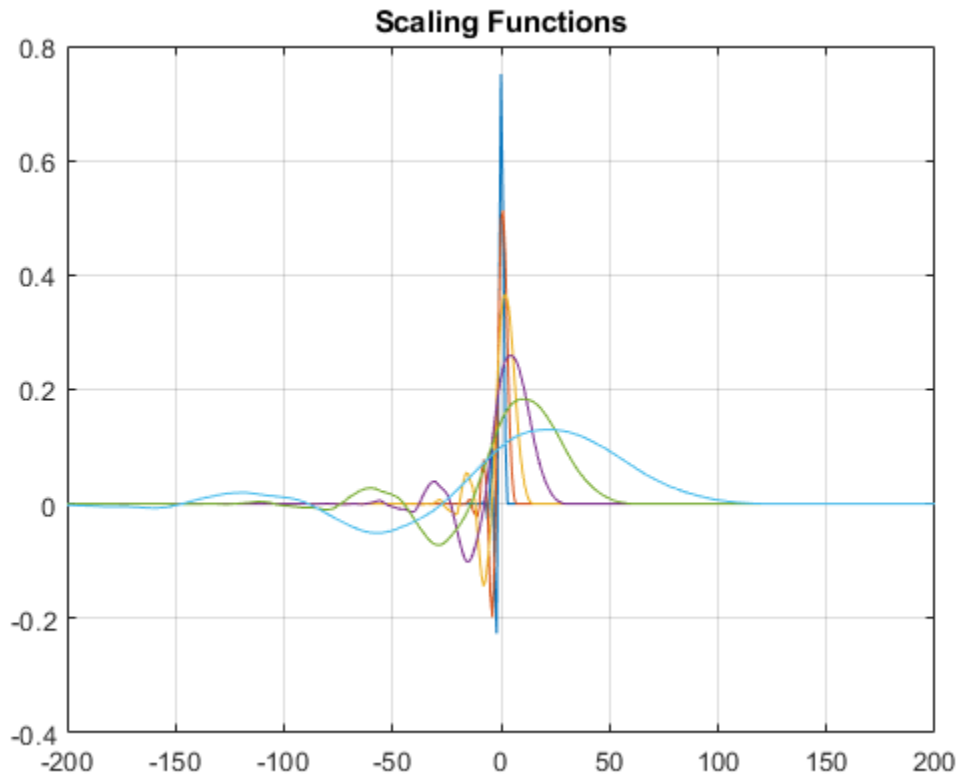
```
fb = dwtfilterbank('Wavelet','db6');
isOrthogonal(fb)
```

```
ans = logical
      1
```

Obtain the time-domain and centered scaling functions for each level in the filter bank. Obtain the time-domain and centered wavelets corresponding to the wavelet passband

filters. Plot the scaling functions and wavelets. Confirm the scaling functions and wavelets all have norm square equal to 1.

```
[phi,t] = scalingfunctions(fb);  
psi = wavelets(fb);  
figure  
plot(t,phi')  
grid on  
xlim([-200 200])  
title('Scaling Functions')
```

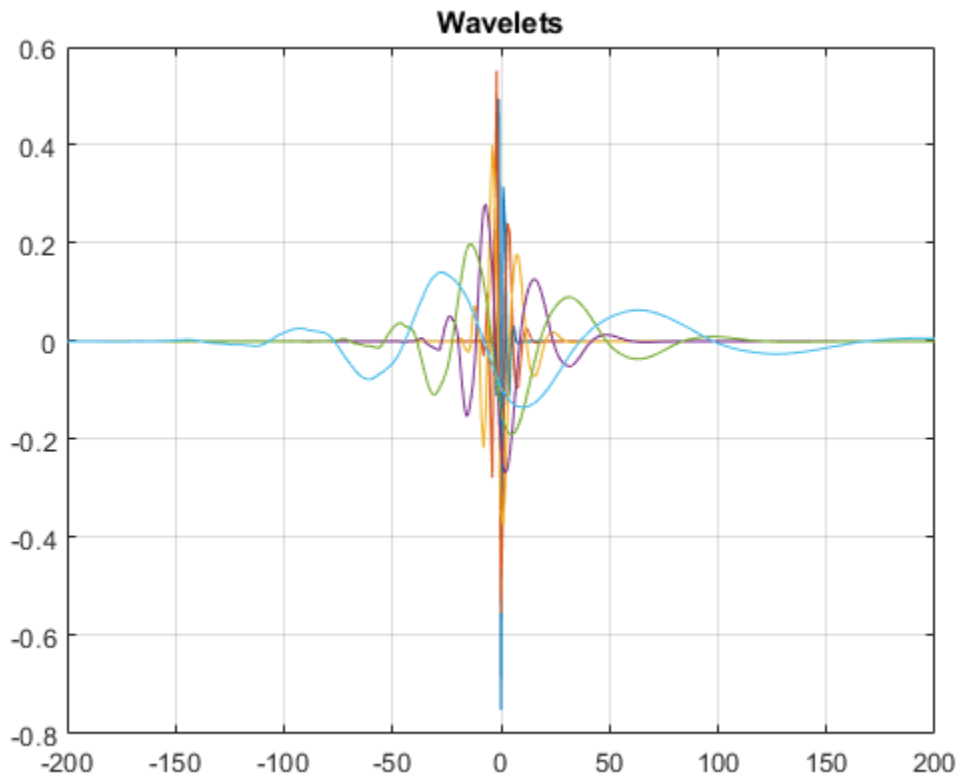


```
sum(phi.^2,2)
```

```
ans = 6×1
```

```
1.0000  
1.0000  
1.0000  
1.0000  
1.0000  
1.0000
```

```
figure  
plot(t,psi')  
grid on  
xlim([-200 200])  
title('Wavelets')
```



```
sum(psi.^2,2)
```

```
ans = 6×1
```

```
1.0000  
1.0000  
1.0000  
1.0000  
1.0000  
1.0000
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

tol — Tolerance

10^{-2} (default) | positive scalar

Tolerance to use to determine orthogonality of the filter bank, specified as a positive scalar in the interval $(0, 10^{-2}]$.

See Also

`dwtfilterbank`

Introduced in R2018a

istnode

Terminal nodes indices test

Syntax

```
R = istnode(T,N)
```

Description

`istnode` is a tree-management utility.

`R = istnode(T,N)` returns ranks (in left to right terminal nodes ordering) for terminal nodes N belonging to the tree T , and 0's for others.

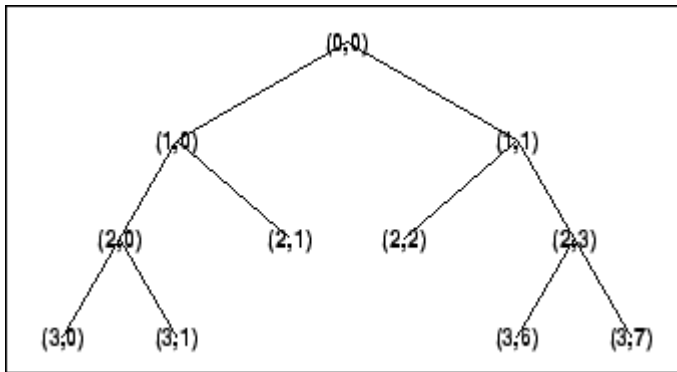
N can be a column vector containing the indices of nodes or a matrix that contains the depths and positions of nodes.

In the last case, $N(i,1)$ is the depth of the i -th node and $N(i,2)$ is the position of the i -th node.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

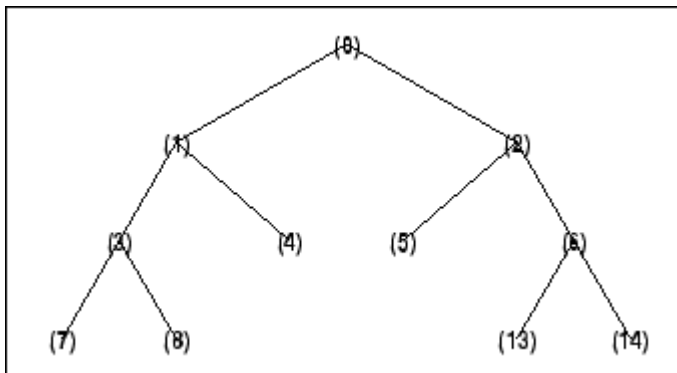
```
% Create initial tree.  
ord = 2;  
t = ntree(ord,3); % binary tree of depth 3.  
t = nodejoin(t,5);  
t = nodejoin(t,4);  
plot(t)
```



```

% Change Node Label from Depth_Position to Index
% (see the plot function)x.

```



```

% Find terminal nodes and return indices for terminal
% nodes in the tree.

```

```
istnode(t,[14])
```

```
ans =
     6
```

```
istnode(t,[15])
```

```
ans =
     0
```

```
istnode(t,[1;7;14;25])
```

```
ans =
     0
     1
```

```
      6
      0
istnode(t,[1 0;3 1;4 5])
ans =
      0
      2
      0
```

See Also

isnode | wtreemgr

Introduced before R2006a

iswt

Inverse discrete stationary wavelet transform 1-D

Syntax

```
X = iswt(SWC, 'wname')
X = iswt(SWA, SWD, 'wname')
X = iswt(SWA(end, :), SWD, 'wname')
X = iswt(SWC, Lo_R, Hi_R)
X = iswt(SWA, SWD, Lo_R, Hi_R)
X = iswt(SWA(end, :), SWD, Lo_R, Hi_R)
```

Description

`iswt` performs a multilevel 1-D stationary wavelet reconstruction using either an orthogonal or a biorthogonal wavelet. Specify the wavelet using its name (`'wname'`, see `wfilters` for more information) or its reconstruction filters (`Lo_R` and `Hi_R`).

`X = iswt(SWC, 'wname')` or `X = iswt(SWA, SWD, 'wname')` or `X = iswt(SWA(end, :), SWD, 'wname')` reconstructs the signal `X` based on the multilevel stationary wavelet decomposition structure `SWC` or `[SWA, SWD]` (see `swt` for more information).

`X = iswt(SWC, Lo_R, Hi_R)` or `X = iswt(SWA, SWD, Lo_R, Hi_R)` or `X = iswt(SWA(end, :), SWD, Lo_R, Hi_R)` reconstruct as above, using filters that you specify.

- `Lo_R` is the reconstruction low-pass filter.
- `Hi_R` is the reconstruction high-pass filter.

`Lo_R` and `Hi_R` must be the same length.

Examples

Multilevel Stationary Wavelet Reconstruction

Demonstrate perfect reconstruction using `swt` and `iswt` with a biorthogonal wavelet.

```
load noisbloc
[Lo_D,Hi_D,Lo_R,Hi_R] = wfilters('bior3.5');
[swa,swd] = swt(noisbloc,3,Lo_D,Hi_D);
recon = iswt(swa,swd,Lo_R,Hi_R);
norm(noisbloc-recon)
```

```
ans = 1.1386e-13
```

References

Nason, G.P.; B.W. Silverman (1995), "The stationary wavelet transform and some statistical applications," *Lecture Notes in Statistics*, 103, pp. 281-299.

Coifman, R.R.; Donoho D.L. (1995), "Translation invariant de-noising," *Lecture Notes in Statistics*, 103, pp 125-150.

Pesquet, J.C.; H. Krim, H. Carfatan (1996), "Time-invariant orthonormal wavelet representations," *IEEE Trans. Sign. Proc.*, vol. 44, 8, pp. 1964-1970.

See Also

`idwt` | `swt` | `waverec`

Introduced before R2006a

iswt2

Inverse discrete stationary wavelet transform 2-D

Syntax

```
X = iswt2(SWC, 'wname')
X = iswt2(A, H, V, D, wname)
X = iswt2(A(:, :, end), H, V, D, 'wname')
X = iswt2(A(:, :, 1, :), H, V, D, 'wname')
X = iswt2(SWC, Lo_R, Hi_R)
X = iswt2(A, H, V, D, Lo_R, Hi_R)
X = iswt2(A(:, :, end), H, V, D, Lo_R, Hi_R)
X = iswt2(A(:, :, 1, :), H, V, D, 'wname')
```

Description

`iswt2` performs a multilevel 2-D stationary wavelet reconstruction using either an orthogonal or a biorthogonal wavelet. Specify the wavelet using its name (`'wname'`, see `wfilters` for more information) or its reconstruction filters (`Lo_R` and `Hi_R`).

`X = iswt2(SWC, 'wname')` or `X = iswt2(A, H, V, D, wname)` reconstructs the signal `X`, based on the multilevel stationary wavelet decomposition structure `SWC` or `[A, H, V, D]` (see `swt2`).

If multilevel stationary wavelet decomposition structure `SWC` or `[A, H, V, D]` was generated from a 2-D matrix, the syntax `X = iswt2(A(:, :, end), H, V, D, 'wname')` reconstructs the signal `X`.

If the stationary wavelet decomposition structure `SWC` or `[A, H, V, D]` was generated from a single level stationary wavelet decomposition of a 3-D matrix, `X = iswt2(A(:, :, 1, :), H, V, D, 'wname')` reconstructs the signal `X`.

`X = iswt2(SWC, Lo_R, Hi_R)` or `X = iswt2(A, H, V, D, Lo_R, Hi_R)` or `X = iswt2(A(:, :, end), H, V, D, Lo_R, Hi_R)` or `X = iswt2(A(:, :, 1, :), H, V, D, 'wname')` reconstructs as in the previous syntax, using filters that you specify:

- Lo_R is the reconstruction low-pass filter.
- Hi_R is the reconstruction high-pass filter.

Lo_R and Hi_R must be the same length.

Note

- `iswt2` synthesizes X from the coefficient arrays generated by `swt2`. `swt2` uses double-precision arithmetic internally and returns double-precision coefficient matrices. `swt2` warns if there is a loss of precision when converting to double.
- To distinguish a single-level decomposition of a truecolor image from a multilevel decomposition of an indexed image, the approximation and detail coefficient arrays of truecolor images are 4-D arrays. See “Distinguish Single-Level Truecolor Image from Multilevel Indexed Image Decompositions” on page 1-663. Also see examples “Stationary Wavelet Transform of an Image” on page 1-655 and “Inverse Stationary Wavelet Transform of an Image” on page 1-659.

If an K -level decomposition is performed, the dimensions of the A , H , V , and D coefficient arrays are m -by- n -by-3-by- K .

If a single-level decomposition is performed, the dimensions of the A , H , V , and D coefficient arrays are m -by- n -by-1-by-3. Since MATLAB removes singleton last dimensions by default, the third dimension of the coefficient arrays is singleton.

Examples

Multilevel Two-Dimensional Stationary Wavelet Reconstruction

Demonstrate perfect reconstruction using `swt2` and `iswt2` with an orthogonal wavelet.

```
load woman
[Lo_D,Hi_D,Lo_R,Hi_R] = wfilters('db6');
[ca,chd,cvd,cdd] = swt2(X,3,Lo_D,Hi_D);
recon = iswt2(ca,chd,cvd,cdd,Lo_R,Hi_R);
norm(X-recon)
```

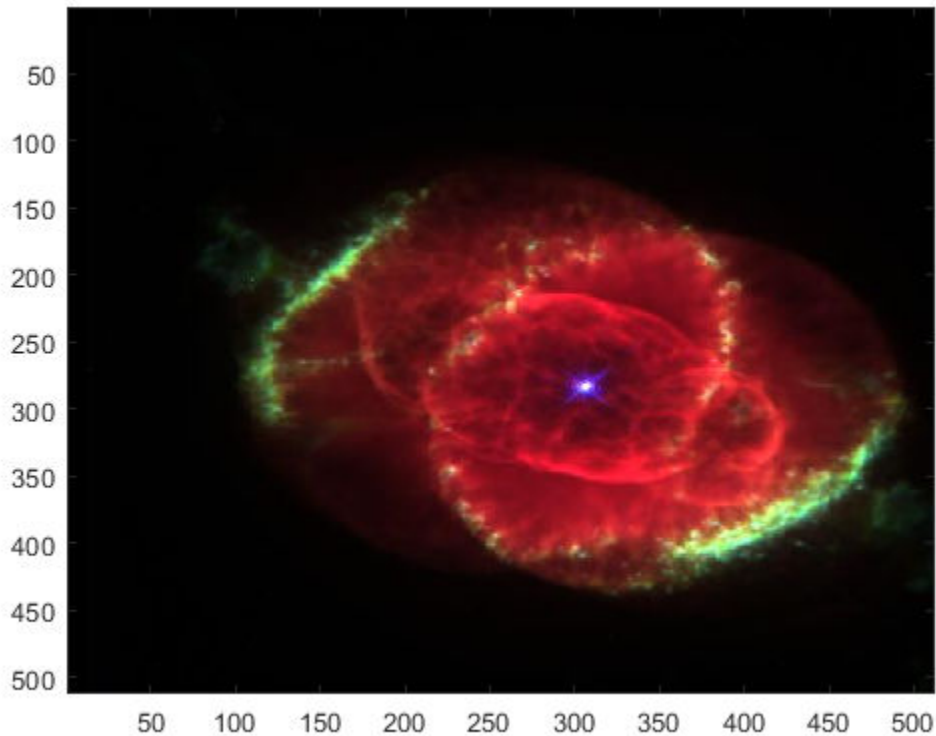
```
ans = 1.0126e-08
```

Stationary Wavelet Transform of an Image

In this example you obtain single-level and multilevel stationary wavelet decompositions of a truecolor image. You view results of each decomposition.

Load in a truecolor image. The image is a 3-D array of type `uint8`. Since `swt2` requires the first and second dimensions both be divisible by a power of 2, extract a portion of the image and view it.

```
imdata = imread('ngc6543a.jpg');  
x = imdata(1:512,1:512,:);  
imagesc(x)
```



Obtain the 4-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and the horizontal, vertical, and detail coefficients as separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,4,'db4');  
size(a)
```

```
ans = 1×4
```

```
512 512 3 4
```

```
size(h)
```

```
ans = 1×4
```

```
512 512 3 4
```

```
size(v)
```

```
ans = 1×4
```

```
512 512 3 4
```

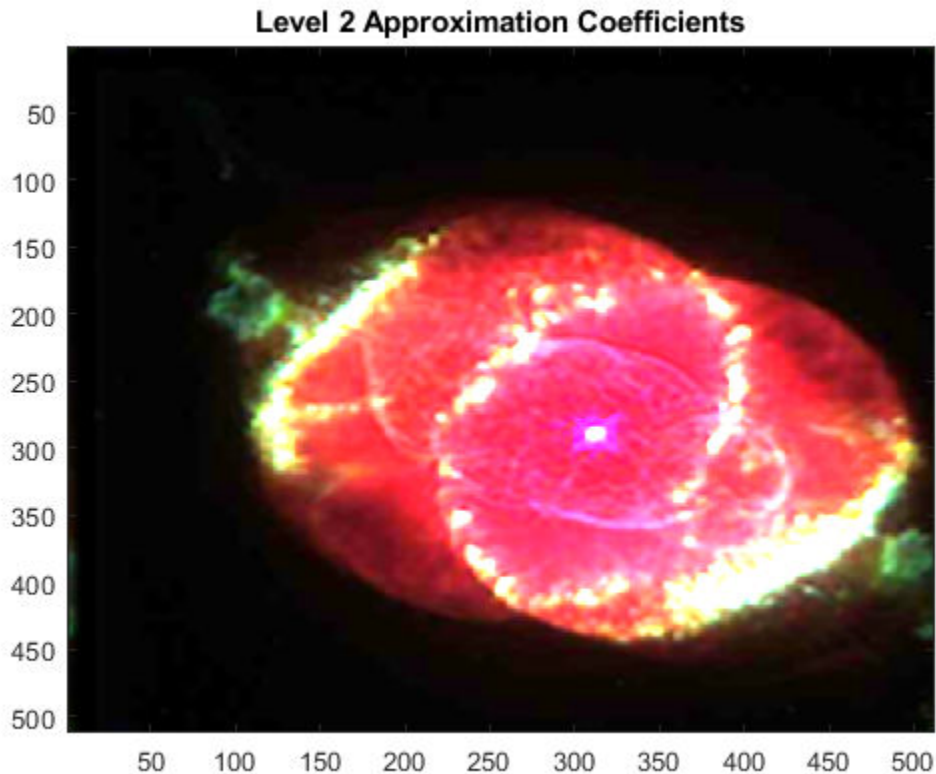
```
size(d)
```

```
ans = 1×4
```

```
512 512 3 4
```

The output arrays are all of type `double`. View the level 2 approximation coefficients. Since the approximation coefficients are of type `double`, cast them as `uint8`, which is the datatype of the image.

```
figure  
imagesc(uint8(a(:,:,2)))  
title('Level 2 Approximation Coefficients')
```



Reconstruct the image by performing the inverse transform. Compute the difference between the original image and reconstruction. Since the reconstruction is of type double, cast the reconstruction as type uint8 before computing the difference.

```
rec = iswt2(a,h,v,d,'db4');
maxdiff = max(abs(uint8(rec(:))-x(:)));
disp(['maximum difference = ' num2str(maxdiff)])
```

```
maximum difference = 0
```

Obtain the single-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and horizontal, vertical, and detail coefficients in separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,1, 'db4');  
size(a)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(h)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(v)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(d)
```

```
ans = 1×4
```

```
512 512 1 3
```

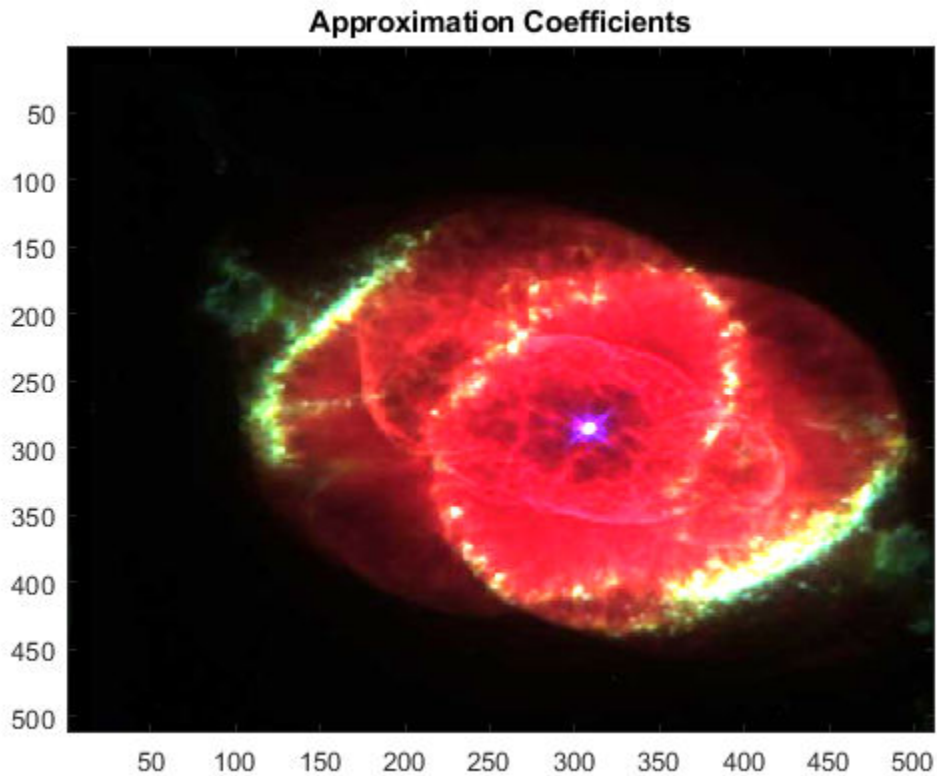
View the approximation coefficients. To prevent an error when using `imagesc`, squeeze the approximation coefficients array to remove the singleton dimension.

```
asqueeze = squeeze(a);  
size(asqueeze)
```

```
ans = 1×3
```

```
512 512 3
```

```
figure  
imagesc(uint8(asqueeze))  
title('Approximation Coefficients')
```

Reconstruct the image by performing the inverse transform. Compute the difference between the original image and reconstruction. Since the reconstruction is of type double, cast the reconstruction as type uint8 before computing the difference.

```
rec = iswt2(a,h,v,d,'db4');  
maxdiff = max(abs(uint8(rec(:))-x(:)));  
disp(['maximum difference = ' num2str(maxdiff)])
```

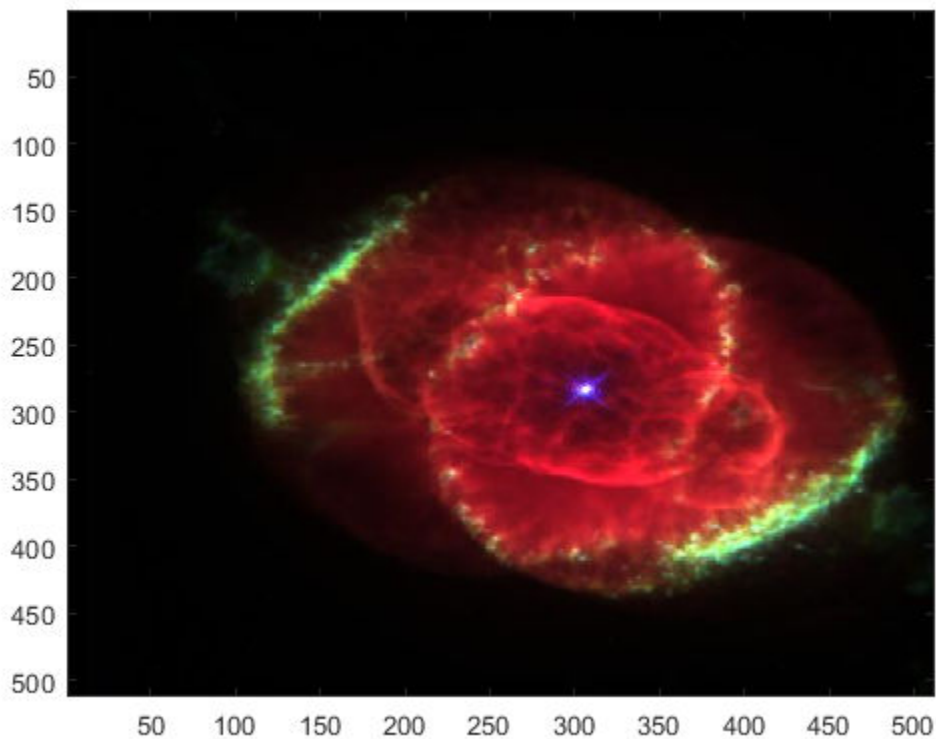
```
maximum difference = 0
```

Inverse Stationary Wavelet Transform of an Image

This example shows how to reconstruct a truecolor image from a single-level stationary wavelet decomposition using 3-D approximation and detail coefficient arrays.

Load in a truecolor image. The image is a 3-D array of type `uint8`. Since `swt2` requires the first and second dimensions both be divisible by a power of 2, extract a portion of the image and view it.

```
imdata = imread('ngc6543a.jpg');  
x = imdata(1:512,1:512,:);  
imagesc(x)
```



Obtain the single-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and horizontal, vertical, and detail coefficients in separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,1,'db4');  
size(a)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(h)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(v)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(d)
```

```
ans = 1×4
```

```
512 512 1 3
```

Squeeze the coefficient arrays to remove their singleton dimensions. Note the dimensions of the squeezed arrays.

```
asq = squeeze(a);  
hsq = squeeze(h);  
vsq = squeeze(v);  
dsq = squeeze(d);  
size(asq)
```

```
ans = 1×3
```

```
512 512 3
```

```
size(hsq)
ans = 1×3
    512    512     3
```

```
size(vsq)
ans = 1×3
    512    512     3
```

```
size(dsq)
ans = 1×3
    512    512     3
```

So that `iswt2` can properly reconstruct the true image from the new coefficient arrays, insert a singleton dimension by reshaping the squeezed arrays. Reconstruct the image with the reshaped coefficient arrays.

```
a2 = reshape(asq,[512,512,1,3]);
h2 = reshape(hsq,[512,512,1,3]);
v2 = reshape(vsq,[512,512,1,3]);
d2 = reshape(dsq,[512,512,1,3]);
rec = iswt2(a2,h2,v2,d2,'db4');
```

Compute the difference between the original image and reconstruction. Since the reconstruction is of type `double`, cast the reconstruction as type `uint8` before computing the difference.

```
maxdiff = max(abs(uint8(rec(:))-x(:)));
disp(['maximum difference = ' num2str(maxdiff)])

maximum difference = 0
```

Tips

If SWC or (cA,cH,cV,cD) are obtained from an indexed image analysis or a truecolor image analysis, then X is an m-by-n matrix or an m-by-n-by-3 array, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Compatibility Considerations

Distinguish Single-Level Truecolor Image from Multilevel Indexed Image Decompositions

Behavior changed in R2017b

To distinguish a single-level decomposition of a truecolor image from a multilevel decomposition of an indexed image, the approximation and detail coefficient arrays of truecolor images are 4-D arrays.

Migrate from Previous Releases to R2017b

Depending on the original input data type and level of wavelet decomposition, you might have to take different steps to make `swt2` coefficient arrays from previous releases compatible with R2017b coefficient arrays. The steps depend on whether you have a single coefficient array or separate approximation and detail coefficient arrays.

Single Coefficient Array	Multiple Coefficient Arrays
Input: Index image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues 	Input: Index image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues

Single Coefficient Array	Multiple Coefficient Arrays
<p>Input: Truecolor image</p> <ul style="list-style-type: none"> Single-level: If <code>swc</code> is the output of <code>swt2</code> from a previous release, execute: <code>swc1 = double(swc);</code> Multi-level: If <code>swc</code> is the output of <code>swt2</code> from a previous release, execute: <code>swc1 = double(swc);</code> 	<p>Input: Truecolor image</p> <ul style="list-style-type: none"> Single-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from a previous release, execute: <code>ca1 = double(ca);</code> <code>chd1 = double(chd);</code> <code>cvd1 = double(cvd);</code> <code>cdd1 = double(cdd);</code> <code>ca2 = reshape(ca1, [m,n,1,3]);</code> <code>chd2 = reshape(chd1, [m,n,1,3]);</code> <code>cvd2 = reshape(cvd1, [m,n,1,3]);</code> <code>cdd2 = reshape(cdd1, [m,n,1,3]);</code> Multi-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from a previous release, execute: <code>ca1 = double(ca);</code> <code>chd1 = double(chd);</code> <code>cvd1 = double(cvd);</code> <code>cdd1 = double(cdd);</code>

Migrate from R2017b to Previous Releases

Depending on the original input data type and level of wavelet decomposition, you might have to take different steps to make R2017b `swt2` coefficient arrays compatible with the coefficient arrays from previous releases. The steps depend on whether you have a single coefficient array or separate approximation and detail coefficient arrays.

Single Coefficient Array	Multiple Coefficient Arrays
<p>Input: Index image</p> <ul style="list-style-type: none"> Single-level: No compatibility issues Multi-level: No compatibility issues 	<p>Input: Index image</p> <ul style="list-style-type: none"> Single-level: No compatibility issues Multi-level: No compatibility issues

Single Coefficient Array	Multiple Coefficient Arrays
Input: Truecolor image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues 	Input: Truecolor image <ul style="list-style-type: none"> • Single-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from R2017b, execute: <pre>ca1 = single(squeeze(ca)); chd1 = single(squeeze(chd)); cvd1 = single(squeeze(cvd)); cdd1 = single(squeeze(cdd));</pre> • Multi-level: No compatibility issues

References

Nason, G.P.; B.W. Silverman (1995), "The stationary wavelet transform and some statistical applications," *Lecture Notes in Statistics*, 103, pp. 281-299.

Coifman, R.R.; Donoho D.L. (1995), "Translation invariant de-noising," *Lecture Notes in Statistics*, 103, pp. 125-150.

Pesquet, J.C.; H. Krim, H. Carfatan (1996), "Time-invariant orthonormal wavelet representations," *IEEE Trans. Sign. Proc.*, vol. 44, 8, pp. 1964-1970.

See Also

`idwt2` | `swt2` | `waverec2`

Introduced before R2006a

iwssst

Inverse wavelet synchrosqueezed transform

Syntax

```
xrec = iwssst(sst)
xrec = iwssst(sst,f,freqrange)
xrec = iwssst(sst,iridge)
xrec = iwssst( ____,wav)
xrec = iwssst( ____,iridge,'NumFrequencyBins',numBins)
```

Description

`xrec = iwssst(sst)` inverts the input synchrosqueezed transform, `sst`, and returns the inverse in vector `xrec`. To obtain the `sst` input, use the `wssst` function. The `iwssst` function assumes that you obtain `sst` using the analytic Morlet wavelet.

Note The wavelet transform does not preserve a nonzero mean. After inverting the synchrosqueezed transform, you must add back the original signal mean.

`xrec = iwssst(sst,f,freqrange)` inverts the synchrosqueezed transform for a specified range of frequencies, `freqrange`, contained in the frequency vector, `f`. The frequency vector, `f`, is the output of `wssst`.

`xrec = iwssst(sst,iridge)` inverts the synchrosqueezed transform along the time-frequency ridges specified by `iridge`, the index column vector. `iridge` is the output of `wssstridge`. The `xrec` output is the same size as `iridge`.

`xrec = iwssst(____,wav)` uses the analytic wavelet specified by `wav` to invert the synchrosqueezed transform. This wavelet must be the same wavelet as used in `wssst`. You can include any of the input arguments from previous syntaxes.

`xrec = iwssst(____,iridge,'NumFrequencyBins',numBins)` returns the inverse synchrosqueezed transform with `numBins`-many additional frequency bins included on either side of each `iridge` index bin.

Examples

Inverse Synchrosqueezed Transform of Chirp

Obtain the wavelet synchrosqueezed transform of a quadratic chirp using default values. Then reconstruct the signal using the inverse wavelet synchrosqueezed transform.

```
load quadchirp;
sst = wsst(quadchirp);
xrec = iwsst(sst);
```

Synchrosqueezed and Inverse Synchrosqueezed Transform of Chirp

Obtain the wavelet synchrosqueezed transform of a quadratic chirp sampled at 1000 Hz. Then reconstruct the chirp.

Load the chirp and obtain the synchrosqueezed transform.

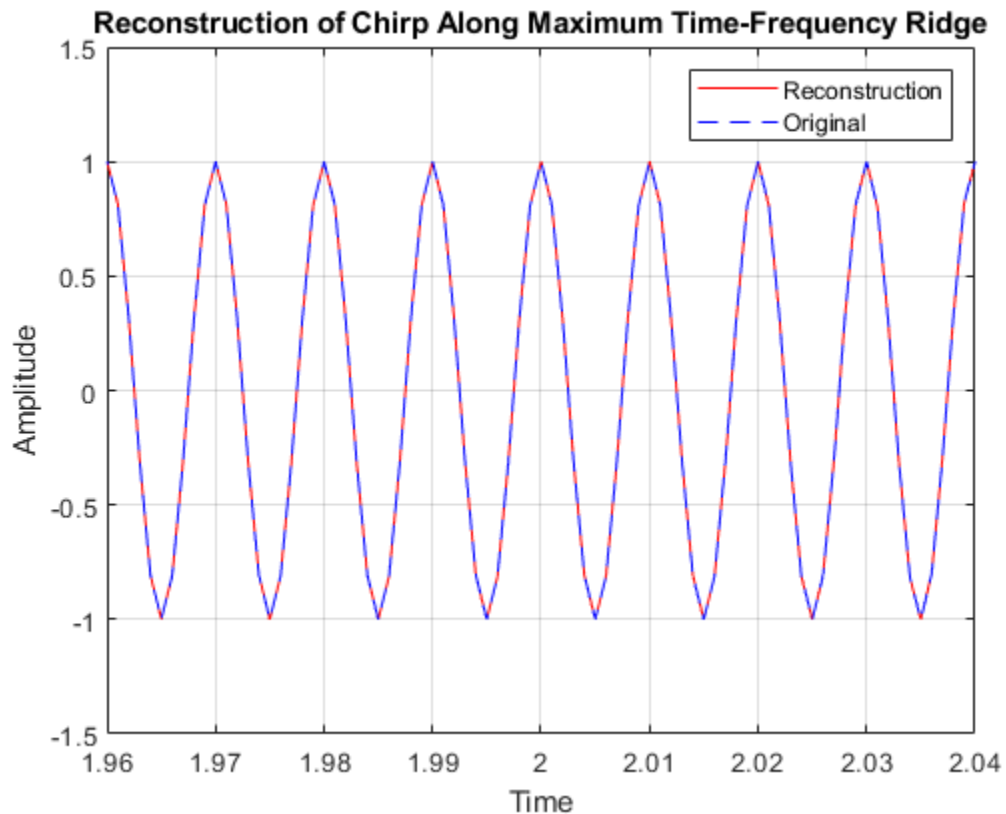
```
load quadchirp;
sstchirp = wsst(quadchirp, 'ExtendSignal', true);
```

Extract the maximum energy time-frequency ridge and reconstruct the signal mode along the ridge.

```
[~,iridge] = wsstridge(sstchirp);
xrec = iwsst(sstchirp,iridge);
```

Plot and zoom in on the original and reconstructed signal.

```
plot(tquad,xrec,'r');
hold on;
plot(tquad,quadchirp,'b--');
xlabel('Time'); ylabel('Amplitude');
set(gca,'ylim',[-1.5 1.5]);
legend('Reconstruction','Original');
grid on;
title('Reconstruction of Chirp Along Maximum Time-Frequency Ridge');
zoom xon
zoom(50)
```



Inverse Synchrosqueezed Transform of Range of Frequencies

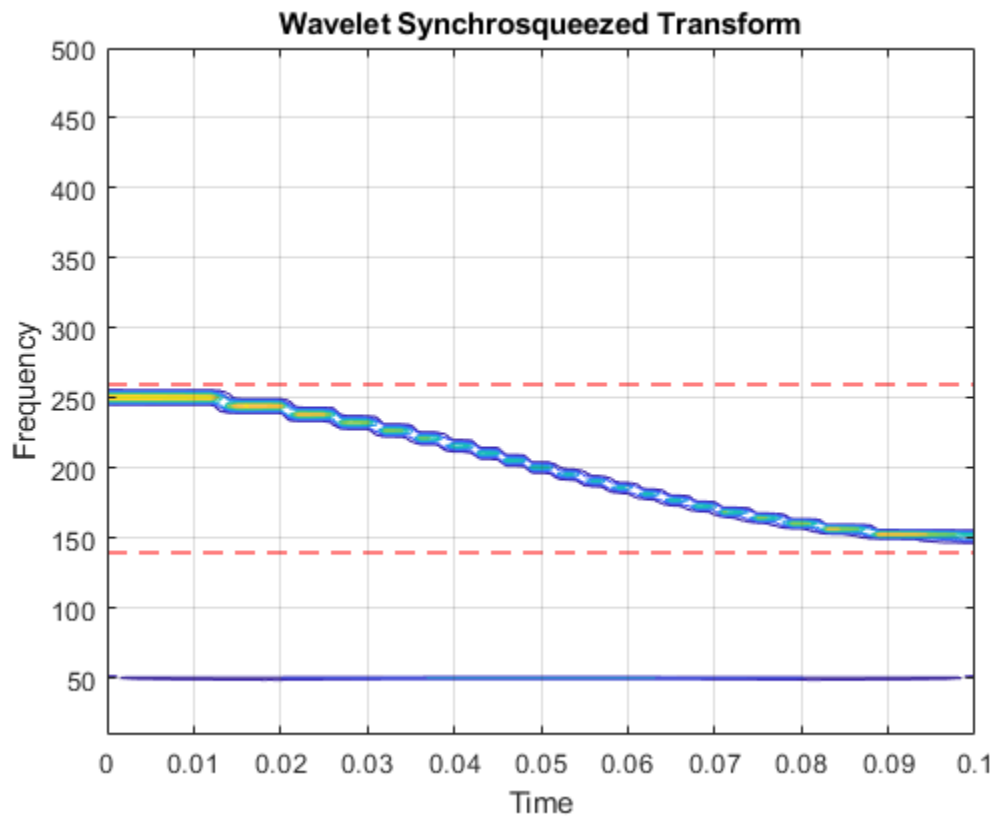
Obtain the inverse synchrosqueezed transform for a specified frequency range of a two-component signal. The input is a combination of an amplitude-modulated and a frequency-modulated signal.

Create the signal.

```
t = 0:0.001:0.1;  
x1 = (2+0.5*cos(2*pi*10*t)).*cos(2*pi*200*t+10*sin(2*pi*5*t));  
x2 = cos(2*pi*50*t);  
sig = x1+x2;
```

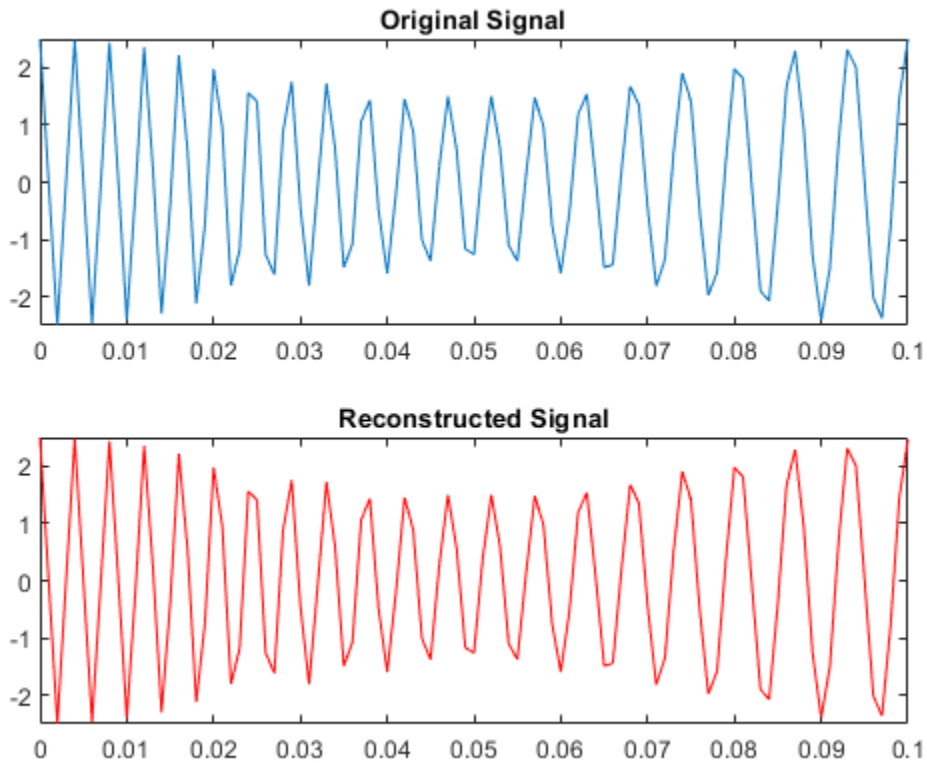
Obtain the wavelet synchrosqueezed transform and plot the resulting two frequency components.

```
[sst,f] = wssst(sig,1000,'ExtendSignal',true);  
contour(t,f,abs(sst));  
grid on;  
title('Wavelet Synchrosqueezed Transform');  
ylabel('Frequency');  
xlabel('Time');  
hold on  
plot(t,140*ones(size(t)),'r--');  
plot(t,260*ones(size(t)),'r--');
```



Obtain the inverse synchrosqueezed transform for frequencies from 140 Hz to 260 Hz. Plot the result.

```
xrec = iwsst(sst,f,[140,260]);  
subplot(2,1,1);  
plot(t,x1);  
title('Original Signal');  
subplot(2,1,2);  
plot(t,xrec,'r');  
title('Reconstructed Signal');
```



Synchrosqueezed and Inverse Synchrosqueezed Transform of Speech Signal

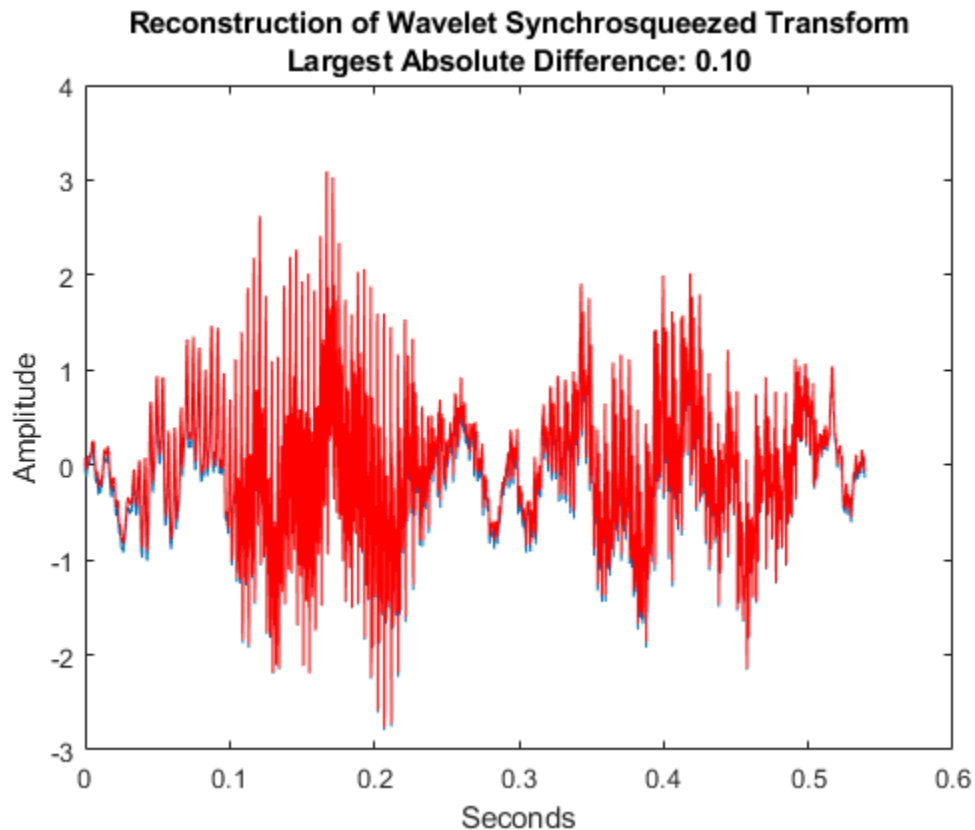
Obtain the wavelet synchrosqueezed transform and inverse synchrosqueezed transform of a speech sample using the bump wavelet.

Load the speech signal and obtain the synchrosqueezed transform and inverse synchrosqueezed transform.

```
load mtlb
dt = 1/Fs;
t = 0:dt:numel(mtlb)*dt-dt;
Txmtlb = wssst(mtlb,'bump');
xrec = iwsst(Txmtlb,'bump');
```

Obtain the L-infinity norm of the difference between the original waveform and the reconstruction. Plot the results.

```
Linf = norm(abs(mtlb-xrec),Inf);
plot(t,mtlb)
hold on
xlabel('Seconds')
ylabel('Amplitude')
plot(t,xrec,'r')
title({'Reconstruction of Wavelet Synchrosqueezed Transform';...
      ['Largest Absolute Difference: ' num2str(Linf, '%1.2f')]});
```



Synchrosqueezed Transform Using Specified Number of Bins for Chirp

This example shows how to invert the wavelet synchrosqueezed transform using a specified number of frequency bins for a quadratic chirp. The chirp is sampled at 1000 Hz.

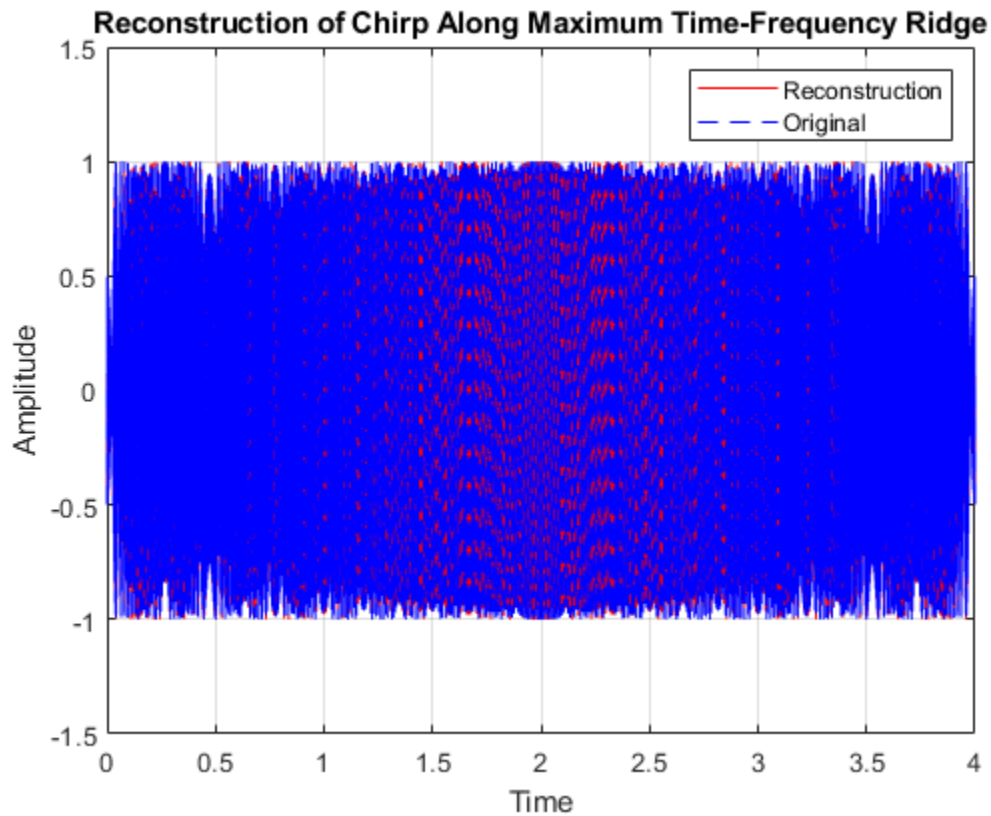
```
load quadchirp;  
sstchirp = wssst(quadchirp, 'ExtendSignal', true);
```

Extract the maximum energy time-frequency ridge using 10 bins on each side of the ridge index and reconstruct the signal mode along the ridge.

```
[~,iridge] = wsstridge(sstchirp);  
xrec = iwsst(sstchirp,iridge,'NumFrequencyBins',10);
```

Plot the original and reconstructed signal.

```
plot(tquad,xrec,'r');  
hold on;  
plot(tquad,quadchirp,'b--');  
xlabel('Time'); ylabel('Amplitude');  
set(gca,'ylim',[-1.5 1.5]);  
legend('Reconstruction','Original');  
grid on;  
title('Reconstruction of Chirp Along Maximum Time-Frequency Ridge');
```



Input Arguments

sst — Synchrosqueezed transform

matrix

Synchrosqueezed transform, specified as a matrix. `sst` is the output from the `wsst` function.

f — Synchrosqueezed transform frequencies

vector

Synchrosqueezed transform frequencies corresponding to the rows of the synchrosqueezed transform, specified as a vector. The number of elements in the frequency vector is equal to the number of rows in the `sst` input. If you specify `f`, you must also specify `freqrange`.

freqrange — Frequency range

two-element vector

Frequency range for which to return inverse synchrosqueezed transform values, specified as a two-element vector. The values of `freqrange` must be in the range of the values of the frequencies, `f`. The first and second elements of `freqrange` define the start and end of the frequency range, where the frequency values in that range must be positive and strictly increasing. If you specify `freqrange`, you must also specify `f`.

iridge — Time-frequency ridge row indices

vector or matrix

Time-frequency ridge row indices of the synchrosqueezed transform specified as a vector or matrix. `iridge` is the output of the `wsstridge` function. If `iridge` is a matrix, `iwsst` inverts the synchrosqueezed transform along the first column of `iridge`. Then, it iteratively reconstructs along subsequent columns of `iridge`. The sizes of `iridge` and the `xrec` output are the same.

wav — Analytic wavelet

'amor' (default) | 'bump'

Analytic wavelet used to compute the inverse synchrosqueezed transform, specified as one of the following:

- 'amor' — Analytic Morlet wavelet

- 'bump' — Bump wavelet

You must use the same wavelet in the reconstruction that you used to compute the synchrosqueezed transform, `sst`.

numBins — Number of additional frequency bins

16 (default) | positive integer

Number of additional frequency bins to include on either side of each `iridge` index bin, specified as a positive integer. If the number of additional bins exceeds the number of frequency bins available at a particular time step, `iwsst` truncates the reconstruction at the first or last frequency bin. The default, 16, is one half the default number of voices per octave.

To specify this argument, you also specify `iridge`, which is the output of `wsstridge`. You cannot include a frequency `f` and frequency range `freqrange`, if you include the number of frequency bins.

Output Arguments

xrec — Inverse synchrosqueezed transform

vector or matrix

Inverse synchrosqueezed transform, returned as a vector or matrix. If you do not specify an `iridge` input, `xrec` is a column vector with the same number of rows as `sst`. If you specify an `iridge` input, `xrec` is the same size as `iridge`.

References

- [1] Daubechies, I., J. Lu, and H. T. Wu. "Synchrosqueezed Wavelet Transforms: an Empirical Mode Decomposition-like Tool." *Applied and Computational Harmonic Analysis*, Vol. 30, Number 2, 2011, pp. 243-261.
- [2] Thakur, G., E. Brevdo, N. S. Fučkar, and H. T. Wu. "The Synchrosqueezing algorithm for time-varying spectral analysis: robustness properties and new paleoclimate applications." *Signal Processing*, Vol. 93, Number 5, 2013, pp. 1079-1094.

See Also

`wsst` | `wsstridge`

Topics

“Time-Frequency Reassignment and Mode Extraction with Synchrosqueezing”

“Wavelet Synchrosqueezing”

Introduced in R2016a

labelDefinitionsHierarchy

Get hierarchical list of label and sublabel names

Syntax

```
str = labelDefinitionsHierarchy(lbldefs)  
str = labelDefinitionsHierarchy(lss)
```

Description

`str = labelDefinitionsHierarchy(lbldefs)` returns a character array with a hierarchical list of label and sublabel names contained in `lbldefs`, a vector of `signalLabelDefinition` objects.

`str = labelDefinitionsHierarchy(lss)` returns a character array with a hierarchical list of label and sublabel names contained in the `labeledSignalSet` object `lss`.

Examples

Label Hierarchy

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss  
  
lss =  
  labeledSignalSet with properties:  
    Source: {2x1 cell}  
    NumMembers: 2  
    TimeInformation: "sampleRate"  
    SampleRate: 4000
```

```
Labels: [2x3 table]
Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Visualize the label hierarchy of the set.

```
labelDefinitionsHierarchy(lss)
```

```
ans =
    'WhaleType
      Sublabels: []
    MoanRegions
      Sublabels: []
    TrillRegions
      Sublabels: TrillPeaks
    ,
```

Input Arguments

lbldefs — Signal label definitions

`signalLabelDefinition` object | vector of `signalLabelDefinition` objects

Signal label definitions, specified as a `signalLabelDefinition` object or a vector of `signalLabelDefinition` objects.

Example:

`signalLabelDefinition("Asleep", 'LabelType', 'roi', 'LabelDataType', 'logical')` can label a region of a signal in which a patient is asleep.

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1) randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

Output Arguments

str — List of label and sublabel names

character array

List of label and sublabel names, returned as a character array.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

labelDefinitionsSummary

Get summary table of signal label definitions

Syntax

```
T = labelDefinitionsSummary(lbldefs)
```

```
T = labelDefinitionsSummary(lss)
```

```
T = labelDefinitionsSummary( ____, lblname)
```

```
T = labelDefinitionsSummary( ____, lblname, 'sublbls')
```

Description

T = labelDefinitionsSummary(lbldefs) returns a table, T, with the properties of the label definitions contained in lbldefs, a vector of signalLabelDefinition objects.

T = labelDefinitionsSummary(lss) returns a table, T, with the properties of the label definitions contained in the labeledSignalSet object lss.

T = labelDefinitionsSummary(____, lblname) returns a table with the properties of the label lblname.

T = labelDefinitionsSummary(____, lblname, 'sublbls') returns a table of the properties of the sublabels defined for lblname.

Examples

Label Properties

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =
  labeledSignalSet with properties:
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Visualize the label properties of the set.

```
labelDefinitionsSummary(lss)
```

```
ans=3x9 table
```

LabelName	LabelType	LabelDataType	Categories	ValidationFunction
"WhaleType"	"attribute"	"categorical"	[3x1 string]	["N/A"]
"MoanRegions"	"roi"	"logical"	["N/A"]	[]
"TrillRegions"	"roi"	"logical"	["N/A"]	[]

Visualize the properties of the `TrillRegions` label.

```
labelDefinitionsSummary(lss,"TrillRegions")
```

```
ans=1x9 table
```

LabelName	LabelType	LabelDataType	Categories	ValidationFunction
"TrillRegions"	"roi"	"logical"	["N/A"]	[]

Visualize the properties of the `TrillRegions` sublabels.

```
labelDefinitionsSummary(lss,"TrillRegions",'sublbls')
```

```
ans=1x8 table
```

LabelName	LabelType	LabelDataType	Categories	ValidationFunction
-----------	-----------	---------------	------------	--------------------

"TrillPeaks" "point" "numeric" ["N/A"] []

Input Arguments

lbldefs — Signal label definitions

signalLabelDefinition object | vector of signalLabelDefinition objects

Signal label definitions, specified as a `signalLabelDefinition` object or a vector of `signalLabelDefinition` objects.

Example:

`signalLabelDefinition("Asleep", 'LabelType', 'roi', 'LabelDataType', 'logical')` can label a region of a signal in which a patient is asleep.

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1), randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

Output Arguments

T – Summary table

table

Summary table with the properties of a label.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

labeledSignalSet

Create labeled signal set

Description

Use `labeledSignalSet` to store labeled signals along with the label definitions. Create signal label definitions using `signalLabelDefinition`.

Creation

Syntax

```
lss = labeledSignalSet
```

```
lss = labeledSignalSet(src)
```

```
lss = labeledSignalSet(src, lbldefs)
```

```
lss = labeledSignalSet(src, lbldefs, 'MemberNames', mnames)
```

```
lss = labeledSignalSet(src, lbldefs, Name, Value)
```

Description

`lss = labeledSignalSet` creates an empty labeled signal set. Use `addMembers` to add signals to the set. Use `addLabelDefinitions` to add label definitions to the set.

`lss = labeledSignalSet(src)` creates a labeled signal set for the input data source `src`. Use `addLabelDefinitions` to add label definitions to the set.

`lss = labeledSignalSet(src, lbldefs)` creates a labeled signal set for the input data source `src` using the signal label definitions `lbldefs`. Use `signalLabelDefinition` to create signal label definitions.

`lss = labeledSignalSet(src, lbldefs, 'MemberNames', mnames)` creates a labeled signal set for the input data source `src` and specifies names for the members of the set. Use `setMemberNames` to modify the member names. `lbldefs` is optional.

`lss = labeledSignalSet(src, lbldefs, Name, Value)` sets “Properties” on page 1-685 using name-value pairs. You can specify multiple name-value pairs. Enclose each property name in quotes. `lbldefs` is optional.

Input Arguments

src — Input data source

matrix | cell array | timetable | audio datastore

Input data source, specified as a matrix, a timetable, a cell array, or an audio datastore. `src` specifies the number of members of the set, the number of signals in each member, and the data in each signal.

lbldefs — Label definitions

vector of `signalLabelDefinition` objects

Label definitions, specified as a vector of `signalLabelDefinition` objects.

mnames — Member names

character vector | string scalar | cell array of character vectors | string array

Member names, specified as a character vector, a string scalar, a cell array of character vectors, or a string array.

Example: `labeledSignalSet({randn(100,1) randn(10,1)}, 'MemberNames', {'llama' 'alpaca'})` specifies a set of random signals with two members, 'llama' and 'alpaca'.

Properties

Description — Labeled signal set description

character vector | string scalar

Labeled signal set description, specified as a character vector or string scalar.

Example: 'Description', 'Sleep test patients by sex and age'

Data Types: `char` | `string`

SampleRate — Sample rate values

positive scalar | vector

This property is read-only.

Sample rate values, specified as a positive scalar or a vector. This property is valid only when the data source does not contain inherent time information.

- Set `SampleRate` to a positive numeric scalar to specify the same sample rate for all signals in the labeled set.
- Set `SampleRate` to a vector to specify that each member of the labeled set has signals sampled at the same rate, but the sample rates differ from member to member. The vector must have a number of elements equal to the number of members of the set. If a member of a set has signals with different sample rates, then specify the sample rates using timetables.

Example: `'SampleRate', [1e2 1e3]` specifies that the signals in the first member of a set are sampled at a rate of 100 Hz and the signals in the second member are sampled at 1 kHz.

SampleTime — Sample time values

positive scalar | vector | `duration` scalar | `duration` vector

This property is read-only.

Sample time values, specified as a positive scalar, a vector, a `duration` scalar, or a `duration` vector. This property is valid only when the data source does not contain inherent time information.

- Set `SampleTime` to a numeric or `duration` scalar to specify the same sample time for all signals in the labeled set.
- Set `SampleTime` to a numeric or `duration` vector to specify that each member of the labeled set has signals with the same time interval between samples, but the intervals differ from member to member. The vector must have a number of elements equal to the number of members of the set. If a member of a set has signals with different sample times, then specify the sample times using timetables.

Example: `'SampleTime', seconds([1e-2 1e-3])` specifies that the signals in the first member of a set have 0.01 second between samples, and the signals in the second member have 1 millisecond between samples.

TimeValues — Time values

vector | duration vector | matrix | cell array

This property is read-only.

Time values, specified as a vector, a duration vector, a matrix, or a cell array. This property is valid only when the data source does not contain inherent time information. Time values must be unique and increasing.

- Set `TimeValues` to a numeric or duration vector to specify the same time values for all signals in the labeled set. The vector must have the same length as all the signals in the set.
- Set `TimeValues` to a numeric or duration matrix or cell array to specify that each member of the labeled set has signals with the same time values, but the time values differ from member to member.
 - If `TimeValues` is a matrix, then it must have a number of columns equal to the number of members of the set. All signals in the set must have a length equal to the number of rows of the matrix.
 - If `TimeValues` is a cell array, then it must contain a number of vectors equal to the number of members of the set. All signals in a member must have a length equal to the number of elements of the corresponding vector in the cell array.

If a member of a set has signals with different time values, then specify the time values using timetables.

Example: `'TimeValues', [1:1000;0:1/500:2-1/500]'` specifies that the signals in the first member of a set are sampled 1 Hz for 1000 seconds. The signals in the second member are sampled at 500 Hz for 2 seconds.

Example: `'TimeValues', seconds([1:1000;0:1/500:2-1/500])'` specifies that the signals in the first member of a set are sampled 1 Hz for 1000 seconds. The signals in the second member are sampled at 500 Hz for 2 seconds.

Example: `'TimeValues', {1:1000,0:1/500:2-1/500}` specifies that the signals in the first member of a set are sampled 1 Hz for 1000 seconds. The signals in the second member are sampled at 500 Hz for 2 seconds.

Example: `'TimeValues', {seconds(1:1000), seconds(0:1/500:2-1/500)}` specifies that the signals in the first member of a set are sampled 1 Hz for 1000 seconds. The signals in the second member are sampled at 500 Hz for 2 seconds.

NumMembers — Number of members in set

positive integer

This property is read-only.

Number of members in set, specified as a positive integer.

Labels — Labels table

table

This property is read-only.

Labels table, specified as a MATLAB table. Each variable of `Labels` corresponds to a label defined for the set. Each row of `Labels` corresponds to a member of the data source. The row names of `Labels` are the member names.

Data Types: table

TimeInformation — Time information of source

'None' | 'SampleRate' | 'SampleTime' | 'TimeValues' | 'Inherent'

Time information of source, specified as one of the following:

- 'None' — The signals in the source have no time information.
- 'SampleRate' — The signals in the source are sampled at a specified rate.
- 'SampleTime' — The signals in the source have a specified time interval between samples.
- 'TimeValues' — The signals in the source have a time value corresponding to each sample.
- 'Inherent' — The signals in the source contain inherent time information. MATLAB timetables are an example of such signals.

Data Types: char | string

Source — Data source of labeled signal set

matrix | cell array | timetable

This property is read-only.

Data source of labeled signal set, specified as a matrix, a timetable, a cell array, or an audio datastore.

- If `Source` is a numeric matrix, then the labeled signal set has one member that contains a number of signals equal to the number of matrix columns.

Example: `labeledSignalSet(randn(10,3))` has one member that contains three 10-sample signals.

- If `Source` is a cell array of matrices, then the labeled signal set has a number of members equal to the number of matrices in the cell array. Each member contains a number of signals equal to the number of columns of the corresponding matrix.

Example: `labeledSignalSet({randn(10,3), randn(17,9)})` has two members. The first member contains three 10-sample signals. The second member contains nine 17-sample signals.

- If `Source` is a cell array, and each element of the cell array is a cell array of numeric vectors, then the labeled signal set has a number of members equal to the number of cell array elements. Each signal within a member can have any length.

Example: `labeledSignalSet({{randn(10,1)}, {randn(17,1), randn(27,1)}})` has two members. The first member contains one 10-sample signal. The second member contains a 17-sample signal and a 27-sample signal.

- If `Source` is a timetable with variables containing numeric values, then the labeled signal set has one member that contains a number of signals equal to the number of variables. The time values of the timetable must be of type `duration`, unique, and increasing.

Example: `labeledSignalSet(timetable(seconds(1:10)', randn(10,3)))` has one member that contains three signals sampled at 1 Hz for 10 seconds.

- If `Source` is a cell array of timetables, and each timetable has an arbitrary number of variables with numeric values, then the labeled signal set has a number of members equal to the number of timetables. Each member contains a number of signals equal to the number of variables in the corresponding timetable.

Example:

`labeledSignalSet({timetable(seconds(1:10)', randn(10,3)), timetable(seconds(1:5)', randn(5,13))})` has two members. The first member contains three signals sampled at 1 Hz for 10 seconds. The second member contains 13 signals sampled at 1 Hz for 5 seconds.

- If `Source` is a cell array, and each element of the cell array is a cell array of timetables, then the labeled signal set has a number of members equal to the number

of cell array elements. Each member can have any number of timetables, and each timetable within a member can have any number of variables.

Example:

`labeledSignalSet({{timetable(seconds(1:10) ', randn(10,3)), timetable(seconds(1:7) ', randn(7,2))}, {timetable(seconds(1:3) ', randn(3,1))})` has two members. The first member contains three signals sampled at 1 Hz for 10 seconds and two signals sampled at 1 Hz for 7 seconds. The second member contains one signal sampled at 1 Hz for 3 seconds.

- If the input data source, `src`, is an audio datastore, then the labeled signal set has a number of members equal to the number of files to which the datastore points. The `Source` property contains a cell array of character vectors with the file names. Each member contains all the signals returned by the read of the corresponding datastore file.

Object Functions

<code>addLabelDefinitions</code>	Add label definitions to labeled signal set
<code>addMembers</code>	Add members to labeled signal set
<code>concatenate</code>	Concatenate two or more labeled signal sets
<code>editLabelDefinition</code>	Edit label definition properties
<code>getLabelDefinitions</code>	Get label definitions in labeled signal set
<code>getLabeledSignal</code>	Get labeled signals from labeled signal set
<code>getLabelNames</code>	Get label names in labeled signal set
<code>getLabelValues</code>	Get label values from labeled signal set
<code>getMemberNames</code>	Get member names in labeled signal set
<code>getSignal</code>	Get signals from labeled signal set
<code>head</code>	Get top rows of labels table
<code>labelDefinitionsHierarchy</code>	Get hierarchical list of label and sublabel names
<code>labelDefinitionsSummary</code>	Get summary table of signal label definitions
<code>removeLabelDefinition</code>	Remove label definition from labeled signal set
<code>removeMembers</code>	Remove members from labeled signal set
<code>removePointValue</code>	Remove row from point label
<code>removeRegionValue</code>	Remove row from ROI label
<code>resetLabelValues</code>	Reset labels to default values
<code>setLabelValue</code>	Set label value in labeled signal set
<code>setMemberNames</code>	Set member names in labeled signal set
<code>subset</code>	Get new labeled signal set with subset of members

Examples

Label Definitions for Whale Songs

Consider a set of whale sound recordings. The recorded whale sounds consist of trills and moans. You want to look at each signal and label it to identify the whale type, the trill regions, and the moan regions. For each trill region, you also want to label the signal peaks higher than a certain threshold.

Signal Label Definitions

Define an attribute label to store whale types. The possible categories are blue whale, humpback whale, and white whale.

```
dWhaleType = signalLabelDefinition('WhaleType',...
    'LabelType','attribute',...
    'LabelDataType','categorical',...
    'Categories', ["blue" "humpback" "white"],...
    'Description','Whale type');
```

Define a region-of-interest (ROI) label to capture moan regions. Define another ROI label to capture trill regions.

```
dMoans = signalLabelDefinition('MoanRegions',...
    'LabelType','roi',...
    'LabelDataType','logical',...
    'Description','Regions where moans occur');

dTrills = signalLabelDefinition('TrillRegions',...
    'LabelType','roi',...
    'LabelDataType','logical',...
    'Description','Regions where trills occur');
```

Finally, define a point label to capture the trill peaks. Set this label as a sublabel of the dTrills definition.

```
dTrillPeaks = signalLabelDefinition('TrillPeaks',...
    'LabelType','point',...
    'LabelDataType','numeric',...
    'Description','Trill peaks');

dTrills.Sublabels = dTrillPeaks;
```

Labeled Signal Set

Create a `labeledSignalSet` with the whale signals and the label definitions. Add label values to identify the whale type, the moan and trill regions, and the peaks of the trills.

```
load labelwhalesignals
lbldefs = [dWhaleType dMoans dTrills];

lss = labeledSignalSet({whale1 whale2},lbldefs,...
    'SampleRate',Fs,'Description','Characterize wave song regions');
```

Visualize the label hierarchy and label properties using `labelDefinitionsHierarchy` and `labelDefinitionsSummary`.

```
labelDefinitionsHierarchy(lss)
```

```
ans =
    'WhaleType
      Sublabels: []
    MoanRegions
      Sublabels: []
    TrillRegions
      Sublabels: TrillPeaks
    '
```

```
labelDefinitionsSummary(lss)
```

```
ans=3×9 table
    LabelName      LabelType      LabelDataType      Categories      ValidationFunction
    _____      _____      _____      _____      _____
    "WhaleType"     "attribute"    "categorical"     [3×1 string]   ["N/A"]
    "MoanRegions"  "roi"         "logical"         ["N/A" ]       []
    "TrillRegions" "roi"         "logical"         ["N/A" ]       []
```

The signals in the loaded data correspond to songs of two blue whales. Set the 'WhaleType' values for both signals.

```
setLabelValue(lss,1,'WhaleType','blue');
setLabelValue(lss,2,'WhaleType','blue');
```

Visualize the 'Labels' property. The table has the newly added 'WhaleType' values for both signals.

```
lss.Labels
```

```
ans=2x3 table
```

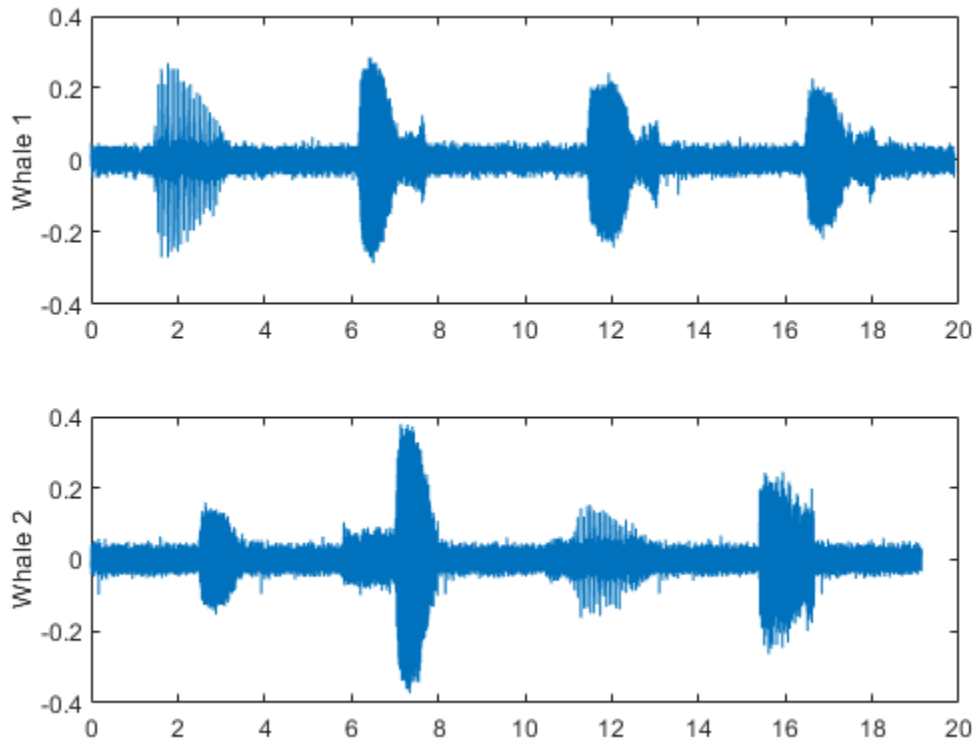
	WhaleType	MoanRegions	TrillRegions
Member{1}	blue	[0x2 table]	[0x3 table]
Member{2}	blue	[0x2 table]	[0x3 table]

Visualize Regions and Points

Visualize the whale songs to identify the trill and moan regions.

```
subplot(2,1,1)
plot((0:length(whale1)-1)/Fs,whale1)
ylabel('Whale 1')
hold on
```

```
subplot(2,1,2)
plot((0:length(whale2)-1)/Fs,whale2)
ylabel('Whale 2')
hold on
```



Add the moan and trill regions to the labeled set. For ROI labels, specify the ROI limits in seconds and the label values. Label the different regions in the plots using an auxiliary function.

```
moanRegionsWhale1 = [6.1 7.7; 11.4 13.1; 16.5 18.1];  
setLabelValue(lss,1,'MoanRegions',moanRegionsWhale1,[true true true]);  
  
trillRegionWhale1 = [1.4 3.1];  
setLabelValue(lss,1,'TrillRegions',trillRegionWhale1,true);  
  
subplot(2,1,1)  
labelIntervals([moanRegionsWhale1;trillRegionWhale1])  
  
moanRegionsWhale2 = [2.5 3.5; 5.8 8; 15.4 16.7];
```

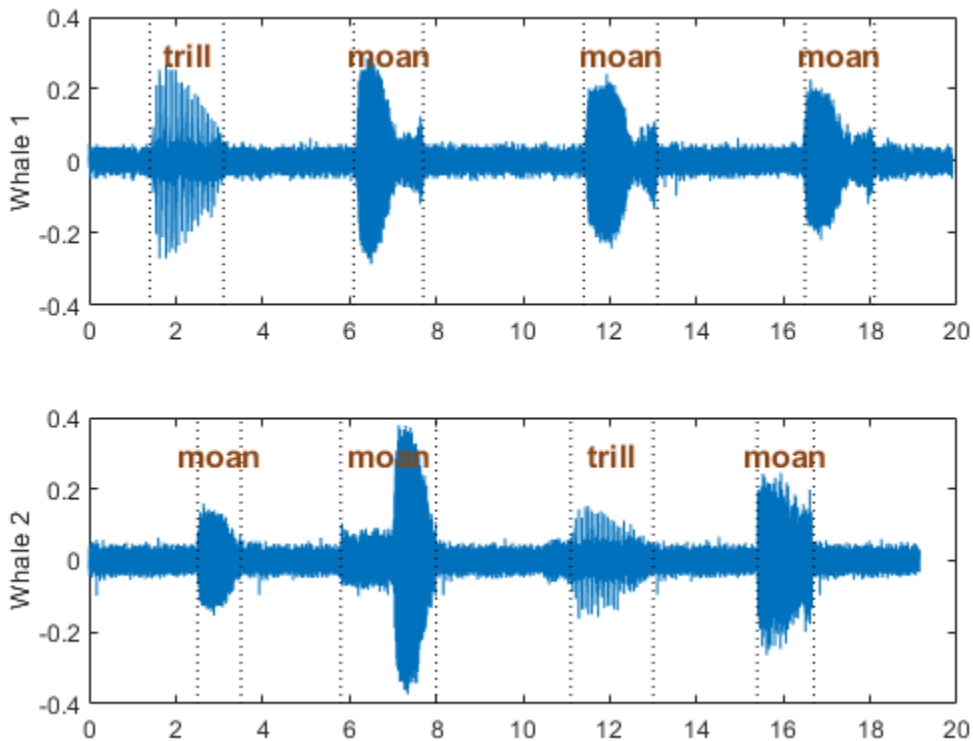
```

setLabelValue(lss,2,'MoanRegions',moanRegionsWhale2,[true true true]);

trillRegionWhale2 = [11.1 13];
setLabelValue(lss,2,'TrillRegions',trillRegionWhale2,true);

subplot(2,1,2)
labelIntervals([moanRegionsWhale2;trillRegionWhale2])

```



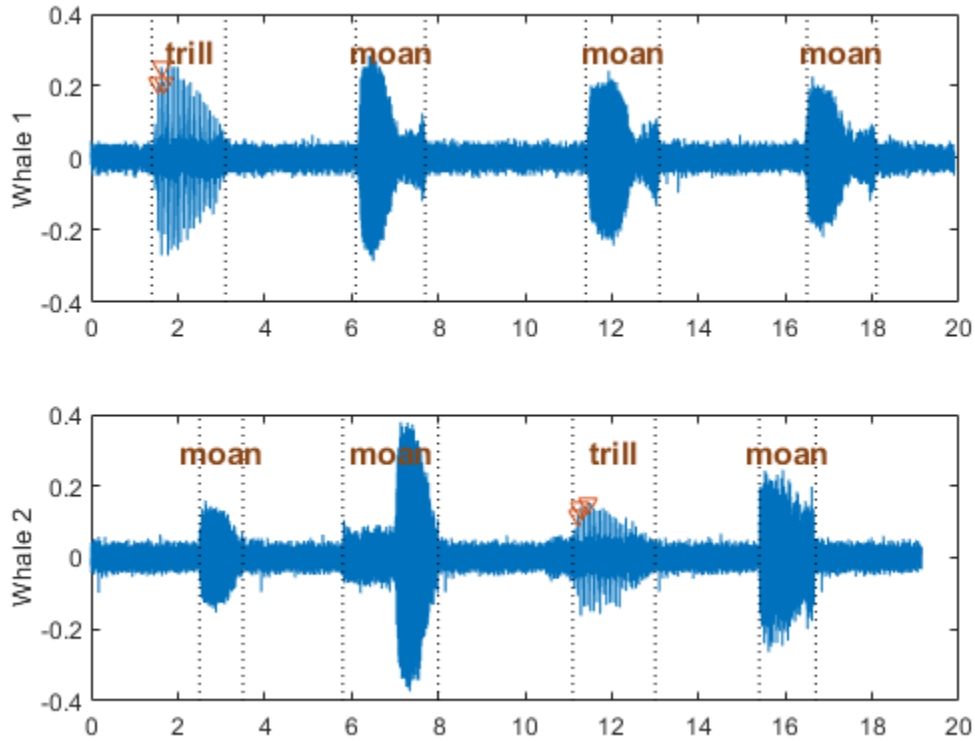
Label three peaks for each trill region. For point labels, you specify the point locations and the label values. In this example, the point locations are in seconds.

```

peakLocsWhale1 = [1.553 1.626 1.7];
peakValsWhale1 = [0.211 0.254 0.211];

```

```
setLabelValue(lss,1,["TrillRegions" "TrillPeaks"],...  
    peakLocsWhale1,peakValsWhale1,'LabelRowIndex',1);  
  
subplot(2,1,1)  
plot(peakLocsWhale1,peakValsWhale1,'v')  
hold off  
  
peakLocsWhale2 = [11.214 11.288 11.437];  
peakValsWhale2 = [0.119 0.14 0.15];  
  
setLabelValue(lss,2,["TrillRegions" "TrillPeaks"],...  
    peakLocsWhale2,peakValsWhale2,'LabelRowIndex',1);  
  
subplot(2,1,2)  
plot(peakLocsWhale2,peakValsWhale2,'v')  
hold off
```



Explore Label Values

Explore the label values using `getLabelValues`.

```
getLabelValues(lss)
```

```
ans=2x3 table
```

	WhaleType	MoanRegions	TrillRegions
Member{1}	blue	[3x2 table]	[1x3 table]
Member{2}	blue	[3x2 table]	[1x3 table]

Retrieve the moan regions for the first member of the labeled set.

```
getLabelValues(lss,1, 'MoanRegions')
```

```
ans=3x2 table
  ROIlimits      Value
  _____      _____
    6.1      7.7      [1]
   11.4     13.1     [1]
   16.5     18.1     [1]
```

Use a second output argument to list the sublabels of a label.

```
[value,valueWithSublabel] = getLabelValues(lss,1, 'TrillRegions')
```

```
value=1x2 table
  ROIlimits      Value
  _____      _____
    1.4      3.1      [1]
```

```
valueWithSublabel=1x3 table
  ROIlimits      Value      Sublabels
  _____      _____      _____
    1.4      3.1      [1]      [3x2 table]
```

To retrieve the values in a sublabel, express the label name as a two-element array.

```
getLabelValues(lss,1,["TrillRegions" "TrillPeaks"])
```

```
ans=3x2 table
  Location      Value
  _____      _____
    1.553     [0.2110]
    1.626     [0.2540]
    1.7       [0.2110]
```

Find the value of the third trill peak corresponding to the second member of the set.


```
getLabelValues(lss,2,["TrillRegions" "TrillPeaks"], ...
    'LabelRowIndex',1,'SublabelRowIndex',3)
```

```
ans=1x2 table
    Location      Value
    _____  _____
    11.437        [0.1500]
```

```
function labelIntervals(moansTrills)
% Auxiliary function to label moan and trill regions in plots
[X,Y] = meshgrid(moansTrills,ylim);
plot(X,Y,'k:')
topts = {'HorizontalAlignment','center','FontWeight','bold', ...
    'FontSize',12,'Color',[139 69 19]/255};
text((X(1,1:4)+X(1,5:end))/2,Y(2,5:end)-0.1, ...
    ["moan" "moan" "moan" "trill"],topts{:})
end
```

See Also

signalLabelDefinition

Introduced in R2018b

laurmat

Laurent matrices constructor

Syntax

```
M = laurmat(V)
```

Description

`M = laurmat(V)` returns the Laurent matrix object `M` associated with `V` which can be a cell array (at most two dimensional) of Laurent polynomials (see `laurpoly`) or an ordinary matrix.

Examples

```
% Define Laurent matrices.  
M1 = laurmat(eye(2,2))
```

$$M1 = \begin{array}{|c|cc|} \hline & 1 & 0 \\ \hline & & \\ \hline & 0 & 1 \\ \hline \end{array}$$

```
Z = laurpoly(1,1);  
M2 = laurmat({1 Z;0 1})
```

$$M2 = \begin{array}{|c|cc|} \hline & 1 & z^{(+1)} \\ \hline & & \\ \hline & 0 & 1 \\ \hline \end{array}$$

```
% Calculus on Laurent polynomials.  
P = M1 * M2
```

$$P = \begin{vmatrix} 1 & z^{(+1)} \\ 0 & 1 \end{vmatrix}$$

$$d = \det(P)$$

$$d(z) = 1$$

References

Strang, G.; T. Nguyen (1996), *Wavelets and filter banks*, Wellesley-Cambridge Press.

Sweldens, W. (1998), "The Lifting Scheme: a Construction of Second Generation of Wavelets," *SIAM J. Math. Anal.*, 29 (2), pp. 511-546.

See Also

laurpoly

Introduced before R2006a

laurpoly

Laurent polynomials constructor

Syntax

```
P = laurpoly(C,d)
P = laurpoly(C,'dmin',d)
P = laurpoly(C,'dmax',d)
P = laurpoly(C,d)
```

Description

`P = laurpoly(C,d)` returns a Laurent polynomial object. C is a vector whose elements are the coefficients of the polynomial P and d is the highest degree of the monomials of P .

If m is the length of the vector C , P represents the following Laurent polynomial:

$$P(z) = C(1)*z^d + C(2)*z^{(d-1)} + \dots + C(m)*z^{(d-m+1)}$$

`P = laurpoly(C,'dmin',d)` specifies the lowest degree instead of the highest degree of monomials of P . The corresponding output P represents the following Laurent polynomial:

$$P(z) = C(1)*z^{(d+m-1)} + \dots + C(m-1)*z^{(d+1)} + C(m)*z^d$$

`P = laurpoly(C,'dmax',d)` is equivalent to `P = laurpoly(C,d)`.

Examples

```
% Define Laurent polynomials.
P = laurpoly([1:3],2);
P = laurpoly([1:3],'dmax',2)

P(z) = + z^{+2} + 2*z^{+1} + 3

P = laurpoly([1:3],'dmin',2)
```

$$P(z) = z^{+4} + 2z^{+3} + 3z^{+2}$$

% Calculus on Laurent polynomials.

$$Z = \text{laurpoly}(1,1)$$

$$Z(z) = z^{+1}$$

$$Q = Z * P$$

$$Q(z) = z^{+5} + 2z^{+4} + 3z^{+3}$$

$$R = Z^1 - Z^{-1}$$

$$R(z) = z^{+1} - z^{-1}$$

References

Strang, G.; T. Nguyen (1996), *Wavelets and filter banks*, Wellesley-Cambridge Press.

Sweldens, W. (1998), "The Lifting Scheme: a Construction of Second Generation of Wavelets," *SIAM J. Math. Anal.*, 29 (2), pp. 511-546.

See Also

laurmat

Introduced before R2006a

leaves

Determine terminal nodes

Syntax

```
N = leaves(T)
[N,K] = leaves(T,'sort')
N = leaves(T,'dp')
[N,K] = leaves(T,'sortdp')
[N,K] = leaves(T,'sdp')
```

Description

`N = leaves(T)` returns the indices of terminal nodes of the tree T where N is a column vector.

The nodes are ordered from left to right as in tree T .

`[N,K] = leaves(T,'s')` or `[N,K] = leaves(T,'sort')` returns sorted indices. $M = N(K)$ are the indices reordered as in tree T , from left to right.

`N = leaves(T,'dp')` returns a matrix N , which contains the depths and positions of terminal nodes.

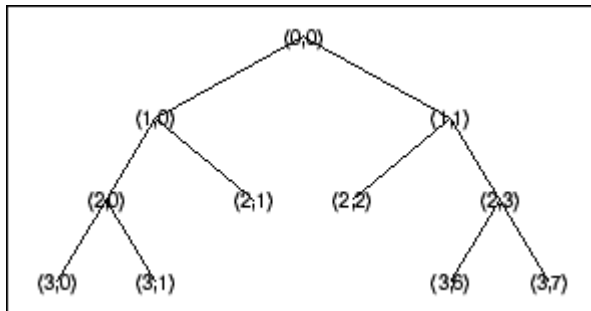
$N(i,1)$ is the depth of the i -th terminal node, and $N(i,2)$ is the position of the i -th terminal node.

`[N,K] = leaves(T,'sortdp')` or `[N,K] = leaves(T,'sdp')` returns sorted nodes.

Examples

```
% Create initial tree.
ord = 2;
t = ntree(ord,3);           % binary tree of depth 3.
t=nodejoin(t,5);
```

```
t=nodejoin(t,4);
plot(t)
```



```
% List terminal nodes (index).
tnodes_ind = leaves(t)
tnodes_ind =
    7
    8
    4
    5
   13
   14

% List terminal nodes (sorted on index).
[tnodes_ind,Ind] = leaves(t,'sort')
tnodes_ind =
    4
    5
    7
    8
   13
   14

Ind =
    3
    4
    1
    2
    5
    6

% List terminal nodes (Depth_Position).
tnodes_depo = leaves(t,'dp')
```

```
tnodes_depo =  
    3     0  
    3     1  
    2     1  
    2     2  
    3     6  
    3     7  
  
% List terminal nodes (sorted on Depth_Position).  
[tnodes_depo,Ind] = leaves(t,'sortdp')  
tnodes_depo =  
    2     1  
    2     2  
    3     0  
    3     1  
    3     6  
    3     7  
  
Ind =  
    3  
    4  
    1  
    2  
    5  
    6
```

See Also

noleaves | tnodes

Introduced before R2006a

liftfilt

Apply elementary lifting steps on quadruplet of filters

Syntax

```
[LoDN,HiDN,LoRN,HiRN] = liftfilt(LoD,HiD,LoR,HiR,ELS)  
liftfilt(LoD,HiD,LoR,HiR,ELS,TYPE,VALUE)
```

Description

`[LoDN,HiDN,LoRN,HiRN] = liftfilt(LoD,HiD,LoR,HiR,ELS)` returns the four filters `LoDN`, `HiDN`, `LoRN`, and `HiRN` obtained by an elementary lifting step (ELS) starting from the four filters `LoD`, `HiD`, `LoR`, and `HiR`. The four input filters verify the perfect reconstruction condition.

ELS is a structure such that

- `TYPE = ELS.type` contains the type of the elementary lifting step. The valid values for `TYPE` are 'p' (primal) or 'd' (dual).
- `VALUE = ELS.value` contains the Laurent polynomial `T` associated with the elementary lifting step (see `laurpoly`). If `VALUE` is a vector, the associated Laurent polynomial `T` is equal to `laurpoly(VALUE,0)`.

In addition, ELS may be a scaling step. In that case, `TYPE` is equal to 's' (scaling) and `VALUE` is a scalar different from zero.

`liftfilt(LoD,HiD,LoR,HiR,ELS,TYPE,VALUE)` gives the same outputs.

Note If `TYPE = 'p'`, `HiD` and `LoR` are unchanged.

If `TYPE = 'd'`, `LoD` and `HiR` are unchanged.

If `TYPE = 's'`, the four filters are changed.

If ELS is an array of elementary lifting steps, `liftfilt(...,ELS)` performs each step successively.

`liftfilt(..., FLAGPLOT)` plots the successive biorthogonal pairs—scaling function and wavelet.

Examples

```
% Get Haar filters.
[LoD,HiD,LoR,HiR] = wfilters('haar');

% Lift the Haar filters.
twoels(1) = struct('type','p','value',...
    laurpoly([0.125 -0.125],0));
twoels(2) = struct('type','p','value',...
    laurpoly([0.125 -0.125],1));
[LoDN,HiDN,LoRN,HiRN] = liftfilt(LoD,HiD,LoR,HiR,twoels);

% The biorthogonal wavelet bior1.3 is obtained up to
% an insignificant sign.
[LoDB,HiDB,LoRB,HiRB] = wfilters('bior1.3');
samewavelet = ...
isequal([LoDB,HiDB,LoRB,HiRB],[LoDN,-HiDN,LoRN,HiRN])

samewavelet =

    1
```

See Also

`laurpoly`

Introduced before R2006a

liftwave

Lifting schemes

Syntax

```
LS = liftwave(WNAME)
LS = liftwave(WNAME, 'Int2Int')
```

Description

`LS = liftwave(WNAME)` returns the lifting scheme associated with the wavelet specified by `WNAME`. `LS` is a structure, not an integer, and used by `lwt`, `ilwt`, `lwt2`, etc.

`LS = liftwave(WNAME, 'Int2Int')` performs an integer to integer wavelet transform. Using `'Int2Int'` produces an `LS` such that when you use `[CA,CD] = lwt(X,LS)` or `Y = lwt(X,LS)` and `X` is a vector of integers, the resulting `CA`, `CD`, and `Y` are vectors of integers. If you omit `'Int2Int'` then `lwt` produces vectors of real numbers.

The valid values for `WNAME` are

WNAME Values
'lazy'
'haar'
'db1', 'db2', 'db3', 'db4', 'db5', 'db6', 'db7', 'db8'
'sym2', 'sym3', 'sym4', 'sym5', 'sym6', 'sym7', 'sym8'
Cohen-Daubechies-Feauveau wavelets
'cdf1.1', 'cdf1.3', 'cdf1.5'
'cdf3.1', 'cdf3.3', 'cdf3.5'
'cdf5.1', 'cdf5.3', 'cdf5.5'
'cdf2.2', 'cdf2.4', 'cdf2.6'
'cdf4.2', 'cdf4.4', 'cdf4.6'
'cdf6.2', 'cdf6.4', 'cdf6.6'

WNAME Values
'biorX.Y'
'rbioX.Y'
'bs3'
'rbs3'
'9.7'
'r9.7'

For more information about lifting schemes, see `lsinfo`.

Examples

```
% Start from the db2 wavelet and get the  
% corresponding lifting scheme.  
lsdb2 = liftwave('db2');
```

```
% Visualize the obtained lifting scheme.  
displs(lsdb2);
```

```
lsdb2 = {...  
'd'          [ -1.73205081]          [0]  
'p'          [ -0.06698730  0.43301270] [1]  
'd'          [  1.00000000]          [-1]  
[  1.93185165] [  0.51763809]          []  
};
```

See Also

`laurpoly`

Introduced before R2006a

littlewoodPaleySum

Littlewood-Paley sum

Syntax

```
lpsum = littlewoodPaleySum(sf)
lpsum = littlewoodPaleySum(sf,fb)
[lpsum,f] = littlewoodPaleySum( ___ )
```

Description

`lpsum = littlewoodPaleySum(sf)` returns the Littlewood-Paley sum for the scattering filter banks in `sf`, the scattering decomposition framework. `lpsum` is an M -by- L matrix, where M is the number of elements in the Fourier transform of the scattering filters, and L is the number of scattering filter banks. The columns of `lpsum` are ordered by the position of the filter bank in the scattering decomposition. For example, the first column of `lpsum` corresponds to the filter bank used for the first-order scattering coefficients.

Since the scattering transform is contractive, the Littlewood-Paley sums will not exceed one.

`lpsum = littlewoodPaleySum(sf,fb)` returns the Littlewood-Paley sum for the specified filter bank `fb` in `sf`. The argument `fb` is a positive integer between 1 and the number of filter banks in `sf` inclusive. The number of filter banks in `sf` is equal to the number of specified `QualityFactors` in `sf`.

`[lpsum,f] = littlewoodPaleySum(___)` returns the frequencies for the Littlewood-Paley sum. If you specify a sampling frequency in `sf`, `f` is in hertz. If you do not specify a sampling frequency, `f` is in cycles/sample. You can use these output arguments with any of the input syntaxes shown previously.

Examples

Plot Littlewood-Paley Sum

Create a scattering decomposition framework with three filter banks for data sampled at 25 Hz.

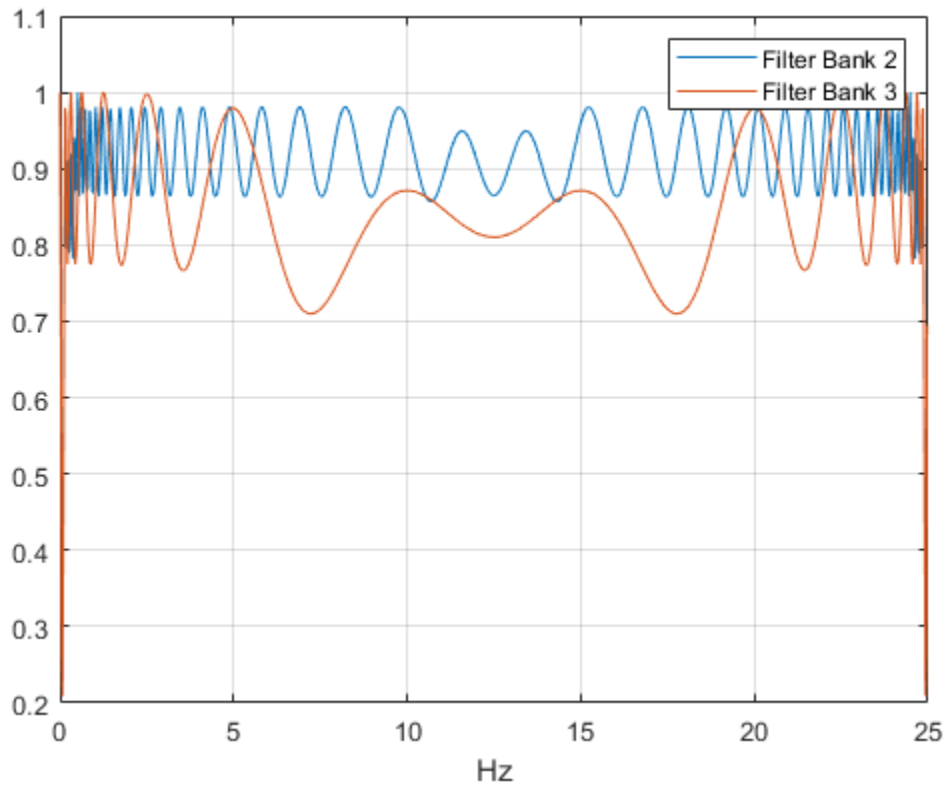
```
sf = waveletScattering('QualityFactors',[8 4 1],...  
    'SamplingFrequency',25)
```

```
sf =  
    waveletScattering with properties:
```

```
    SignalLength: 1024  
    InvarianceScale: 20.4800  
    QualityFactors: [8 4 1]  
        Boundary: "periodic"  
    SamplingFrequency: 25  
        Precision: "double"  
    OversamplingFactor: 0
```

Plot the Littlewood-Paley sums for the second and third filter banks. Note that the sums do not exceed 1. This shows the filters have been normalized so that the scattering transform is contractive.

```
[lpsum,f] = littlewoodPaleySum(sf);  
plot(f,lpsum(:,2:3))  
grid on  
legend('Filter Bank 2','Filter Bank 3')  
xlabel('Hz')
```



Input Arguments

sf — Scattering decomposition framework

waveletScattering object

Scattering decomposition framework, specified as a waveletScattering object.

fb — Filter bank index

positive integer

Filter bank index in the scattering decomposition framework, specified as a positive integer between 1 and the number of filter banks in `sf` inclusive. The number of filter banks in `sf` is equal to the number of specified `QualityFactors` in `sf`.

Data Types: `double`

Output Arguments

lpsum — Littlewood-Paley sum

real-valued matrix

Littlewood-Paley sum for the filter banks in the scattering decomposition framework `sf`, returned as a real-valued matrix. `lpsum` is an M -by- L matrix, where M is the number of elements in the Fourier transform of the scattering filters and L is the number of scattering filter banks. For example, the first column of `lpsum` corresponds to the filter bank used for the first-order scattering coefficients.

f — Frequencies

real-valued vector

Frequencies for the Littlewood-Paley sum, returned as a real-valued vector. If you specify a sampling frequency in `sf`, `f` is in hertz. If you do not specify a sampling frequency, `f` is in cycles/sample.

Data Types: `double`

See Also

`waveletScattering`

Introduced in R2018b

littlewoodPaleySum

Littlewood-Paley sum

Syntax

```
lpsum = littlewoodPaleySum(sf)
lpsum = littlewoodPaleySum(sf,fb)
[lpsum,f] = littlewoodPaleySum( ___ )
```

Description

`lpsum = littlewoodPaleySum(sf)` returns the Littlewood-Paley sum for the 2-D filter banks in the 2-D wavelet scattering decomposition `sf`. `lpsum` is an M -by- N -by- Nfb matrix, where M -by- N is the matrix size of the padded filters and Nfb is the number of filter banks.

Since the scattering transform is contractive, the Littlewood-Paley sums do not exceed 1.

`lpsum = littlewoodPaleySum(sf,fb)` returns the Littlewood-Paley sum for the specified filter banks `fb`. `fb` is a positive integer or vector of positive integers between 1 and `numfilterbanks(sf)` inclusive. `lpsum` is an M -by- N -by- L matrix, where L is the number of unique elements in `fb`.

`[lpsum,f] = littlewoodPaleySum(___)` returns the spatial frequencies for the Littlewood-Paley sum. `f` is a two-column matrix with the first column containing the spatial frequencies in the x -direction and the second column containing the spatial frequencies in the y -direction.

Examples

Littlewood-Paley Sum of Image Scattering Framework

This example shows how to obtain and display the Littlewood-Paley sum of an image scattering framework.

Create a scattering framework with two filter banks and quality factors of 2 and 1, respectively.

```
sf = waveletScattering2('QualityFactors',[2 1]);
```

Obtain the Littlewood-Paley sums and spatial frequencies of the two filter banks. Display the maximum value of the sums. Since the scattering transform is contractive, the sums do not exceed 1.

```
[lpsum,f] = littlewoodPaleySum(sf);  
max(max(lpsum(:,:,1)))  
max(max(lpsum(:,:,2))))
```

```
ans =
```

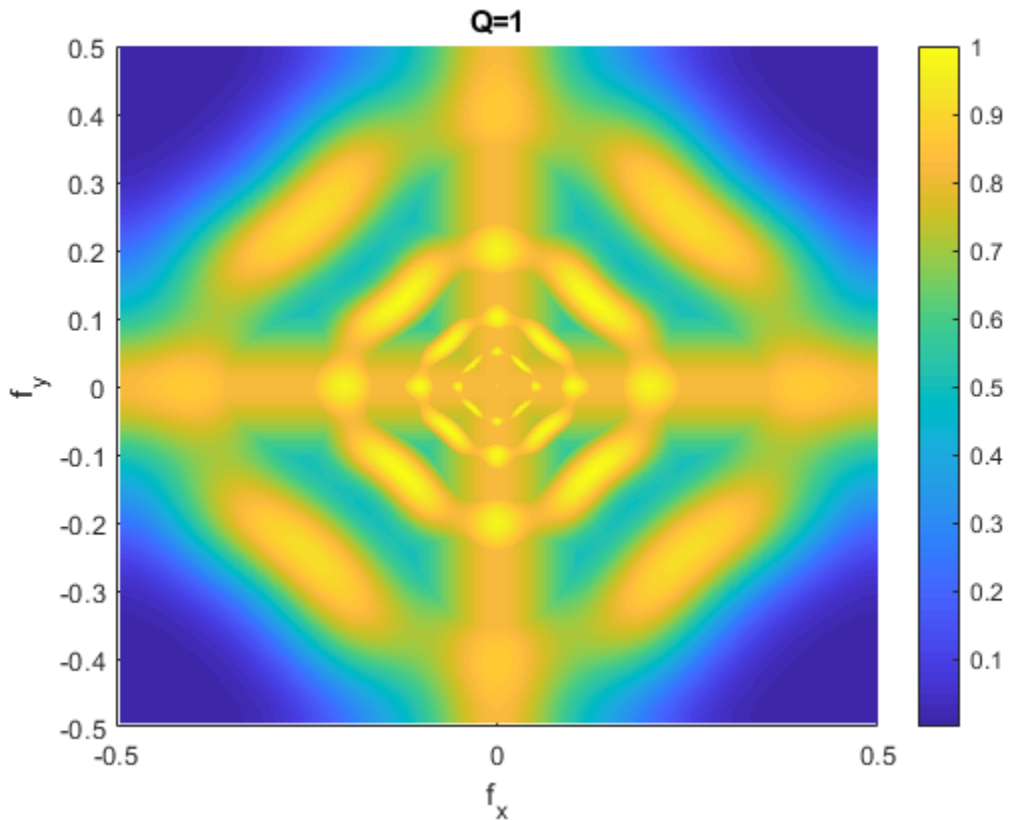
```
1.0000
```

```
ans =
```

```
1.0000
```

Display the Littlewood-Paley sum of the second filter bank with the zero frequency centered. Note the 2-D Morlet filter bank used in the scattering transform is not designed to capture the highest spatial frequencies jointly in the x- and y-directions.

```
f(f>1/2) = f(f>1/2)-1;  
surf(fftshift(f(:,1)),fftshift(f(:,2)),fftshift(lpsum(:,:,2)))  
shading interp  
view(0,90)  
xlabel('f_x')  
ylabel('f_y')  
colorbar  
title('Q=1')
```



Input Arguments

sf – Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

fb – Filter bank index

positive integer | vector of positive integers

Filter bank index in the image scattering decomposition framework, specified as a positive integer or vector of positive integers between 1 and numfilterbanks(sf)

inclusive. The number of filter banks in `sf` is equal to the number of specified `QualityFactors` in `sf`.

Output Arguments

lpsum — Littlewood-Paley sum

real-valued 3-D matrix

Littlewood-Paley sum for the filter banks in the image scattering decomposition framework `sf`, returned as a real-valued 3-D matrix. `lpsum` is an M -by- N -by- L matrix, where M -by- N is the matrix size of the padded filters and L does not exceed the number of filter banks in `sf`.

f — Frequencies

real-valued two-column matrix

Frequencies for the Littlewood-Paley sum, returned as a real-valued two-column matrix. Frequencies are in cycles per pixel. The first column of `f` contains the spatial frequencies in the x -direction, and the second column contains the spatial frequencies in the y -direction. In this convention, the Fourier transform is 1-periodic in both Fourier variables.

See Also

`waveletScattering2`

Introduced in R2019a

localmax

Identify and chain local maxima

Syntax

```
[lmaxima,indices] = localmax(inputmatrix)
[lmaxima,indices] = localmax(inputmatrix,initrow)
[lmaxima,indices] = localmax(inputmatrix,initrow,regflag)
```

Description

[lmaxima,indices] = localmax(inputmatrix) identifies and chains the local maxima in the rows of inputmatrix.

[lmaxima,indices] = localmax(inputmatrix,initrow) initializes the chaining of local maxima beginning with row initrow. If there are no local maxima in initrow, all rows in lmaxima with indices less than initrow consist of only zeros.

[lmaxima,indices] = localmax(inputmatrix,initrow,regflag) replaces initrow of inputmatrix with the level-5 approximation (scaling) coefficients obtained with the sym4 wavelet.

Input Arguments

inputmatrix

inputmatrix is a matrix of real or complex numbers. Most often, inputmatrix is a matrix of continuous wavelet transform (CWT) coefficients, and you use localmax to identify maxima lines. localmax operates on the absolute values of inputmatrix.

initrow

Initialization row for chaining local maxima. The chaining algorithm begins at initrow and decrements the row index by 1 until the first row of the matrix is reached. By specifying initrow, you can exclude rows from the chaining algorithm.

Default: `size(inputmatrix,1)`

regflag

Regularization flag. If you set `regflag` to `true`, the row of `inputmatrix` corresponding to `initrow` is replaced by the level-5 approximation (scaling) coefficients obtained with the `sym4` wavelet.

Default: `true`

Output Arguments

lmaxima

Matrix with local maxima chains. `lmaxima` only has nonzero entries at the locations of local maxima in the absolute values of `inputmatrix`. Denote the row index of `lmaxima` by `R`. You can determine the value of `lmaxima` at a local maximum in row `R` as follows:

- If `R>initRow`, the value of `lmaxima` at a local maximum is 1.
- If `R=initRow`, the value of `lmaxima` at a local maximum is the column index in row `R`.
- If `R<initRow`, the value of `lmaxima` at a local maximum in row `R` is the column index of the nearest local maximum in row `R+1`.

To illustrate this, if `inputmatrix` is:

```
3   2   5   3
4   6   3   2
4   4   7   4
4   6   2   2
```

`lmaxima` with `initRow = 4` and `regflag = false` is:

```
0   0   2   0
0   3   0   0
0   0   2   0
0   2   0   0
```

`lmaxima` with `initRow = 3` and `regflag = false` is:

```
0   0   2   0
0   3   0   0
```

```

0    0    3    0
0    1    0    0

```

- If the local maximum in row R lies between two local maxima in row R+1, the value of the local maximum in row R is the higher column index in row R+1.

To illustrate this, if `inputmatrix` is:

```

0    0    1    0    0    0
0    1    0    1    0    0

```

`lmaxima` with `initRow = 2` and `regflag = false` is:

```

0    0    4    0    0    0
0    2    0    4    0    0

```

`lmaxima` with `initRow = 1` and `regflag = false` is:

```

0    0    3    0    0    0
0    1    0    1    0    0

```

indices

Linear indices of the nonzero values of `lmaxima`. Use `ind2sub` to convert the linear indices to matrix row and column indices.

Examples

Local Maxima of a Matrix

Construct a 4-by-4 matrix with local maxima at the following row-column indices: (4,2), (3,3), (2,2), and (1,3). Set `initrow` to 4 and `regflag` to false.

```

inputmatrix = ...
[3    2    5    3
 4    6    3    2
 4    4    7    4
 4    6    2    2];
[lmaxima,indices] = localmax(inputmatrix,4,false);
lmaxima

```

Because `localmax` operates on the absolute values of `inputmatrix`, setting `inputmatrix(4,2) = -inputmatrix(4,2)` produces an identical `lmaxima`.

```
inputmatrix(4,2) = -inputmatrix(4,2);  
[lmaxima1,indices1] = localmax(inputmatrix,4,false);  
isequal(lmaxima,lmaxima1)
```

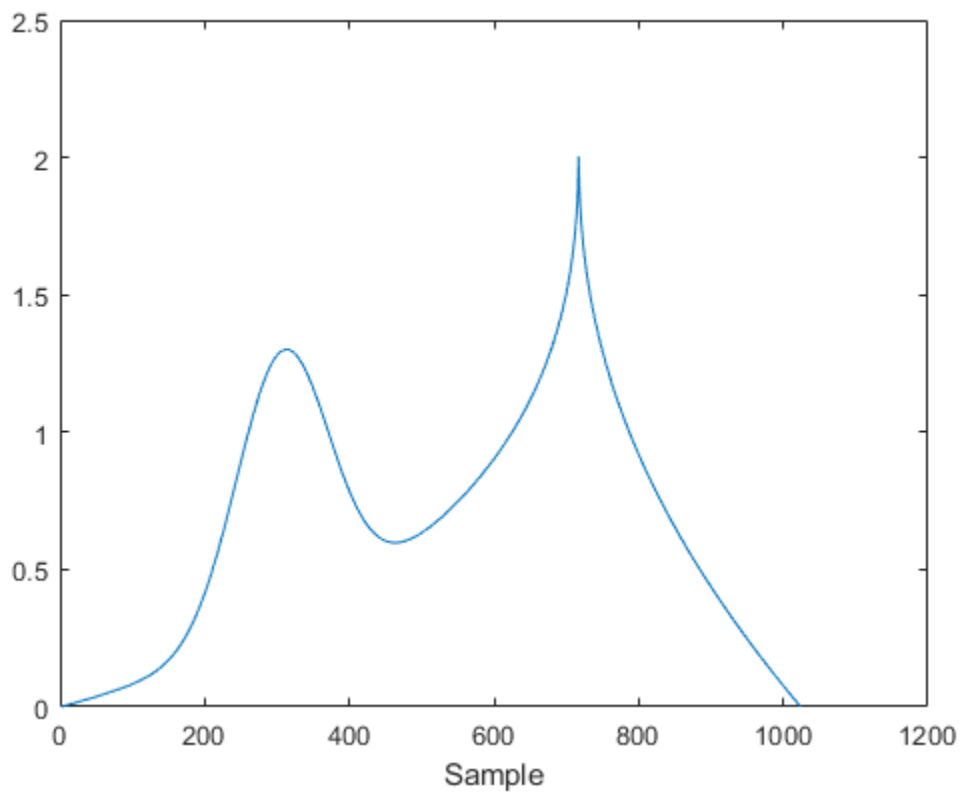
CWT Coefficient Moduli and Maxima Lines

Determine the local maxima from the CWT of the `cuspmamax` signal using the default Morse wavelet. Plot the CWT coefficient moduli and maxima lines.

```
load cuspmamax;
```

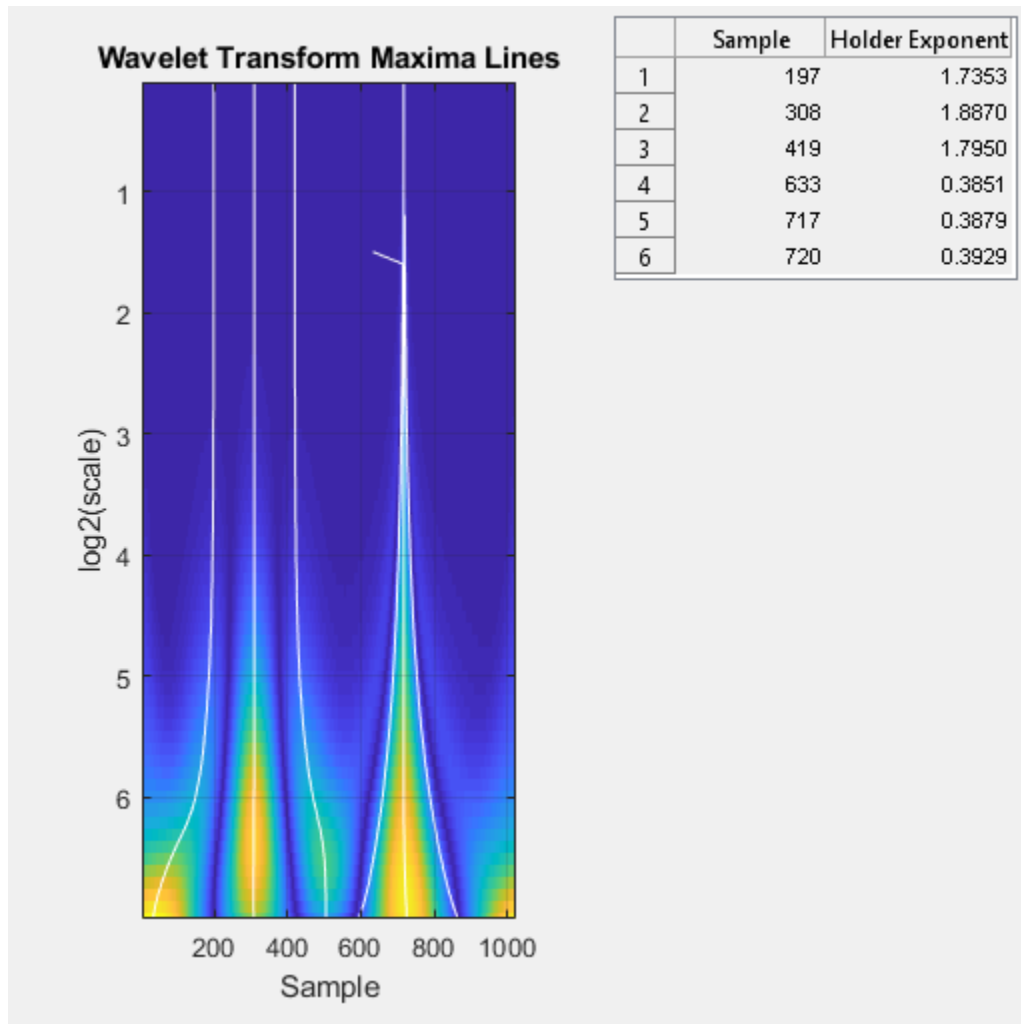
Plot the `cuspmamax` signal and notice the shape of the signal near samples 300 and 700. The signal shows a cusp near sample 700.

```
plot(cuspmamax);  
xlabel('Sample');
```

Plot the wavelet transform modulus maxima and note the local Holder exponent values at samples 308 and 717.

```
wtmm(cuspsmax, 'ScalingExponent', 'local');
```



Holder exponent values indicate the strength of the singularities in a signal. Signal locations where the local Holder exponent is 0 are discontinuous at that location. Locations with Holder exponents greater than or equal to 1 are differentiable. Holder exponent values less than but close to 1 indicate that the signal at the location is almost differentiable. The closer the Holder exponent value is to 0, the stronger the singularity.

The Holder exponent at sample 308 is 1.9 and at sample 717 is 0.39. The low Holder value at sample 717 confirms that the signal is not differentiable and has a fairly strong singularity at that point.

Introduced in R2008a

log

Natural logarithm of scattering transform

Syntax

```
slog = log(sf,s)
ulog = log(sf,u)
xlog = log(sf,x)
```

Description

`slog = log(sf,s)` returns the natural logarithm of the scattering coefficients in the cell array `s`. `s` is the output of `scatteringTransform` and is a cell array of structure arrays with a `signals` field.

The precision of `slog` depends on the precision specified in the framework `sf`.

`ulog = log(sf,u)` returns the natural logarithm of the scalogram coefficients in the cell array `u`. `u` is the output of `scatteringTransform` and is a cell array of structure arrays with a `coefficients` field.

The precision of `ulog` depends on the precision specified in the framework `sf`.

`xlog = log(sf,x)` returns the natural logarithm of the 2-D matrix or 3-D array `x`. `x` is the output of `featureMatrix`.

The precision of `xlog` depends on the precision specified in the framework `sf`.

Examples

Natural Logarithm of Scattering Coefficients

This example shows how to obtain the natural logarithm of scattering coefficients.

Load a noisy Doppler signal and create a scattering decomposition framework that can be used with the signal. Return the scattering coefficients.

```
load noisdopp
sf = waveletScattering('SignalLength', numel(noisdopp));
S = scatteringTransform(sf, noisdopp);
```

Calculate the natural logarithm of the scattering coefficients. Display the number of rows in the table containing the first-order scattering coefficients.

```
slog = log(sf, S);
coefOrder = 1;
display(['Number of rows: ', ...
        num2str(size(S{coefOrder+1}, 1))])
```

```
Number of rows: 40
```

Choose a row from the first-order scattering coefficients table. Take the natural logarithm of the absolute value of the scattering coefficients in that row. Compare with the corresponding row in `slog` and confirm they are equal.

```
row = 23;
tmp1 = slog{coefOrder+1}.signals{row};
tmp2 = log(abs(S{coefOrder+1}.signals{row}));
disp(['Max Difference of Scattering Coefficients: ', ...
      num2str(max(abs(tmp1(:) - tmp2(:)))])])
```

```
Max Difference of Scattering Coefficients: 0
```

Input Arguments

sf — Scattering decomposition framework

`waveletScattering` object

Scattering decomposition framework, specified as a `waveletScattering` object.

s — Scattering coefficients

cell array

Scattering coefficients, specified as a cell array of structure arrays. `s` is the output of `scatteringTransform` for the scattering decomposition framework `sf`.

u — Scalogram coefficients

cell array

Scalogram coefficients, specified as a cell array of structure arrays. `u` is the output of `scatteringTransform` for the scattering decomposition framework `sf`.

x — Scattering feature matrix

real-valued matrix | real-valued array

Scattering feature matrix, specified as a real-valued 2-D matrix or 3-D array. `x` is the output of `featureMatrix` for the scattering decomposition framework `sf`.

Output Arguments

slog — Natural logarithm of scattering coefficients

cell array

Natural logarithm of scattering coefficients, returned as a cell array. The dimensions of `slog` are equal to the dimensions of `s`.

The precision of `slog` depends on the precision specified in the framework `sf`.

u'log — Natural logarithm of scalogram coefficients

cell array

Natural logarithm of scalogram coefficients, returned as a cell array. The dimensions of `u'log` are equal to the dimensions of `u`.

The precision of `u'log` depends on the precision specified in the framework `sf`.

xlog — Natural logarithm of scattering feature matrix

real-valued matrix | real-valued array

Natural logarithm of scattering feature matrix, returned as a real-valued matrix or array. The dimensions of `xlog` are equal to the dimensions of `x`.

The precision of `xlog` depends on the precision specified in the framework `sf`.

Algorithms

log returns the natural logarithm of the absolute value of the input argument.

See Also

scatteringTransform | waveletScattering

Introduced in R2018b

log

Natural logarithm of 2-D scattering transform

Syntax

```
slog = log(sf,s)  
ulog = log(sf,u)  
xlog = log(sf,x)
```

Description

`slog = log(sf,s)` returns the natural logarithm of the scattering coefficients in the cell array `s`. `s` is the output of `scatteringTransform` and is a cell array of structure arrays with an `images` field.

The precision of `slog` depends on the precision specified in the framework `sf`.

`ulog = log(sf,u)` returns the natural logarithm of the scalogram coefficients in the cell array `u`. `u` is the output of `scatteringTransform` and is a cell array of structure arrays with a `coefficients` field.

The precision of `ulog` depends on the precision specified in the framework `sf`.

`xlog = log(sf,x)` returns the natural logarithm of the 3-D matrix or 4-D tensor `x`. `x` is the output of `featureMatrix`.

The precision of `xlog` depends on the precision specified in the framework `sf`.

Examples

Natural Logarithm of Scattering Coefficients

This example shows how to obtain the natural logarithm of scattering coefficients.

Load the xbox image. Create an image scattering framework that can be applied to the image.

```
load xbox
sf = waveletScattering2('ImageSize',size(xbox),...
    'InvarianceScale',min(size(xbox)))
```

```
sf =
    waveletScattering2 with properties:
```

```
        ImageSize: [128 128]
    InvarianceScale: 128
        NumRotations: [6 6]
    QualityFactors: [1 1]
        Precision: "single"
    OversamplingFactor: 0
        OptimizePath: 1
```

Obtain the scattering transform of the image and then the natural logarithm of the scattering coefficients. Display the number of rows in the table containing the first-order scattering coefficients.

```
S = scatteringTransform(sf,xbox);
Slog = log(sf,S);
coefOrder = 1;
display(['Number of rows: ',num2str(size(S{coefOrder+1},1))])
```

```
Number of rows: 30
```

Choose a row from the first-order scattering coefficients table. Take the natural logarithm of the absolute value of the scattering coefficients in that row. Compare with the corresponding row in Slog and confirm they are equal.

```
row = 11;
tmp1 = Slog{coefOrder+1}.images{row};
tmp2 = log(abs(S{coefOrder+1}.images{row}));
disp(['Max Difference of Scattering Coefficients: '...
    num2str(max(abs(tmp1(:))-tmp2(:)))])
```

```
Max Difference of Scattering Coefficients: 0
```

Input Arguments

sf — Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

s — Scattering coefficients

cell array

Scattering coefficients, specified as a cell array of structure arrays. **s** is the output of scatteringTransform for the image scattering decomposition framework **sf**.

u — Scalogram coefficients

cell array

Scalogram coefficients, specified as a cell array of structure arrays. **u** is the output of scatteringTransform for the image scattering decomposition framework **sf**.

x — Scattering feature matrix

real-valued matrix | real-valued 4-D tensor

Scattering feature matrix, specified as a real-valued 3-D matrix or a real-valued 4-D tensor. **x** is the output of featureMatrix for the image scattering decomposition framework **sf**.

Output Arguments

slog — Natural logarithm of scattering coefficients

cell array

Natural logarithm of scattering coefficients, returned as a cell array. The dimensions of **slog** are equal to the dimensions of **s**.

The precision of **slog** depends on the precision specified in the framework **sf**.

u log — Natural logarithm of scalogram coefficients

cell array

Natural logarithm of scalogram coefficients, returned as a cell array. The dimensions of **u log** are equal to the dimensions of **u**.

The precision of `u$log` depends on the precision specified in the framework `sf`.

`x$log` – Natural logarithm of scattering feature matrix

real-valued 3-D matrix | real-valued 4-D tensor

Natural logarithm of scattering feature matrix, returned as a real-valued matrix or tensor. The dimensions of `x$log` are equal to the dimensions of `x`.

The precision of `x$log` depends on the precision specified in the framework `sf`.

Algorithms

`log` returns the natural logarithm of the absolute value of the input argument.

See Also

`featureMatrix` | `scatteringTransform` | `waveletScattering2`

Introduced in R2019a

ls2filt

Transform lifting scheme to quadruplet of filters

Syntax

```
[LoD,HiD,LoR,HiR] = ls2filt(LS)
```

Description

`[LoD,HiD,LoR,HiR] = ls2filt(LS)` returns the four filters LoD, HiD, LoR, and HiR associated with the lifting scheme LS.

Examples

```
% Start from the db2 wavelet and get the  
% corresponding lifting scheme.
```

```
LS = liftwave('db2')
```

```
LS =
```

```
    'd'          [ -1.7321]      [ 0]  
    'p'          [1x2 double]    [ 1]  
    'd'          [         1]    [-1]  
 [1.9319]      [  0.5176]      []
```

```
% Visualize the obtained lifting scheme.
```

```
displs(LS);
```

```
LS = {...  
    'd'          [ -1.73205081]      [0]  
    'p'          [ -0.06698730  0.43301270] [1]  
    'd'          [  1.00000000]    [-1]  
 [ 1.93185165] [  0.51763809]      []  
};
```

```
% Get the filters from the lifting scheme.
[LoD,HiD,LoR,HiR] = ls2filt(LS)

LoD =
    -0.1294    0.2241    0.8365    0.4830

HiD =
    -0.4830    0.8365   -0.2241   -0.1294

LoR =
    0.4830    0.8365    0.2241   -0.1294

HiR =
    -0.1294   -0.2241    0.8365   -0.4830

% Get the db2 filters using wfilters.
% You can check the equality.

[LoDref,HiDref,LoRref,HiRref] = wfilters('db2')

LoDref =
    -0.1294    0.2241    0.8365    0.4830

HiDref =
    -0.4830    0.8365   -0.2241   -0.1294

LoRref =
    0.4830    0.8365    0.2241   -0.1294

HiRref =
    -0.1294   -0.2241    0.8365   -0.4830
```

See Also

[filt2ls](#) | [lsinfo](#)

Introduced before R2006a

linfo

Lifting schemes information

Syntax

linfo

Description

linfo displays the following information about lifting schemes. A lifting scheme LS is a $N \times 3$ cell array. The $N-1$ first rows of the array are elementary lifting steps (ELS). The last row gives the normalization of LS.

Each ELS has this format:

```
{type, coefficients, max_degree}
```

where `type` is 'p' (primal) or 'd' (dual), `coefficients` is a vector C of real numbers defining the coefficients of a Laurent polynomial P described below, and `max_degree` is the highest degree d of the monomials of P .

The Laurent polynomial P is of the form

$$P(z) = C(1)*z^d + C(2)*z^{(d-1)} + \dots + C(m)*z^{(d-m+1)}$$

The lifting scheme LS is such that for

$k = 1:N-1$, $LS\{k, : \}$ is an ELS, where

$LS\{k, 1\}$ is the lifting type 'p' (primal) or 'd' (dual).

$LS\{k, 2\}$ is the corresponding lifting filter.

$LS\{k, 3\}$ is the highest degree of the Laurent polynomial corresponding to the filter $LS\{k, 2\}$.

$LS\{N, 1\}$ is the primal normalization (real number).

LS{N, 2} is the dual normalization (real number).

LS{N, 3} is not used.

Usually, the normalizations are such that $LS\{N, 1\} * LS\{N, 2\} = 1$.

For example, the lifting scheme associated with the wavelet db1 is

```
LS = { ...
      'd'      [ -1] [0]
      'p'      [0.5000] [0]
      [1.4142] [0.7071] []
    }
```

See Also

displs | laurpoly

Introduced before R2006a

lwt

1-D lifting wavelet transform

Syntax

```
[CA,CD] = lwt(X,W)
X_InPlace = lwt(X,W)
lwt(X,W,LEVEL)
X_InPlace = lwt(X,W,LEVEL,'typeDEC',typeDEC)
[CA,CD] = lwt(X,W,LEVEL,'typeDEC',typeDEC)
```

Description

`lwt` performs a 1-D lifting wavelet decomposition with respect to a particular lifted wavelet that you specify.

`[CA,CD] = lwt(X,W)` computes the approximation coefficients vector `CA` and detail coefficients vector `CD`, obtained by a lifting wavelet decomposition, of the vector `X`. `W` is a lifted wavelet name (see `liftwave`).

`X_InPlace = lwt(X,W)` computes the approximation and detail coefficients. These coefficients are stored in place:

`CA = X_InPlace(1:2:end)` and `CD = X_InPlace(2:2:end)`

`lwt(X,W,LEVEL)` computes the lifting wavelet decomposition at level `LEVEL`.

`X_InPlace = lwt(X,W,LEVEL,'typeDEC',typeDEC)` or `[CA,CD] = lwt(X,W,LEVEL,'typeDEC',typeDEC)` with `typeDEC = 'w'` or `'wp'` computes the wavelet or the wavelet packet decomposition using lifting, at level `LEVEL`.

Instead of a lifted wavelet name, you may use the associated lifting scheme `LS`: `lwt(X,LS,...)` instead of `lwt(X,W,...)`.

For more information about lifting schemes, see `lsinfo`.

Examples

```
% Start from the Haar wavelet and get the
% corresponding lifting scheme.
lshaar = liftwave('haar');

% Add a primal ELS to the lifting scheme.
els = {'p',[-0.125 0.125],0};
lsnew = addlift(lshaar,els);

% Perform LWT at level 1 of a simple signal.
x = 1:8;
[cA,cD] = lwt(x,lsnew)

cA =

    1.9445    4.9497    7.7782   10.6066

cD =

    0.7071    0.7071    0.7071    0.7071

% Perform integer LWT of the same signal.
lshaarInt = liftwave('haar','int2int');
lsnewInt = addlift(lshaarInt,els);
[cAint,cDint] = lwt(x,lsnewInt)

cAint =

     1     3     5     7

cDint =

     1     1     1     1
```

Algorithms

This function uses the polyphase algorithm.

`lwt` reduces to `dwt` with zero-padding extension mode and without extra-coefficients.

References

Strang, G.; T. Nguyen (1996), *Wavelets and filter banks*, Wellesley-Cambridge Press.

Sweldens, W. (1998), "The Lifting Scheme: a Construction of Second Generation of Wavelets," *SIAM J. Math. Anal.*, 29 (2), pp. 511-546.

See Also

ilwt

Introduced before R2006a

lwt2

2-D lifting wavelet transform

Syntax

```
[CA,CH,CV,CD] = lwt2(X,W)
X_InPlace = lwt2(X,LS)
lwt2(X,W,LEVEL)
X_InPlace = lwt2(X,W,LEVEL,'typeDEC',typeDEC)
[CA,CH,CV,CD] = LWT2(X,W,LEVEL,'typeDEC',typeDEC)
```

Description

`lwt2` performs a 2-D lifting wavelet decomposition with respect to a particular lifted wavelet that you specify.

`[CA,CH,CV,CD] = lwt2(X,W)` computes the approximation coefficients matrix `CA` and detail coefficients matrices `CH`, `CV`, and `CD`, obtained by a lifting wavelet decomposition, of the matrix `X`. `W` is a lifted wavelet name (see `liftwave`).

`X_InPlace = lwt2(X,LS)` computes the approximation and detail coefficients. These coefficients are stored in place:

- `CA = X_InPlace(1:2:end,1:2:end)`
- `CH = X_InPlace(2:2:end,1:2:end)`
- `CV = X_InPlace(1:2:end,2:2:end)`
- `CD = X_InPlace(2:2:end,2:2:end)`

`lwt2(X,W,LEVEL)` computes the lifting wavelet decomposition at level `LEVEL`.

`X_InPlace = lwt2(X,W,LEVEL,'typeDEC',typeDEC)` or `[CA,CH,CV,CD] = LWT2(X,W,LEVEL,'typeDEC',typeDEC)` with `typeDEC = 'w'` or `'wp'` computes the wavelet or the wavelet packet decomposition using lifting, at level `LEVEL`.

Instead of a lifted wavelet name, you may use the associated lifting scheme `LS`: `lwt2(X,LS,...)` instead of `LWT2(X,W,...)`.

For more information about lifting schemes, see `lsinfo`.

Examples

```
% Start from the Haar wavelet and get the
% corresponding lifting scheme.
lshaar = liftwave('haar');

% Add a primal ELS to the lifting scheme.
els = {'p',[-0.125 0.125],0};
lsnew = addlift(lshaar,els);

% Perform LWT at level 1 of a simple image.
x = reshape(1:16,4,4);
[cA,cH,cV,cD] = lwt2(x,lsnew)

cA =

    5.7500    22.7500
   10.0000    27.0000

cH =

    1.0000    1.0000
    1.0000    1.0000

cV =

    4.0000    4.0000
    4.0000    4.0000

cD =

    0     0
    0     0

% Perform integer LWT of the same image.
lshaarInt = liftwave('haar','int2int');
lsnewInt = addlift(lshaarInt,els);
[cAint,cHint,cVint,cDint] = lwt2(x,lsnewInt)
```

```
cAint =
```

```
    3    11  
    5    13
```

```
cHint =
```

```
    1    1  
    1    1
```

```
cVint =
```

```
    4    4  
    4    4
```

```
cDint =
```

```
    0    0  
    0    0
```

Tips

When X represents an indexed image, X , as well as the output arrays `cA`, `cH`, `cV`, `cD`, or `X_InPlace` are m -by- n matrices. When X represents a truecolor image, it is an m -by- n -by-3 array, where each m -by- n matrix represents a red, green, or blue color plane concatenated along the third dimension.

For more information on image formats, see the `image` and `imfinfo` reference pages .

Algorithms

This function implements the polyphase algorithm.

`lwt` reduces to `dwt` with zero-padding extension mode and without extra-coefficients.

References

Strang, G.; T. Nguyen (1996), *Wavelets and filter banks*, Wellesley-Cambridge Press.

Sweldens, W. (1998), "The Lifting Scheme: a Construction of Second Generation of Wavelets," *SIAM J. Math. Anal.*, 29 (2), pp. 511-546.

See Also

ilwt2

Introduced before R2006a

lwtcoef

Extract or reconstruct 1-D LWT wavelet coefficients

Syntax

```
Y = lwtcoef(TYPE,XDEC,LS,LEVEL,LEVEXT)  
Y = lwtcoef(TYPE,XDEC,W,LEVEL,LEVEXT)
```

Description

`Y = lwtcoef(TYPE,XDEC,LS,LEVEL,LEVEXT)` returns the coefficients or the reconstructed coefficients of level `LEVEXT`, extracted from `XDEC`, the LWT decomposition at level `LEVEL` obtained with the lifting scheme `LS`.

The valid values for `TYPE` are

TYPE Values	Description
'a'	Approximations
'd'	Details
'ca'	Coefficients of approximations
'cd'	Coefficients of details

`Y = lwtcoef(TYPE,XDEC,W,LEVEL,LEVEXT)` returns the same output using `W`, which is the name of a lifted wavelet.

Examples

```
% Start from the Haar wavelet and get the  
% corresponding lifting scheme.  
lshaar = liftwave('haar');  
  
% Add a primal ELS to the lifting scheme.  
els = {'p',[-0.125 0.125],0};
```



```
lsnew = addlift(lshaar,els);

% Perform LWT at level 2 of a simple signal.
x = 1:8;
xDec = lwt(x,lsnew,2)

xDec =

    4.3438    0.7071    2.1250    0.7071   13.0313    0.7071
    2.0000    0.7071

% Extract approximation coefficients of level 1.
ca1 = lwtcoef('ca',xDec,lsnew,2,1)

ca1 =

    1.9445    4.9497    7.7782   10.6066

% Reconstruct approximations and details.
a1 = lwtcoef('a',xDec,lsnew,2,1)

a1 =

    1.3750    1.3750    3.5000    3.5000    5.5000    5.5000
    7.5000    7.5000

a2 = lwtcoef('a',xDec,lsnew,2,2)

a2 =

    2.1719    2.1719    2.1719    2.1719    6.5156    6.5156
    6.5156    6.5156

d1 = lwtcoef('d',xDec,lsnew,2,1)

d1 =

   -0.3750    0.6250   -0.5000    0.5000   -0.5000    0.5000
   -0.5000    0.5000

d2 = lwtcoef('d',xDec,lsnew,2,2)

d2 =
```

```
-0.7969  -0.7969  1.3281  1.3281  -1.0156  -1.0156  
0.9844  0.9844
```

```
% Check perfect reconstruction.  
err = max(abs(x-a2-d2-d1))
```

```
err =
```

```
9.9920e-016
```

See Also

`ilwt` | `lwt`

Introduced before R2006a

lwtcoef2

Extract or reconstruct 2-D LWT wavelet coefficients

Syntax

```
Y = lwtcoef2(TYPE,XDEC,LS,LEVEL,LEVEXT)
Y = lwtcoef2(TYPE,XDEC,W,LEVEL,LEVEXT)
```

Description

`Y = lwtcoef2(TYPE,XDEC,LS,LEVEL,LEVEXT)` returns the coefficients or the reconstructed coefficients of level `LEVEXT`, extracted from `XDEC`, the LWT decomposition at level `LEVEL` obtained with the lifting scheme `LS`.

The valid values for `TYPE` are listed in this table.

TYPE Values	Description
'a'	Approximations
'h'	Horizontal details
'v'	Vertical details
'd'	Diagonal details
'ca'	Coefficients of approximations
'ch'	Coefficients of horizontal details
'cv'	Coefficients of vertical details
'cd'	Coefficients of diagonal details

`Y = lwtcoef2(TYPE,XDEC,W,LEVEL,LEVEXT)` returns the same output using `W`, which is the name of a lifted wavelet.

Examples

```
% Start from the Haar wavelet and get the
% corresponding lifting scheme.
lshaar = liftwave('haar');

% Add a primal ELS to the lifting scheme.
els = {'p',[-0.125 0.125],0};
lsnew = addlift(lshaar,els);

% Perform LWT at level 2 of a simple image.
x = reshape(1:16,4,4);
xDec = lwt2(x,lsnew,2)

xDec =

    27.4375    4.0000    17.0000    4.0000
     1.0000         0     1.0000         0
     4.2500    4.0000    0.0000    4.0000
     1.0000         0     1.0000         0

% Extract approximation coefficients of level 1.
cal = lwtcoef2('ca',xDec,lsnew,2,1)

cal =

     5.7500    22.7500
    10.0000    27.0000

% Reconstruct approximations and details.
a1 = lwtcoef2('a',xDec,lsnew,2,1)

a1 =

     2.8750     2.8750    11.3750    11.3750
     2.8750     2.8750    11.3750    11.3750
     5.0000     5.0000    13.5000    13.5000
     5.0000     5.0000    13.5000    13.5000

a2 = lwtcoef2('a',xDec,lsnew,2,2)

a2 =

     6.8594     6.8594     6.8594     6.8594
```

```
6.8594    6.8594    6.8594    6.8594
6.8594    6.8594    6.8594    6.8594
6.8594    6.8594    6.8594    6.8594
```

```
h1 = lwtcoef2('h',xDec,lsnew,2,1)
```

```
h1 =
```

```
-0.3750   -0.3750   -0.3750   -0.3750
 0.6250    0.6250    0.6250    0.6250
-0.5000   -0.5000   -0.5000   -0.5000
 0.5000    0.5000    0.5000    0.5000
```

```
v1 = lwtcoef2('v',xDec,lsnew,2,1)
```

```
v1 =
```

```
-1.5000    2.5000   -2.0000    2.0000
-1.5000    2.5000   -2.0000    2.0000
-1.5000    2.5000   -2.0000    2.0000
-1.5000    2.5000   -2.0000    2.0000
```

```
d1 = lwtcoef2('d',xDec,lsnew,2,1)
```

```
d1 =
```

```
0    0    0    0
0    0    0    0
0    0    0    0
0    0    0    0
```

```
h2 = lwtcoef2('h',xDec,lsnew,2,2)
```

```
h2 =
```

```
-0.7969   -0.7969   -0.7969   -0.7969
-0.7969   -0.7969   -0.7969   -0.7969
 1.3281    1.3281    1.3281    1.3281
 1.3281    1.3281    1.3281    1.3281
```

```
v2 = lwtcoef2('v',xDec,lsnew,2,2)
```

```
v2 =
```

```
-3.1875  -3.1875   5.3125   5.3125
-3.1875  -3.1875   5.3125   5.3125
-3.1875  -3.1875   5.3125   5.3125
-3.1875  -3.1875   5.3125   5.3125

d2 = lwtcoef2('d',xDec,lsnew,2,2)

d2 =

    1.0e-015 *

    0.2498    0.2498   -0.4163   -0.4163
    0.2498    0.2498   -0.4163   -0.4163
   -0.4163   -0.4163    0.6939    0.6939
   -0.4163   -0.4163    0.6939    0.6939

% Check perfect reconstruction.
err = max(max(abs(x-a2-h2-v2-d2-h1-v1-d1)))

err =

    3.5527e-015
```

Tips

If XDEC is obtained from an indexed image analysis or a truecolor image analysis, it is an m-by-n matrix or an m-by-n-by-3 array, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

See Also

`ilwt2` | `lwt2`

Introduced before R2006a

mdwtcluster

Multisignals 1-D clustering

Syntax

```
S = mdwtcluster(X)
S = mdwtcluster(X, 'PropName1', PropVal1, 'PropName2', PropVal2, ...)
```

Description

`S = mdwtcluster(X)` constructs clusters from a hierarchical cluster tree. The input matrix X is decomposed in row direction using the DWT function with the haar wavelet and the maximum allowed level.

`S = mdwtcluster(X, 'PropName1', PropVal1, 'PropName2', PropVal2, ...)` allows you to modify some properties. The valid choices for *PropName* are:

Note `mdwtcluster` requires the Statistics and Machine Learning Toolbox™

'dirDec'	'r' (row) or 'c' (column). Default value is 'r'.
'level'	Level of the DWT decomposition. Default value is: <code>level=fix(log2(size(X,d)))</code> where $d=1$ or $d=2$, depending on the <code>dirDec</code> value.
'wname'	Wavelet name used for DWT. Default value is 'haar'.
'dwtEXTM'	DWT extension mode (see <code>dwtmode</code>).
'pdist'	See Statistics and Machine Learning Toolbox <code>pdist</code> function. Default value is 'euclidean'.
'linkage'	See Statistics and Machine Learning Toolbox <code>linkage</code> function. Default value is 'ward'.
'maxclust'	Number of clusters. Default value is 6. The input variable can be a vector.

'lst2clu'	<p>Cell array that contains the list of data to classify.</p> <p>If N is the level of decomposition, the allowed name values for the cells are:</p> <ul style="list-style-type: none"> • 's' — Signal • 'aj' — Approximation at level j • 'dj' — Detail at level j • 'caj' — Coefficients of approximation at level j • 'cdj' — Coefficients of detail at level j <p>Default value is {'s'; 'ca1'; ...; 'caN'}.</p>
-----------	--

The output structure S is such that for each partition j:

S.Idx(:,j)	Contains the cluster numbers obtained from the hierarchical cluster tree (see <code>cluster</code> in the Statistics and Machine Learning Toolbox software).
S.Incons(:,j)	Contains the inconsistent values of each non-leaf node in the hierarchical cluster tree (see Statistics and Machine Learning Toolbox software function <code>inconsistent</code>).
S.Corr(j)	Contains the cophenetic correlation coefficients of the partition (see Statistics and Machine Learning Toolbox software function <code>cophenet</code>).

Note If `maxclustVal` is a vector, then `IdxCLU` is a multidimensional array such that `IdxCLU(:,j,k)` contains the cluster numbers obtained from the hierarchical cluster tree for k clusters.

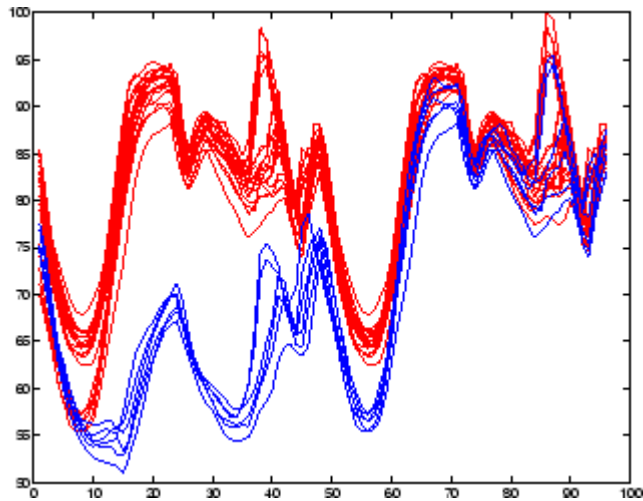
Examples

```
load elecsig10
lst2clu = {'s','ca1','ca3','ca6'};

% Compute the structure resulting from multisignal clustering
S = mdwtcluster(signals,'maxclust',4,'lst2clu',lst2clu)
```



```
S =  
  
    IdxCLU: [70x4 double]  
    Incons: [69x4 double]  
    Corr: [0.7920 0.7926 0.7947 0.7631]  
  
% Retrieve indices of clusters  
IdxCLU = S.IdxCLU;  
  
% Plot the first cluster  
plot(signals(IdxCLU(:,1)==1,:),'r');  
hold on;  
  
% Plot the third clustering  
plot(signals(IdxCLU(:,1)==3,:),'b')
```



```
% Check the equality of partitions  
equalPART = isequal(IdxCLU(:,1),IdxCLU(:,3))  
  
equalPART =  
  
    1
```

```
% So we can see that we obtain the same partitions using
```

% coefficients of approximation at level 3 instead of original
% signals. Much less information is then used.

See Also

mdwtdec | wavedec

Introduced in R2008a

mdwtdec

Multisignal 1-D wavelet decomposition

Syntax

```
DEC = mdwtdec(DIRDEC,X,LEV,WNAME)
DEC = mdwtdec(DIRDEC,X,LEV,LoD,HiD,LoR,HiR)
DEC = mdwtdec(...,'mode',EXTMODE)
```

Description

DEC = mdwtdec(DIRDEC,X,LEV,WNAME) returns the wavelet decomposition at level LEV of each row (if DIRDEC = 'r') or each column (if DIRDEC = 'c') of matrix X, using the wavelet WNAME.

The output DEC is a structure with the following fields:

'dirDec'	Direction indicator: 'r' (row) or 'c' (column)
'level'	Level of the DWT decomposition
'wname'	Wavelet name
'dwtFilters'	Structure with four fields LoD, HiD, LoR and HiR
'dwtEXTM'	DWT extension mode (see dwtmode)
'dwtShift'	DWT shift parameter (0 or 1)
'dataSize'	Size of X
'ca'	Approximation coefficients at level LEV
'cd'	Cell array of detail coefficients, from level 1 to level LEV

Coefficients cA and cD{k} (for k = 1 to LEV) are matrices and are stored in rows if DIRDEC = 'r' or in columns if DIRDEC = 'c'.

DEC = mdwtdec(DIRDEC,X,LEV,LoD,HiD,LoR,HiR) uses the four filters instead of the wavelet name.

`DEC = mdwtdec(..., 'mode', EXTMODE)` computes the wavelet decomposition with the EXTMODE extension mode that you specify (see `dwtmode` for the valid extension modes).

Examples

Decompose Multisignals

This example shows how to return the wavelet decomposition of a multisignal using a wavelet name and wavelet filters.

Load a 1-D multisignal.

```
load thinker
```

Perform a decomposition at level 2 using the db2 wavelet.

```
dec = mdwtdec('r', X, 2, 'db2')  
  
dec = struct with fields:  
    dirDec: 'r'  
    level: 2  
    wname: 'db2'  
    dwtFilters: [1x1 struct]  
    dwtEXTM: 'per'  
    dwtShift: 0  
    dataSize: [192 96]  
    ca: [192x24 double]  
    cd: {[192x48 double] [192x24 double]}
```

Compute the filters associated with the db2 wavelet.

```
[LoD, HiD, LoR, HiR] = wfilters('db2');
```

Perform a decomposition at level 2 using the filters.

```
decBIS = mdwtdec('r', X, 2, LoD, HiD, LoR, HiR)
```

```
decBIS = struct with fields:  
    dirDec: 'r'  
    level: 2  
    wname: ''
```

```

dwtFilters: [1x1 struct]
  dwtEXTM: 'per'
  dwtShift: 0
  dataSize: [192 96]
    ca: [192x24 double]
    cd: {[192x48 double] [192x24 double]}

```

Confirm the approximation and detail coefficients of both decompositions are identical.

```
max(abs(dec.ca(:)-decBIS.ca(:)))
```

```
ans = 0
```

```
max(abs(dec.cd{1}(:)-decBIS.cd{1}(:)))
```

```
ans = 0
```

```
max(abs(dec.cd{2}(:)-decBIS.cd{2}(:)))
```

```
ans = 0
```

References

Daubechies, I. , *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed., 1992.

Mallat, S., "A theory for multiresolution signal decomposition: the wavelet representation," *IEEE Pattern Anal. and Machine Intell.*, vol. 11, no. 7, 1989, pp. 674-693.

Meyer, Y. , *Ondelettes et opérateurs*, Tome 1, Hermann Ed. (English translation: *Wavelets and operators*, Cambridge Univ. Press. 1993.)

See Also

mdwtrec | wavedec

Introduced in R2007a

mdwtrec

Multisignal 1-D wavelet reconstruction

Syntax

```
X = mdwtrec(DEC)
X = mdwtrec(DEC,IDXSIG)
Y = mdwtrec(DEC,TYPE,LEV)
A = mdwtrec(DEC,'a')
A = mdwtrec(DEC,'a',LEVDEC)
D = mdwtrec(DEC,'d')
CA = mdwtrec(DEC,'ca')
CA = mdwtrec(DEC,'ca',LEVDEC)
CD = mdwtrec(DEC,'cd',MODE)
CFS = mdwtrec(DEC,'cfs',MODE)
Y = mdwtrec(...,IDXSIG)
```

Description

`X = mdwtrec(DEC)` returns the original matrix of signals, starting from the wavelet decomposition structure `DEC` (see `mdwtdec`).

`X = mdwtrec(DEC,IDXSIG)` reconstructs the signals whose indices are given by the vector `IDXSIG`.

`Y = mdwtrec(DEC,TYPE,LEV)` extracts or reconstructs the detail or approximation coefficients at level `LEV` depending on the `TYPE` value. The maximum value for `LEV` is `LEVDEC = DEC.level`.

When `TYPE` is equal to:

- `'cd'` or `'ca'`, coefficients of level `LEV` are extracted.
- `'d'` or `'a'`, coefficients of level `LEV` are reconstructed.
- `'a'` or `'ca'`, `LEV` must be such that $0 \leq \text{LEV} \leq \text{LEVDEC}$.

- 'd' or 'cd', LEV must be such that $1 \leq \text{LEV} \leq \text{LEVDEC}$.

$A = \text{mdwtrec}(\text{DEC}, 'a')$ is equivalent to $A = \text{mdwtrec}(\text{DEC}, 'a', \text{LEVDEC})$.

$D = \text{mdwtrec}(\text{DEC}, 'd')$ returns a matrix containing the sum of all the details, so that $X = A + D$.

$CA = \text{mdwtrec}(\text{DEC}, 'ca')$ is equivalent to $CA = \text{mdwtrec}(\text{DEC}, 'ca', \text{LEVDEC})$.

$CD = \text{mdwtrec}(\text{DEC}, 'cd', \text{MODE})$ returns a matrix containing all the detail coefficients.

$CFS = \text{mdwtrec}(\text{DEC}, 'cfs', \text{MODE})$ returns a matrix containing all the coefficients.

For $\text{MODE} = 'descend'$ the coefficients are concatenated from level LEVDEC to level 1 and $\text{MODE} = 'ascend'$ concatenates from level 1 to level LEVDEC). The default is $\text{MODE} = 'descend'$. The concatenation is made row-wise if $\text{DEC.dirDEC} = 'r'$ or column-wise if $\text{DEC.dirDEC} = 'c'$.

$Y = \text{mdwtrec}(\dots, \text{IDXSIG})$ extracts or reconstructs the detail or the approximation coefficients for the signals whose indices are given by the vector IDXSIG .

Examples

Reconstruct Multisignals

This example shows how to reconstruct a multisignal and particular signal within the multisignal.

Load a 1-D multisignal.

```
load thinker
```

Perform a decomposition at level 2 using the **db2** wavelet.

```
dec = mdwtdec('r', X, 2, 'db2');
```

Reconstruct the original matrix of signals, using the decomposition structure **dec**.

```
XR = mdwtrec(dec);
```

Compute the reconstruction error.

```
errREC = max(abs(X(:)-XR(:)))
```

```
errREC = 2.1026e-10
```

Reconstruct the original signal at index 31, the corresponding approximation at level 2, and details at levels 1 and 2.

```
idx = 31;  
Y = mdwtrec(dec,idx);  
A2 = mdwtrec(dec,'a',2,idx);  
D2 = mdwtrec(dec,'d',2,idx);  
D1 = mdwtrec(dec,'d',1,idx);
```

Compute the reconstruction error for signal 31.

```
errREC = max(abs(Y-A2-D2-D1))
```

```
errREC = 6.7946e-14
```

References

Daubechies, I., *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed., 1992.

Mallat, S., "A theory for multiresolution signal decomposition: the wavelet representation," *IEEE Pattern Anal. and Machine Intell.*, vol. 11, no. 7, 1989, pp. 674-693.

Meyer, Y., *Ondelettes et opérateurs*, Tome 1, Hermann Ed. (English translation: *Wavelets and operators*, Cambridge Univ. Press. 1993.)

See Also

mdwtdec | waverec

Introduced in R2007a

measerr

Quality metrics of signal or image approximation

Syntax

```
[PSNR,MSE,MAXERR,L2RAT] = measerr(X,XAPP)
[PSNR,MSE,MAXERR,L2RAT] = measerr(X,XAPP,BPS)
```

Description

`[PSNR,MSE,MAXERR,L2RAT] = measerr(X,XAPP)` returns the peak signal-to-noise ratio, PSNR, mean square error, MSE, maximum squared error, MAXERR, and ratio of squared norms, L2RAT, for an input signal or image, X, and its approximation, XAPP.

`[PSNR,MSE,MAXERR,L2RAT] = measerr(X,XAPP,BPS)` uses the bits per sample, BPS, to determine the peak signal-to-noise ratio.

Examples

Measure Approximation Quality in RGB Image

Approximate an RGB image and compute the quality metrics.

Load an RGB image. Return the image dimensions and minimum and maximum values.

```
X = imread('afrिकासculpt.jpg');
size(X)
```

```
ans = 1×3
```

```
    512    512     3
```

```
[min(X(:)) max(X(:))]
```

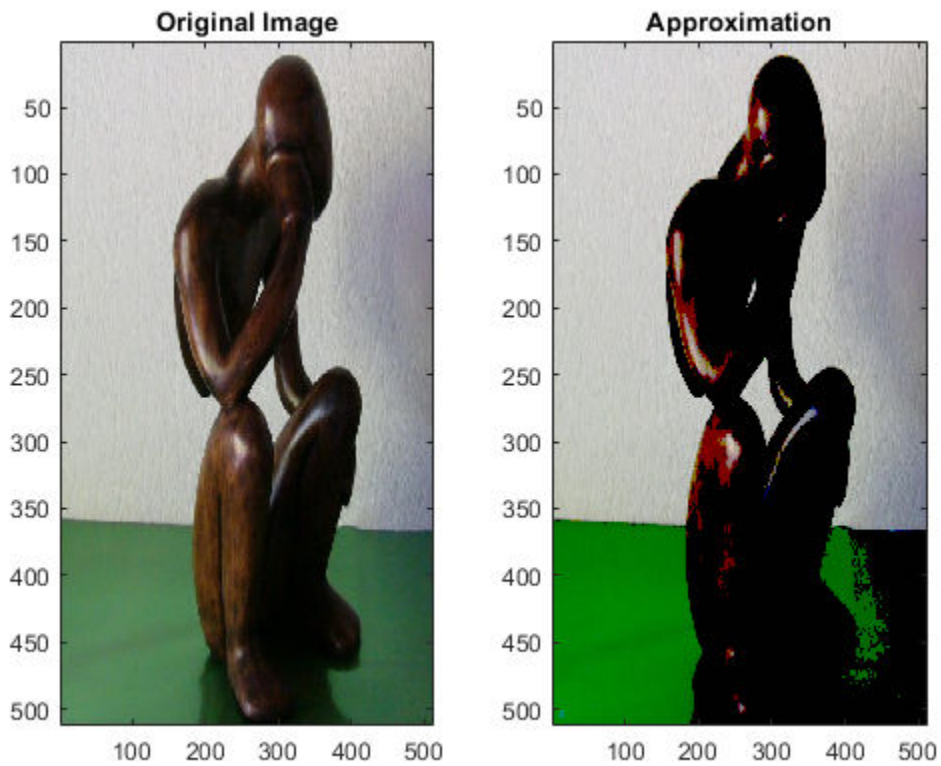
```
ans = 1x2 uint8 row vector  
    0   236
```

Define the image approximation by setting equal to 1 all RGB values less than or equal to 100.

```
Xapp = X;  
Xapp(X<=100) = 1;
```

Display the image and its approximation.

```
subplot(1,2,1)  
image(X)  
title('Original Image')  
subplot(1,2,2)  
image(Xapp)  
title('Approximation')
```



Compute the quality metrics of the image approximation.

```
[psnr,mse,maxerr,L2rat] = measerr(X,Xapp)
```

```
psnr = 17.5287
```

```
mse = 1.1487e+03
```

```
maxerr = 99
```

```
L2rat = 0.9398
```

Measure Approximation Quality in Grayscale Image

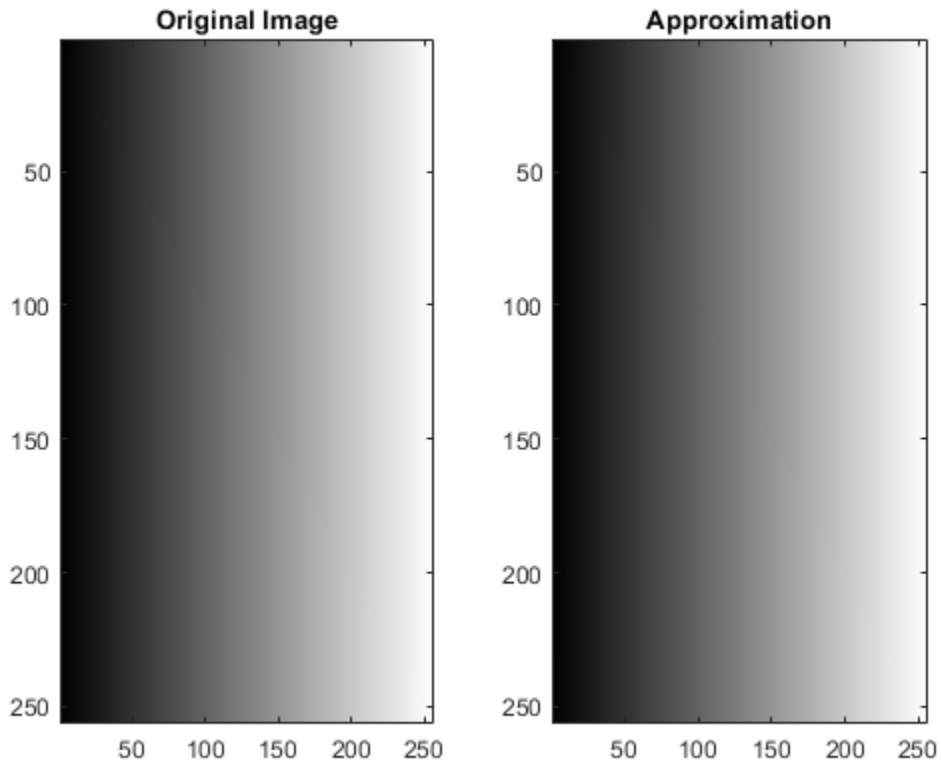
Approximate a grayscale image and calculate approximation quality metrics.

Create a 256-by-256 grayscale image with intensities between 0 and $2^{16} - 1$.

```
val = 0:2^16-1;  
X = reshape(val,256,256);
```

There are 16 bits per sample. Define the image approximation by setting equal to 1 all grayscale values less than or equal to 1000. Display the image and its approximation.

```
Xapp = X;  
Xapp(X<=1000) = 1;  
colormap(gray(2^16))  
subplot(1,2,1)  
image(X)  
title('Original Image')  
subplot(1,2,2)  
image(Xapp)  
title('Approximation')
```



There are 16 bits per sample. Compute the quality metrics of the grayscale approximation.

```
bps = 16;  
[psnr,mse,maxerr,L2rat] = measerr(X,Xapp)  
  
psnr = 11.0733  
mse = 5.0786e+03  
maxerr = 999  
L2rat = 1.0000
```

Input Arguments

X — Input signal or image

real-valued array

Input signal or image, specified as a real-valued array.

XAPP — Approximation of signal or image

real-valued array

Approximation of signal or image X , specified as a real-valued array. $XAPP$ is the same size as X .

BPS — Bits per sample

8 (default) | positive integer

Bits per sample of the input data, specified as a positive integer. The default value is 8, so the maximum possible pixel value of an image (MAXI) is 255. More generally, when samples are represented using linear Pulse Code Modulation with B bits per sample, MAXI is $2^B - 1$.

Output Arguments

PSNR — Peak signal-to-noise ratio

positive real number

Peak signal-to-noise ratio (PSNR) in decibels, returned as a positive real number. The PSNR is only meaningful for data encoded in terms of bits per sample or bits per pixel. For example, an image with 8 bits per pixel contains integers from 0 to 255.

MSE — Mean square error

positive real number

Mean square error, returned as a positive real number. MSE is the squared norm of the difference between X and $XAPP$ divided by the number of elements.

MAXERR — Maximum absolute squared deviation

positive real number

Maximum absolute squared deviation of the data X from the approximation $XAPP$, returned as a positive real number.

L2RAT — Energy ratio

positive real number

Energy ratio between the approximation XAPP and input data X, returned as a positive real number. L2RAT is the ratio of the squared norm of XAPP to X.

Definitions

Peak Signal to Noise Ratio

The peak signal-to-noise ratio (PSNR) in decibels between a signal and its approximation is

$$20\log_{10}\left(\frac{2^B - 1}{\sqrt{MSE}}\right)$$

where MSE represents the mean square error, and B represents the bits per sample.

Mean Square Error

The mean square error (MSE) between a signal or image, X , and an approximation, Y , is

$$\frac{\|X - Y\|^2}{N}$$

where N is the number of elements in the signal.

References

- [1] Huynh-Thu, Q. and M. Ghanbari. "Scope of Validity of PSNR in Image/Video Quality Assessment." *Electronics Letters*. Vol. 44, Issue 13, 2008, pp. 800-801.

See Also

wden | wdencomp | wdenoise

Topics

“Wavelet Data Compression”

“Wavelet Denoising and Nonparametric Function Estimation”

Introduced in R2010b

mexihat

Mexican hat (Ricker) wavelet

Syntax

```
[PSI,X] = mexihat(LB,UB,N)
```

Description

`[PSI,X] = mexihat(LB,UB,N)` returns values of the Mexican hat wavelet on an N point regular grid, X, in the interval [LB,UB]. The Mexican hat wavelet is also known as the Ricker wavelet.

Output arguments are the wavelet function PSI computed on the grid X.

This wavelet has [-5 5] as effective support. Although [-5 5] is the correct theoretical effective support, a wider effective support, [-8 8], is used in the computation to provide more accurate results.

This function is proportional to the second derivative function of the Gaussian probability density function.

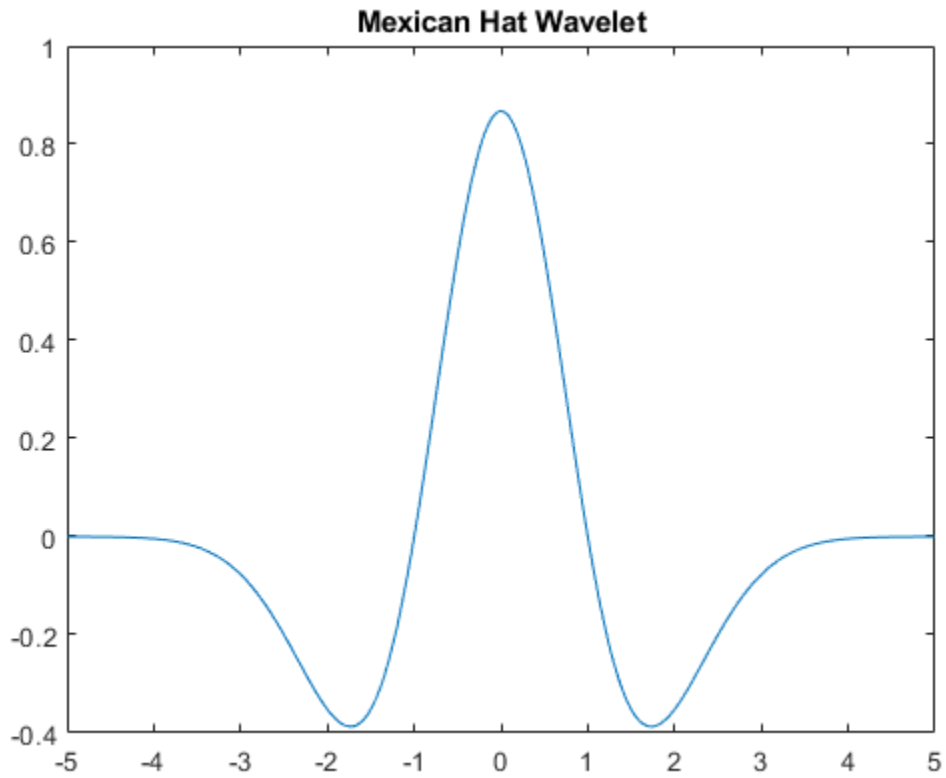
Note You can use `gauswavf` to obtain a second order derivative of a Gaussian wavelet. If you use the negative of this normalized derivative, the resulting wavelet resembles the Mexican hat wavelet.

Examples

Mexican Hat Wavelet

Create a Mexican hat wavelet with support on [-5,5]. Use 1,000 sample points. Plot the result.

```
lb = -5;  
ub = 5;  
N = 1000;  
[psi,xval] = mexihat(lb,ub,N);  
plot(xval,psi)  
title('Mexican Hat Wavelet');
```



See Also

waveinfo

Introduced before R2006a

meyer

Meyer wavelet

Syntax

```
[PHI,PSI,T] = meyer(LB,UB,N)
```

Description

`[PHI,PSI,T] = meyer(LB,UB,N)` returns Meyer scaling and wavelet functions evaluated on an N point regular grid in the interval $[LB,UB]$.

N must be a power of two.

Output arguments are the scaling function `PHI` and the wavelet function `PSI` computed on the grid T . These functions have $[-8,8]$ as effective support.

If only one function is required, a fourth argument is allowed:

```
[PHI,T] = meyer(LB,UB,N,'phi')  
[PSI,T] = meyer(LB,UB,N,'psi')
```

When the fourth argument is used, but not equal to `'phi'` or `'psi'`, outputs are the same as in the main option.

The Meyer wavelet and scaling function are defined in the frequency domain.

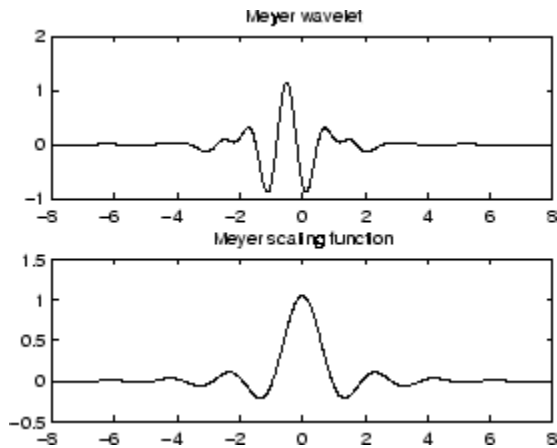
By changing the auxiliary function (see `meyeraux` for more information), you get a family of different wavelets.

Examples

```
% Set effective support and grid parameters.  
lb = -8; ub = 8; n = 1024;
```

```
% Compute and plot Meyer wavelet and scaling functions.
```

```
[phi,psi,x] = meyer(lb,ub,n);
subplot(211), plot(x,psi)
title('Meyer wavelet')
subplot(212), plot(x,phi)
title('Meyer scaling function')
```



Algorithms

Starting from an explicit form of the Fourier transform $\widehat{\phi}$ of ϕ , `meyer` computes the values of $\widehat{\phi}$ on a regular grid, and then the values of ϕ are computed using `instdfft`, the inverse nonstandard discrete FFT.

The procedure for ψ is along the same lines.

References

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics, SIAM Ed., pp. 117-119, 137, 152.

See Also

`meyeraux` | `wavefun` | `waveinfo`

Introduced before R2006a

meyeraux

Meyer wavelet auxiliary function

Syntax

`Y = meyeraux(X)`

Description

`Y = meyeraux(X)` returns values of the auxiliary function used for Meyer wavelet generation evaluated at the elements of the vector or matrix X .

The function is

$$35x^4 - 84x^5 + 70x^6 - 20x^7$$

See Also

`meyer`

Introduced before R2006a

mlpt

Multiscale local 1-D polynomial transform

Syntax

```
[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x,t)
[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x,t,numLevel)
[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x)
[coefs,T,coefsPerLevel,scalingMoments] = mlpt( ____,Name,Value)
```

Description

`[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x,t)` returns the multiscale local polynomial 1-D transform (MLPT) of input signal `x` sampled at the sampling instants, `t`. If `x` or `t` contain NaNs, the union of the NaNs in `x` and `t` is removed before obtaining the `mlpt`.

`[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x,t,numLevel)` returns the transform for `numLevel` resolution levels.

`[coefs,T,coefsPerLevel,scalingMoments] = mlpt(x)` uses uniform sampling instants for `x` as the time instants if `x` does not contain NaNs. If `x` contains NaNs, the NaNs are removed from `x` and the nonuniform sampling instants are obtained from the numeric elements of `x`.

`[coefs,T,coefsPerLevel,scalingMoments] = mlpt(____,Name,Value)` specifies `mlpt` properties using one or more `Name,Value` pair arguments and any of the previous input arguments.

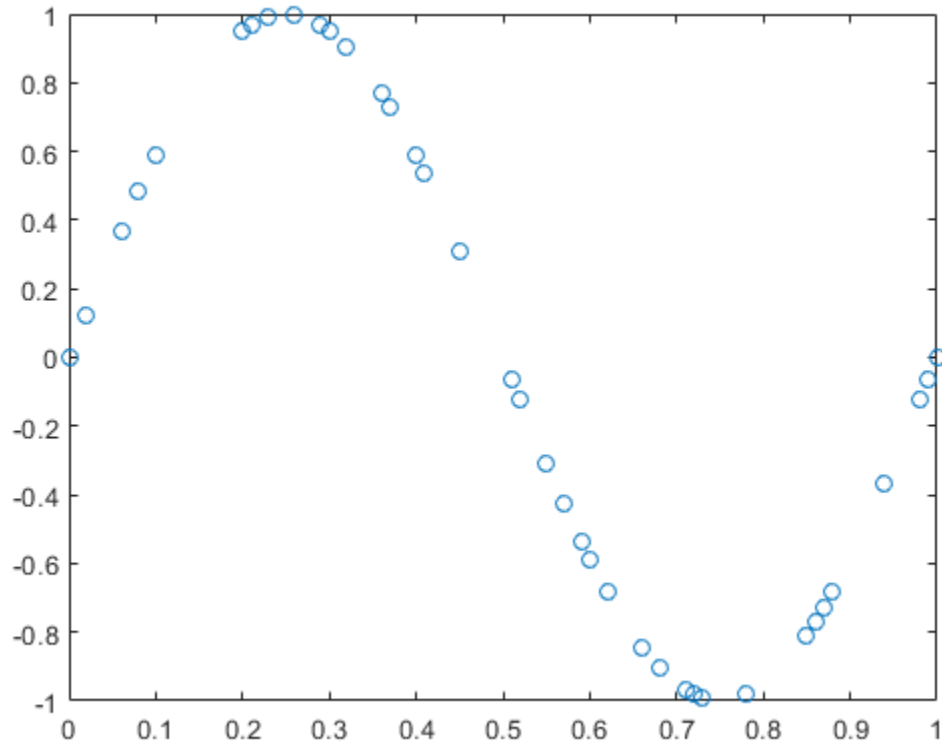
Examples

Multiscale Local 1-D Polynomial Transform and Inverse Transform

Create a signal with nonuniform sampling and verify good reconstruction when performing the mlpt and imlpt.

Create and plot a sine wave with non-uniform sampling.

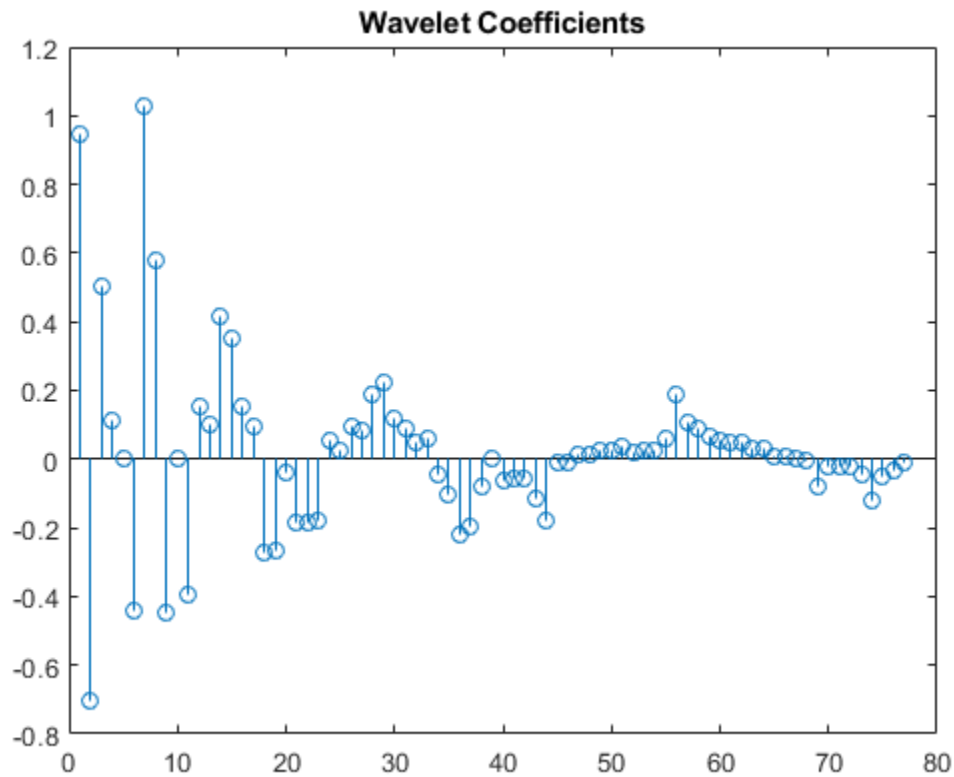
```
timeVector = 0:0.01:1;  
sineWave = sin(2*pi*timeVector)';  
  
samplesToErase = randi(100,100,1);  
sineWave(samplesToErase) = [];  
timeVector(samplesToErase) = [];  
  
figure(1)  
plot(timeVector,sineWave,'o')  
hold on
```



Perform the multiscale local 1-D polynomial transform (`mlpt`) on the signal. Visualize the coefficients.

```
[coefs,T,coefsPerLevel,scalingMoments] = mlpt(sineWave,timeVector);
```

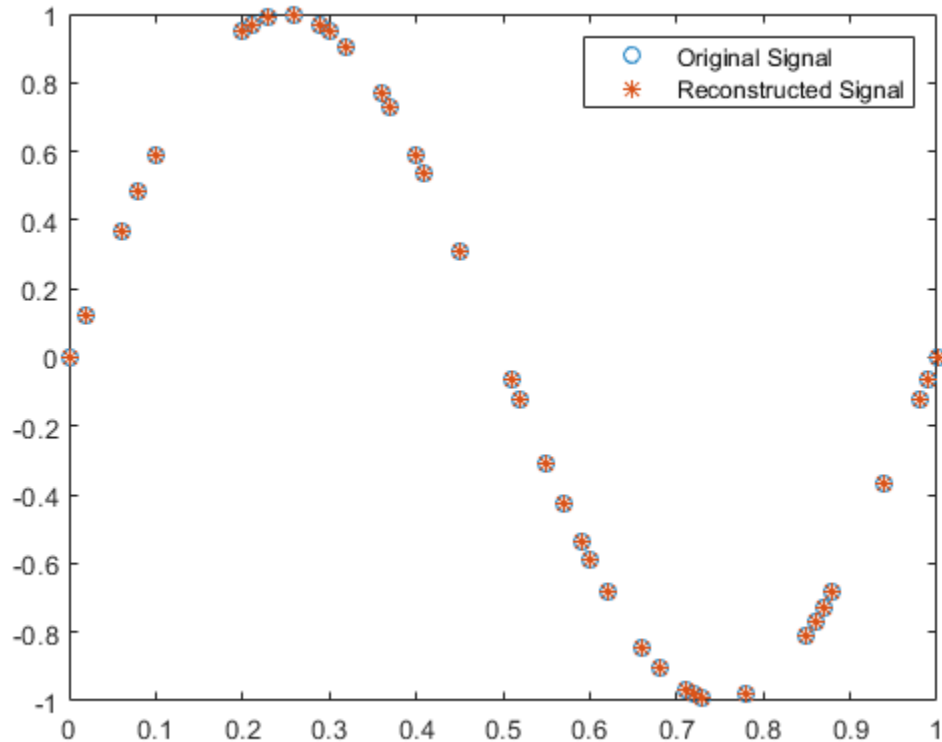
```
figure(2)  
stem(coefs)  
title('Wavelet Coefficients')
```



Perform the inverse multiscale local 1-D polynomial transform (`imlpt`) on the coefficients. Visualize the reconstructed signal.

```
y = imlpt(coefs,T,coefsPerLevel,scalingMoments);
```

```
figure(1)  
plot(T,y,'*')  
legend('Original Signal','Reconstructed Signal')  
hold off
```



Look at the total error to verify good reconstruction.

```
reconstructionError = sum(abs(y-sineWave))
```

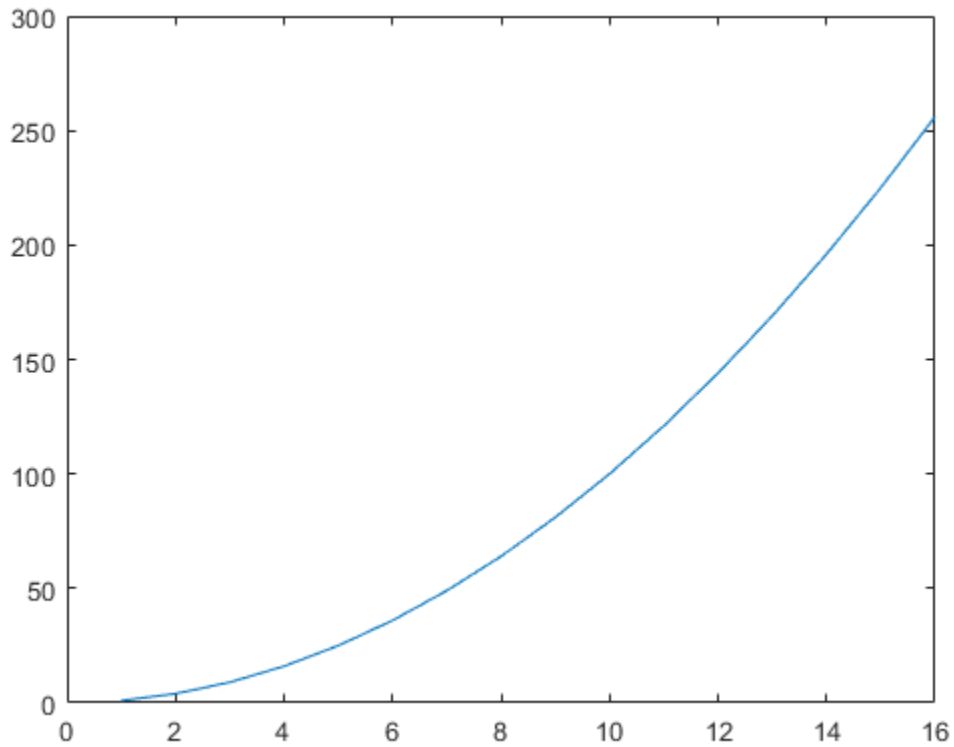
```
reconstructionError = 1.7552e-15
```

Specify Nondefault Dual Moments

Specify nondefault dual moments by using the `mlpt` function. Compare the results of analysis and synthesis using the default and nondefault dual moments.

Create an input signal and visualize it.

```
T = (1:16)';  
x = T.^2;  
plot(x)  
hold on
```



Perform the forward and inverse transform for the input signal using the default and nondefault dual moments.

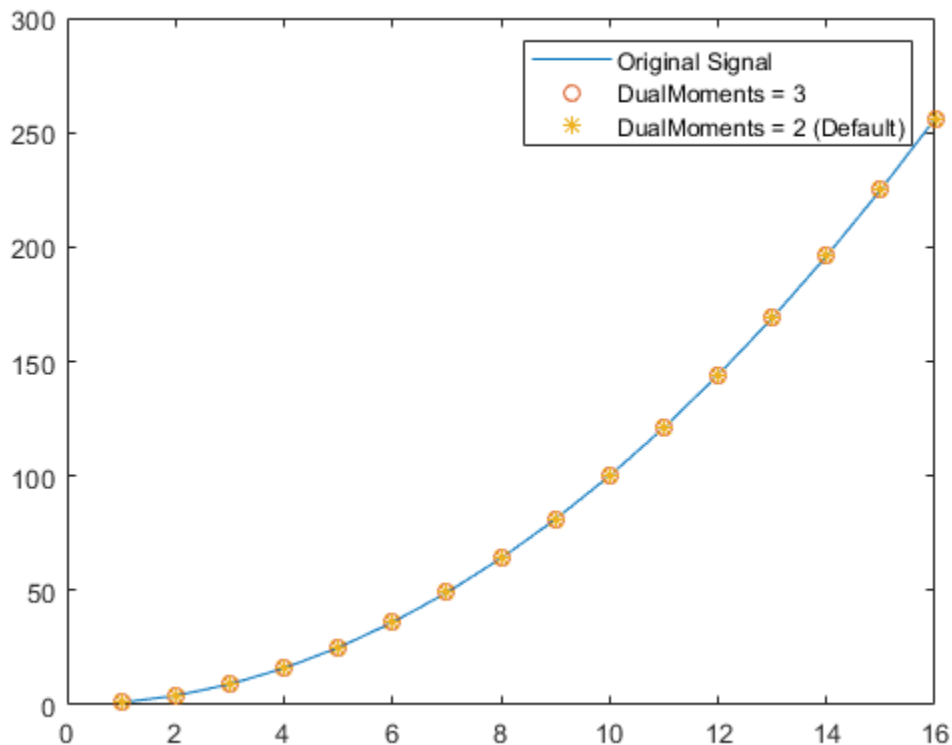
```
[w2,t2,nj2,scalingmoments2] = mlpt(x,T);  
y2 = imlpt(w2,t2,nj2,scalingmoments2);  
[w3,t3,nj3,scalingmoments3] = mlpt(x,T,'dualmoments',3);  
y3 = imlpt(w3,t3,nj3,scalingmoments3,'dualmoments',3);
```

Plot the reconstructed signal and verify perfect reconstruction using both the default and nondefault dual moments.

```
plot(y2,'o')
plot(y3,'*')
legend('Original Signal', ...
       'DualMoments = 3', ...
       'DualMoments = 2 (Default)');

fprintf('\nMean Reconstruction Error:\n');
Mean Reconstruction Error:
fprintf(' - Nondefault dual moments: %0.2f\n',mean(abs(y3-x)));
- Nondefault dual moments: 0.00
fprintf(' - Default dual moments: %0.2f\n\n',mean(abs(y2-x)));
- Default dual moments: 0.00

hold off
```



Specify Nondefault Resolution Levels

Resolution levels are the number of cascaded local polynomial smoothing operations. The details at each resolution level are obtained by predicting one half the samples based on a local polynomial interpolation of the other half. The difference between the predicted and actual values are the details at each resolution level. The scaling coefficients at each coarser resolution level are smoother versions of the higher resolution scaling coefficients. Only the final-level scaling coefficients are retained.

Increasing the number of resolution levels enables you to analyze narrowband coefficients for a computational and memory cost.

Create a dual-tone input signal, x , that contains high and low frequencies.

```
fs = 1000;  
t = (0:1/fs:10)';  
x = sin(499*pi.*t) + sin(2*pi.*t);
```

Use `mlpt` to obtain coefficients for minimum and maximum resolution levels. Print the computation time.

```
tic  
[w1,~,nj1,m1] = mlpt(x,t,1);  
computationTime1 = toc;  
fprintf('Level one computation time: %0.2f\n',computationTime1)
```

```
Level one computation time: 7.02
```

```
tic  
[w13,~,nj13,m13] = mlpt(x,t,13);  
computationTime13 = toc;  
fprintf('Level thirteen computation time: %0.2f\n',computationTime13)
```

```
Level thirteen computation time: 9.95
```

Use Default Time Instants

If your time instants are not known or specified, you can calculate the MLPT using default time instants.

Load a data signal corrupted with NaNs and with unknown time instants. Calculate the MLPT without specifying time instants. The resulting implied time instants is a vector of valid indices of the corrupted signal.

```
load(fullfile(matlabroot, 'examples', 'wavelet', 'CorruptedData.mat'));
```

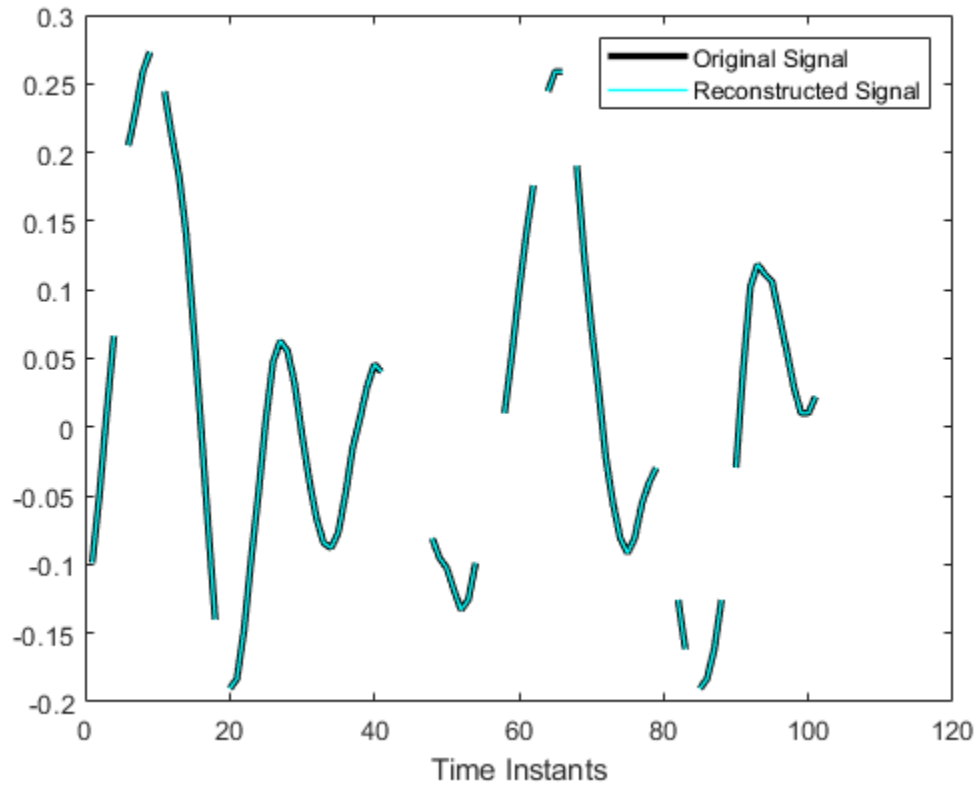
```
[w,t,nj,scalingMoments] = mlpt(yCorrupt);
```

Calculate the inverse MLPT and visualize the results. Reinsert NaNs to visualize gaps in the signal.

```
z = imlpt(w,t,nj,scalingMoments);
```



```
zToPlot = NaN(numel(yCorrupt),1);  
zToPlot(t) = z;  
  
plot(yCorrupt,'k','LineWidth',2.5)  
hold on  
plot(zToPlot,'c','LineWidth',1)  
hold off  
legend('Original Signal','Reconstructed Signal')  
xlabel('Time Instants')
```



Input Arguments

x — Input signal

vector | matrix

Input signal, specified as a vector or matrix.

- **matrix** — **x** must have at least two rows. `mlpt` operates independently on each column of **x**. The number of elements in **t** must equal the row dimension of **x**. Any NaNs in the columns of **x** must occur in the same rows.
- **vector** — **x** and **t** must have the same number of elements.

Data Types: `double`

t — Sampling instants

vector | duration array | datetime array

Sampling instants corresponding to the input signal, specified as a vector, `duration` array, or `datetime` array of monotonically increasing real values. The default value depends on the length of the input signal, **x**.

Data Types: `double` | `duration` | `datetime`

numLevel — Number of resolution levels

positive integer

Number of resolution levels, specified as a positive integer. The maximum value of `numLevel` depends on the shape of the input signal, **x**:

- **matrix** — `floor(log2(size(x,1)))`
- **vector** — `floor(log2(length(x)))`

If `numLevel` is not specified, `mlpt` uses the maximum value.

Data Types: `double`

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name, Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: 'DualMoments',3 computes a transform using three dual vanishing moments.

DualMoments — Number of dual vanishing moments

2 (default) | 3 | 4

Number of dual vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of 'DualMoments' and 2, 3 or 4.

Data Types: double

PrimalMoments — Number of primal vanishing moments

2 (default) | 3 | 4

Number of primal vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of 'PrimalMoments' and 2, 3, or 4.

Data Types: double

Prefilter — Prefilter before mlpt

'Haar' (default) | 'UnbalancedHaar' | 'None'

Prefilter before mlpt operation, specified as the comma-separated pair consisting of 'Prefilter' and 'Haar' [1], 'UnbalancedHaar', or 'None'.

Data Types: char | string

Output Arguments

coefs — MLPT coefficients

vector | matrix

MLPT coefficients, returned as a vector or matrix of coefficients, depending on the level to which the transform is calculated. coefs contains the approximation and detail coefficients.

Data Types: double

T — Sampling instants corresponding to output

vector | duration array

Sampling instants corresponding to output, returned as a vector or duration array of sample times obtained from x and t. The imlpt function requires T as an input. If the

input `t` is a `datetime` or `duration` array, `t` is converted to units that allow for the stable computation of the `mlpt` and `imlpt`. Then `T` is returned as a `duration` array.

Data Types: `double` | `duration`

coefsPerLevel — Coefficients per resolution level

vector

Coefficients per resolution level, returned as a vector containing the number of coefficients at each resolution level in `coefs`. The elements of `coefsPerLevel` are organized as follows:

- `coefsPerLevel(1)` — Number of approximation coefficients at the coarsest resolution level.
- `coefsPerLevel(i)` — Number of detail coefficients at resolution level `i`, where $i = \text{numLevel} - i + 2$ for $i = 2, \dots, \text{numLevel} + 1$.

The smaller the index `i`, the lower the resolution. The MLPT is two times redundant in the number of detail coefficients, but not in the number of approximation coefficients.

Data Types: `double`

scalingMoments — Scaling function moments

matrix

Scaling function moments, returned as a `length(coefs)`-by-`P` matrix, where `P` is the number of primal moments specified by the `PrimalMoments` name-value pair.

Data Types: `double`

Algorithms

Maarten Jansen developed the theoretical foundation of the multiscale local polynomial transform (MLPT) and algorithms for its efficient computation [1][2][3]. The MLPT uses a lifting scheme, wherein a kernel function smooths fine-scale coefficients with a given bandwidth to obtain the coarser resolution coefficients. The `mlpt` function uses only local polynomial interpolation, but the technique developed by Jansen is more general and admits many other kernel types with adjustable bandwidths [2].

References

- [1] Jansen, M. "Multiscale Local Polynomial Smoothing in a Lifted Pyramid for Non-Equispaced Data". *IEEE Transactions on Signal Processing*. Vol. 61, Number 3, 2013, pp. 545-555.
- [2] Jansen, M., and M. Amghar. "Multiscale local polynomial decompositions using bandwidths as scales". *Statistics and Computing* (forthcoming). 2016.
- [3] Jansen, M., and Patrick Oonincx. *Second Generation Wavelets and Applications*. London: Springer, 2005.

See Also

[imlpt](#) | [mlptdenoise](#) | [mlptrecon](#)

Topics

Smoothing Nonuniformly Sampled Data

Introduced in R2017a

mlptdenoise

Denoise signal using multiscale local 1-D polynomial transform

Syntax

```
y = mlptdenoise(x,t)
y = mlptdenoise(x,t,numLevel)
y = mlptdenoise( ____,Name,Value)
[y,T] = mlptdenoise( ____)
[y,T,thresholdedCoefs] = mlptdenoise( ____)
[y,T,thresholdedCoefs,originalCoefs] = mlptdenoise( ____)
```

Description

`y = mlptdenoise(x,t)` returns a denoised version of input signal `x` sampled at the sampling instants, `t`. If `x` or `t` contain NaNs, the union of the NaNs in `x` and `t` is removed before obtaining the `mlpt`.

`y = mlptdenoise(x,t,numLevel)` denoises `x` down to `numLevel`.

`y = mlptdenoise(____,Name,Value)` specifies `mlpt` properties using one or more `Name,Value` pair arguments, and any of the previous syntaxes

`[y,T] = mlptdenoise(____)` also returns the time instants for the denoised signal.

`[y,T,thresholdedCoefs] = mlptdenoise(____)` also returns the thresholded multiscale local 1-D polynomial transform coefficients.

`[y,T,thresholdedCoefs,originalCoefs] = mlptdenoise(____)` also returns the original multiscale local 1-D polynomial transform coefficients.

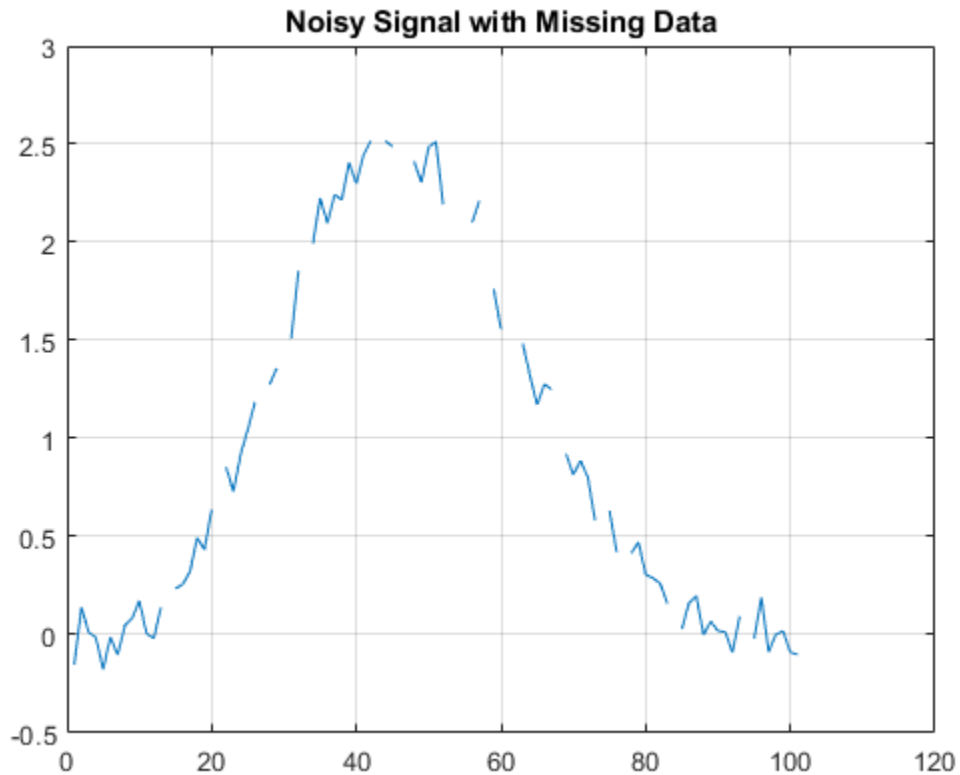
Examples

Specify Nondefault Denoising Method

Denoise a nonuniformly sampled spline signal with added noise using median smoothing and two primal vanishing moments. The nonuniformity of the signal is indicated by NaNs (missing data).

Load the data to your workspace and visualize it.

```
load nonuniformspline
plot(splinenoise)
grid on
title('Noisy Signal with Missing Data')
```

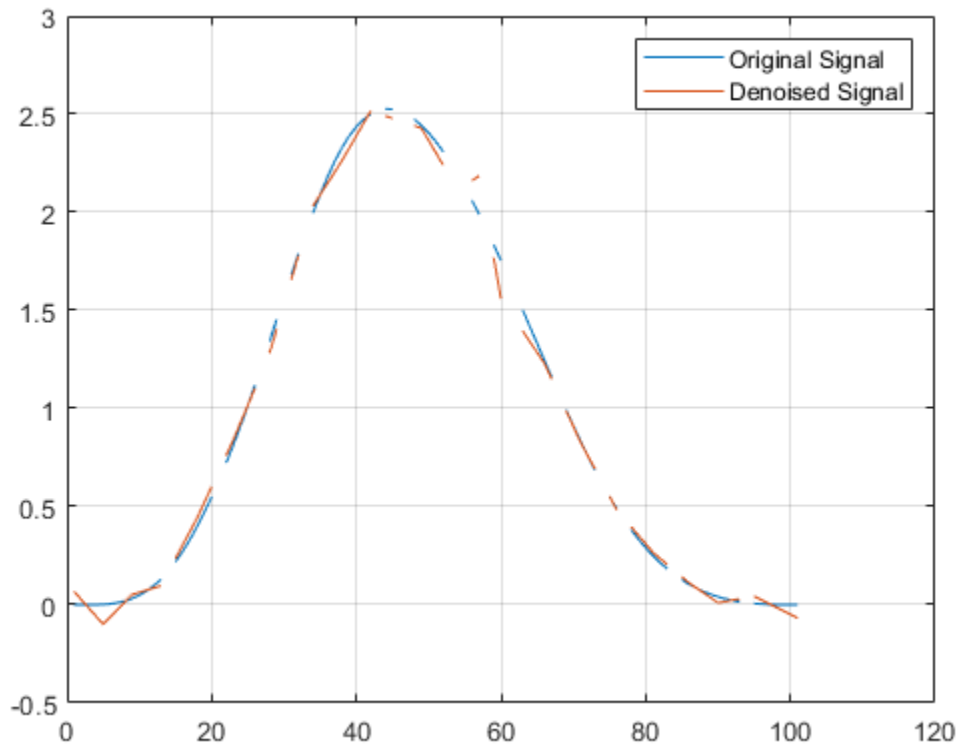


Denoise the data using the median denoising method.

```
xden = mlptdenoise(splinenoise,[], 'DenoisingMethod', 'median');
```

Replace the original missing data in the correct position for plotting purposes. Visualize the original and denoised signals.

```
denoisedsig = NaN(size(splinenoise));  
denoisedsig(~isnan(splinenoise)) = xden;  
figure  
plot([splinesig denoisedsig])  
grid on  
legend('Original Signal', 'Denoised Signal');
```



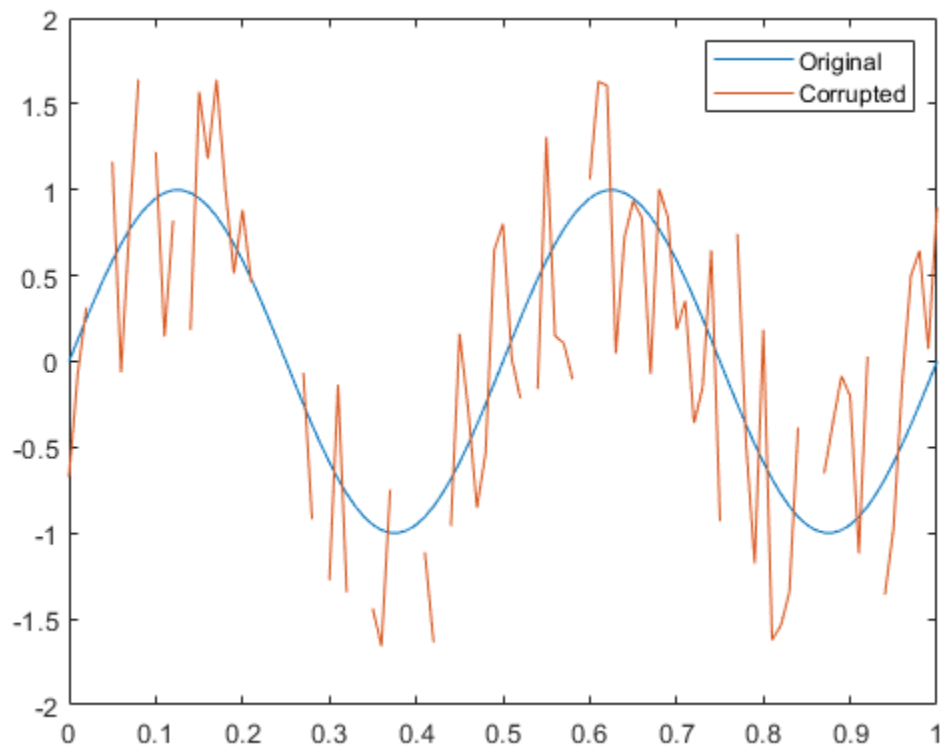
Denoise Using Multiscale Local Polynomial Transform

Reduce noise of signal using the multiscale local polynomial transform (MLPT).

Load a pure sine wave signal with uniform sampling, and a corrupted version of the signal.

```
load(fullfile(matlabroot, 'examples', 'wavelet', 'InputSamples.mat'))
```

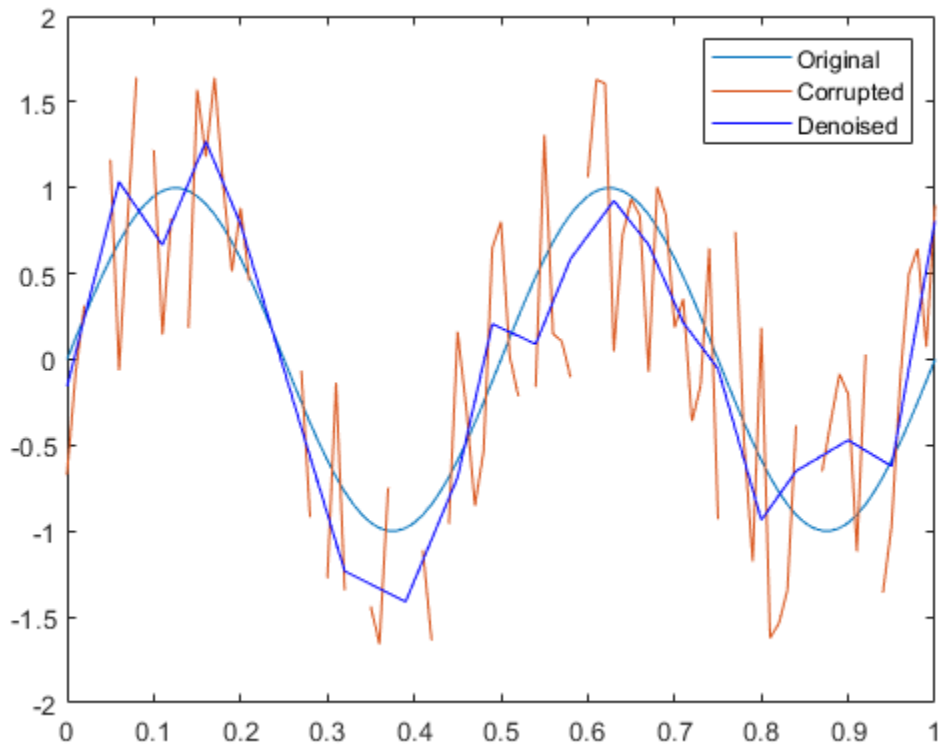
```
plot(t,x)  
hold on  
plot(tCorrupt,xCorrupt)  
legend('Original', 'Corrupted')
```



Use `mlptdenoise` to denoise the corrupted signal. Visually compare the corrupted and denoised signals against the original signal.

```
[xDenoised,tDenoised] = mlptdenoise(xCorrupt,tCorrupt);
```

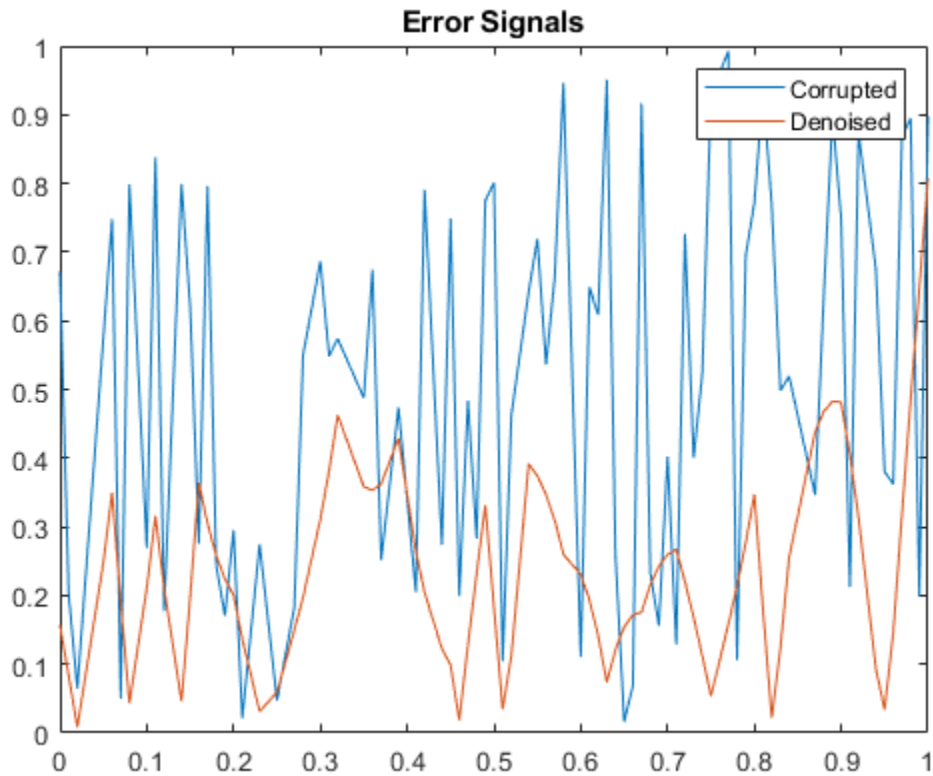
```
plot(tDenoised,xDenoised,'b')  
hold off  
legend('Original','Corrupted','Denoised')
```



Compare the error signals associated with the corrupted signal and the denoised signal. Remove NaNs from the signals for visualization purposes.

```
x(samplesToErase) = [];  
xCorrupt(samplesToErase) = [];
```

```
xCorruptError = abs(diff([x,xCorrupt],[],2));  
yError = abs(diff([x,xDenoised],[],2));  
  
plot(tDenoised,xCorruptError)  
hold on  
plot(tDenoised,yError)  
title('Error Signals')  
legend('Corrupted','Denoised')  
hold off
```



Specify Nondefault Denoising Level

By default, `mlptdenoise` denoises a signal based on the two highest-level detail coefficients. In this example, you denoise a signal to different levels and visualize the effect.

Create a multitone signal.

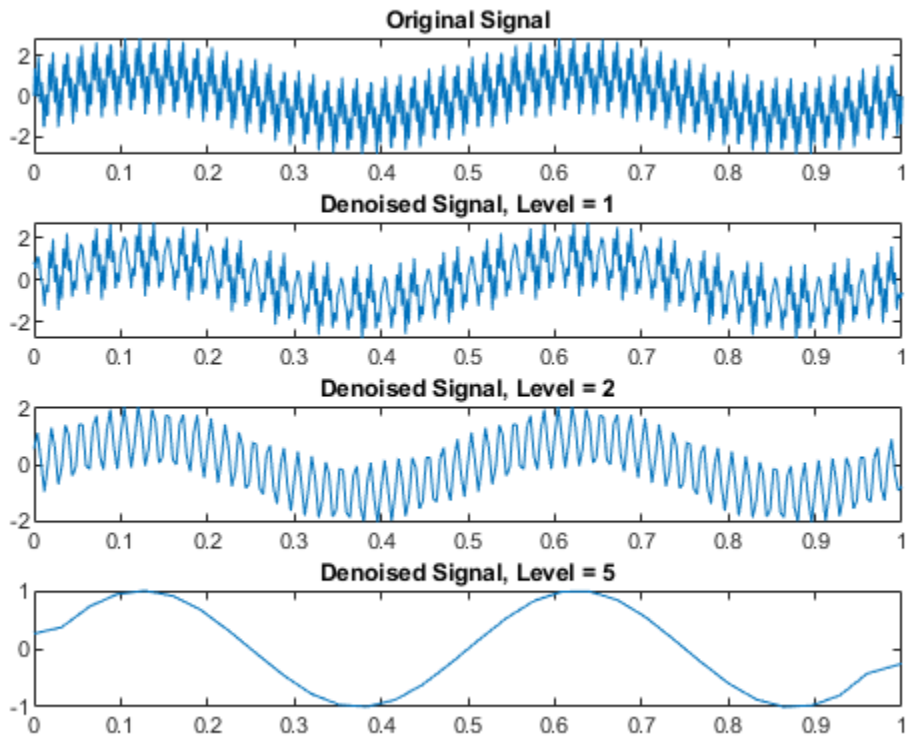
```
fs = 1000;  
t = 0:1/fs:1;  
x = sin(4*pi*t) + sin(120*pi*t) + sin(480*pi*t);
```

Denoise the signal to levels one, two, and five.

```
y1 = mlptdenoise(x,t,1);  
y2 = mlptdenoise(x,t,2);  
y5 = mlptdenoise(x,t,5);
```

Visualize the effect of level on the denoised signal.

```
subplot(4,1,1)  
plot(t,x)  
title('Original Signal')  
  
subplot(4,1,2)  
plot(t,y1)  
title('Denoised Signal, Level = 1')  
  
subplot(4,1,3)  
plot(t,y2)  
title('Denoised Signal, Level = 2')  
  
subplot(4,1,4)  
plot(t,y5)  
title('Denoised Signal, Level = 5')
```



Compare Thresholded and Nonthresholded Coefficients

The `mlptdenoise` function performs the forward MLPT, thresholds the coefficients as specified by the `'DenoisingMethod'` name-value pair. Then `mlptdenoise` performs the inverse MLPT to return a denoised signal in the domain of your original signal.

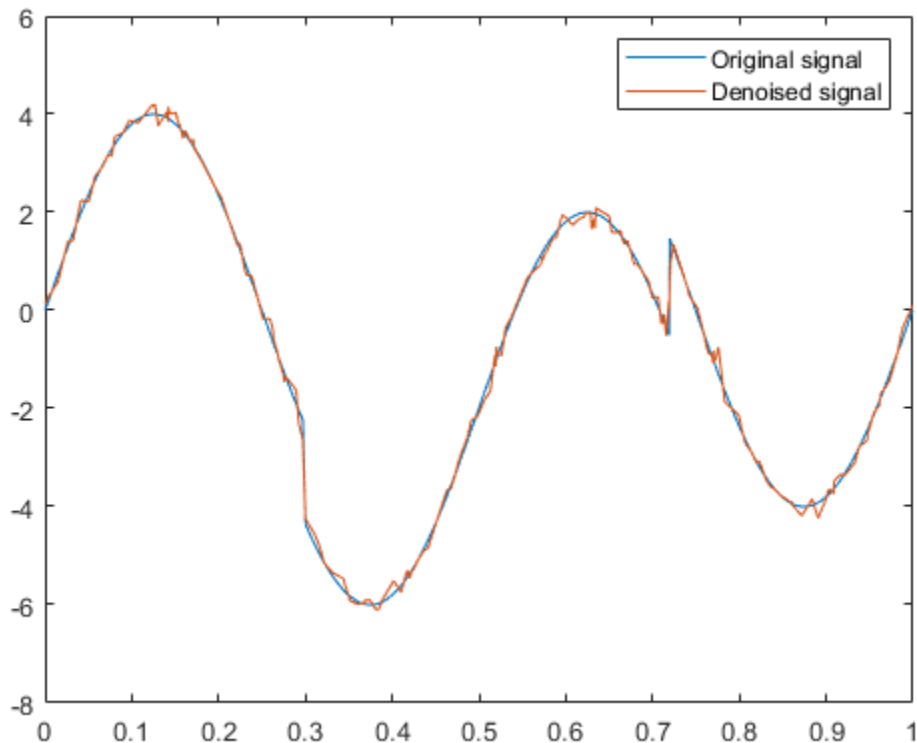
You can optionally return the thresholded and original coefficients for inspection and analysis.

Denoise a nonuniformly sampled signal using Stein's unbiased risk method. Return the denoised signal, the associated time instants, the thresholded MLPT coefficients, and the original MLPT coefficients. Plot the original and denoised signals.

```
load nonuniformheavisine;
```

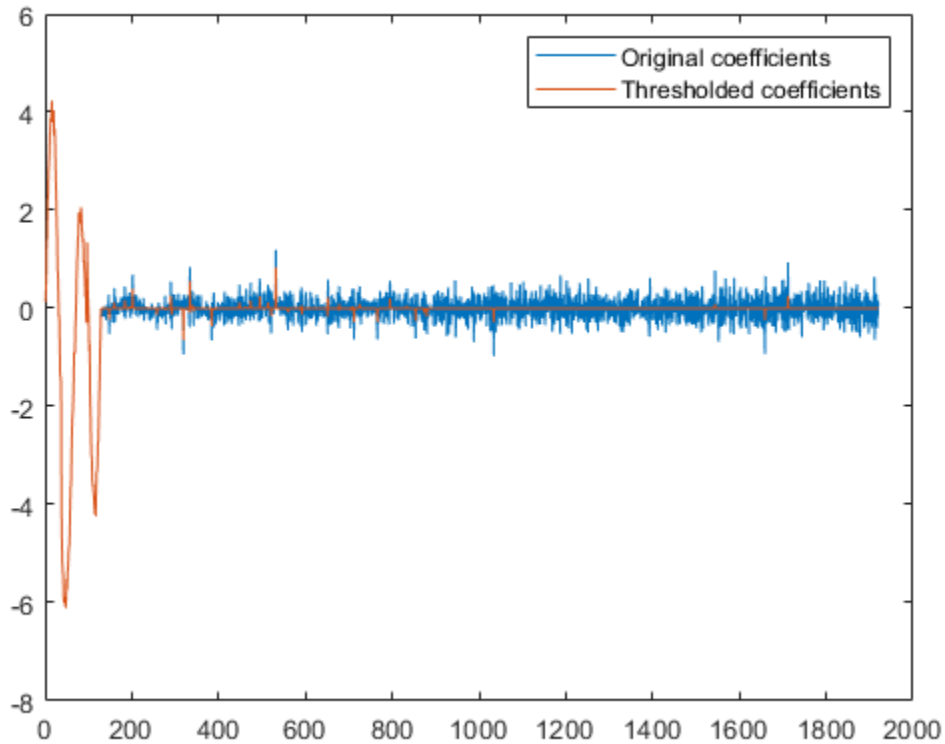
```
[xDenoised,t,wThrolded,w0Original] = mlptdenoise(x,t,3,'denoisingmethod','SURE');
```

```
plot(t,[f,xDenoised])  
legend('Original signal','Denoised signal')
```



Plot the original MLPT coefficients and the thresholded MLPT coefficients for comparison.

```
plot([wOriginal,wThrolded])  
legend('Original coefficients','Thresholded coefficients')
```



Input Arguments

x — Input signal

vector | matrix

Input signal, specified as a vector or matrix.

- **matrix** — `x` must have at least two rows. `mlpt` operates independently on each column of `x`. The number of elements in `t` must equal the row dimension of `x`. Any NaNs in the columns of `x` must occur in the same rows.
- **vector** — `x` and `t` must have the same number of elements.

Data Types: `double`

t — Sampling instants

`vector` | `duration array` | `datetime array`

Sampling instants corresponding to the input signal, specified as a `vector`, `duration array`, or `datetime array` of monotonically increasing real values. The default value depends on the length of the input signal, `x`.

Data Types: `double` | `duration` | `datetime`

numLevel — Number of resolution levels

2 (default) | `positive integer`

Number of resolution levels, specified as a `positive integer`. The maximum value of `numLevel` depends on the shape of the input signal, `x`:

- **matrix** — `floor(log2(size(x,1)))`
- **vector** — `floor(log2(length(x)))`

`mlptdenoise` denoises `x` by thresholding all detail coefficients of an MLPT calculated for `numLevel` resolution levels.

Data Types: `double`

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'DualMoments', 3` computes a transform using three dual vanishing moments.

DualMoments — Number of dual vanishing moments

2 (default) | 3 | 4

Number of dual vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of 'DualMoments' and 2, 3 or 4.

Data Types: double

PrimalMoments — Number of primal vanishing moments

2 (default) | 3 | 4

Number of primal vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of 'PrimalMoments' and 2, 3, or 4.

Data Types: double

Prefilter — Prefilter before mlpt

'Haar' (default) | 'UnbalancedHaar'

Prefilter before mlpt operation, specified as the comma-separated pair consisting of 'Prefilter' and 'Haar' or 'UnbalancedHaar'. If no prefilter is specified, 'Haar' is used by default.

Data Types: char | string

DenoisingMethod — Denoising method applied to MLPT detail coefficients

'Bayesian' (default) | 'Median' | 'SURE' | 'FDR'

Denoising method applied to MLPT detail coefficients, specified as the comma-separated pair consisting of 'DenoisingMethod' and 'Bayesian', 'Median', 'SURE', or 'FDR'.

Note 'FDR' has an optional argument for the Q-value. Q is the proportion of false positives and is specified as a real-valued scalar between zero and one. To specify 'FDR' with a Q-value, use a cell array, where the second element is the Q-value, for example 'DenoisingMethod', {'FDR', 0.01}. If unspecified, Q defaults to 0.05.

Data Types: char | string

Output Arguments

y — Denoised version of the input signal

vector | matrix

Denoised version of the input signal, returned as a vector or matrix. The size of `y` depends on the size of `x` and the union of NaNs in `x` and `t`.

By default, the `mlpt` is denoised based on the two highest resolution detail coefficients, unless `x` has fewer than four samples. If `x` has fewer than four samples, the `mlpt` is denoised based only on the highest resolution detail coefficients.

Data Types: `double`

T — Sampling instants corresponding to output

vector | duration array

Sampling instants corresponding to the output, returned as a vector or duration array obtained from `x` and the input `t`. If the input `t` is a `datetime` or duration array, `t` is converted to units that enable stable `mlpt` and `implt` computation. Then `T` is returned as a duration array.

Data Types: `double` | `duration`

thresholdedCoefs — Thresholded MLPT coefficients

vector | matrix

Thresholded MLPT coefficients, returned as a vector or matrix. The size of `thresholdedCoefs` depends on the size of `x` and the level to which the transform is calculated.

Data Types: `double`

originalCoefs — Original MLPT coefficients

vector | matrix

Original MLPT coefficients, returned as a vector or matrix. The size of `originalCoefs` depends on the size of `x` and the level to which the transform is calculated.

Data Types: `double`

Algorithms

Maarten Jansen developed the theoretical foundation of the multiscale local polynomial transform (MLPT) and algorithms for its efficient computation [1][2][3]. The MLPT uses a lifting scheme, wherein a kernel function smooths fine-scale coefficients with a given bandwidth to obtain the coarser resolution coefficients. The `mlpt` function uses only local

polynomial interpolation, but the technique developed by Jansen is more general and admits many other kernel types with adjustable bandwidths [2].

References

- [1] Jansen, M. "Multiscale Local Polynomial Smoothing in a Lifted Pyramid for Non-Equispaced Data". *IEEE Transactions on Signal Processing*, Vol. 61, Number 3, 2013, pp. 545-555.
- [2] Jansen, M., and M. Amghar. "Multiscale local polynomial decompositions using bandwidths as scales". *Statistics and Computing* (forthcoming). 2016.
- [3] Jansen, M., and Patrick Oonincx. *Second Generation Wavelets and Applications*. London: Springer, 2005.

See Also

`imlpt` | `mlpt` | `mlptrecon`

Topics

Smoothing Nonuniformly Sampled Data

Introduced in R2017a

mlptrecon

Reconstruct signal using inverse multiscale local 1-D polynomial transform

Syntax

```
y = mlptrecon(type,coefs,T,coefsPerLevel,scalingMoments,  
reconstructionLevel)  
y = mlptrecon( ____,Name,Value)
```

Description

`y = mlptrecon(type,coefs,T,coefsPerLevel,scalingMoments, reconstructionLevel)` returns an approximation to the inverse multiscale 1-D polynomial transform (MLPT) of `coefs`.

`y = mlptrecon(____,Name,Value)` specifies `mlptrecon` properties using one or more `Name,Value` pair arguments and the input arguments from the previous syntax.

Examples

Detect and Localize High-Frequency Content

Create a low-frequency signal with high-frequency blips.

```
t = (0:0.01:10)';  
x = sin(2*pi.*t) + 0.5*sin(pi.*t+0.1);  
bliptime = (0:0.01:0.5)';  
blip = sin(50*pi.*bliptime).*triang(numel(bliptime));  
for i = [200,700,900]  
    x(i:i+numel(bliptime)-1) = x(i:i+numel(bliptime)-1)+blip;  
end
```

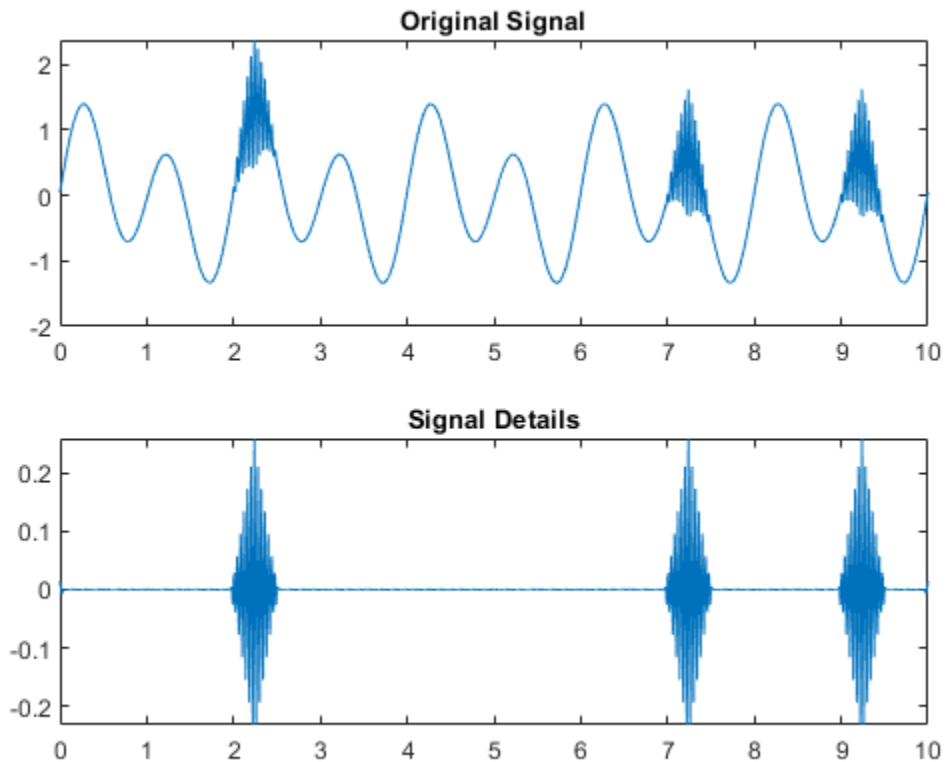
Perform a multilevel polynomial transform. Perform the inverse multilevel polynomial transform using the detail coefficients.

```
[w,t,nj,scalingmoments] = mlpt(x,t);  
yDetails = mlptrecon('d',w,t,nj,scalingmoments,1);
```

Plot the original signal and the processed signal.

```
subplot(2,1,1)  
plot(t,x)  
title('Original Signal')
```

```
subplot(2,1,2)  
plot(t,yDetails)  
title('Signal Details')
```

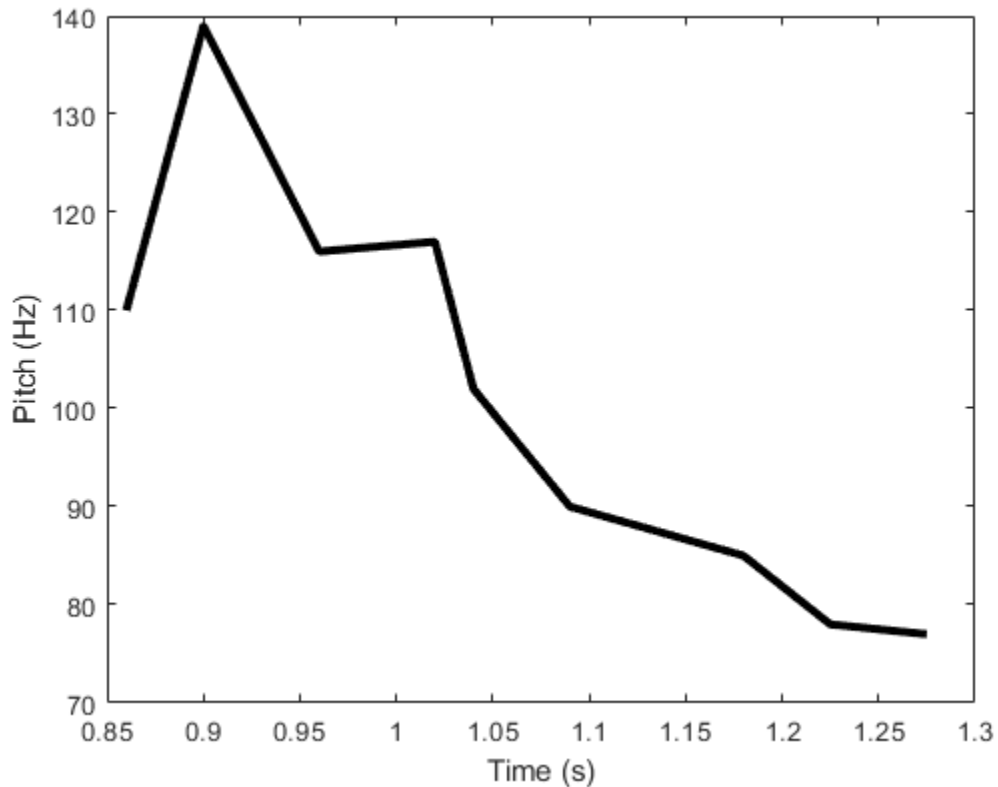


Approximate Data by Choosing Reconstruction Coefficients

Approximate data using multiscale local polynomial transform (MLPT) reconstruction. Use `mlptrecon` to approximate a corrupted and sparsely sampled pitch contour.

Load input data and visualize it.

```
load(fullfile(matlabroot,'examples','wavelet','CorruptedPitchData.mat'));  
plot(time,pitchContour,'k','linewidth',3)  
hold on  
xlabel('Time (s)')  
ylabel('Pitch (Hz)')
```



Compute the MLPT of the pitch contour.

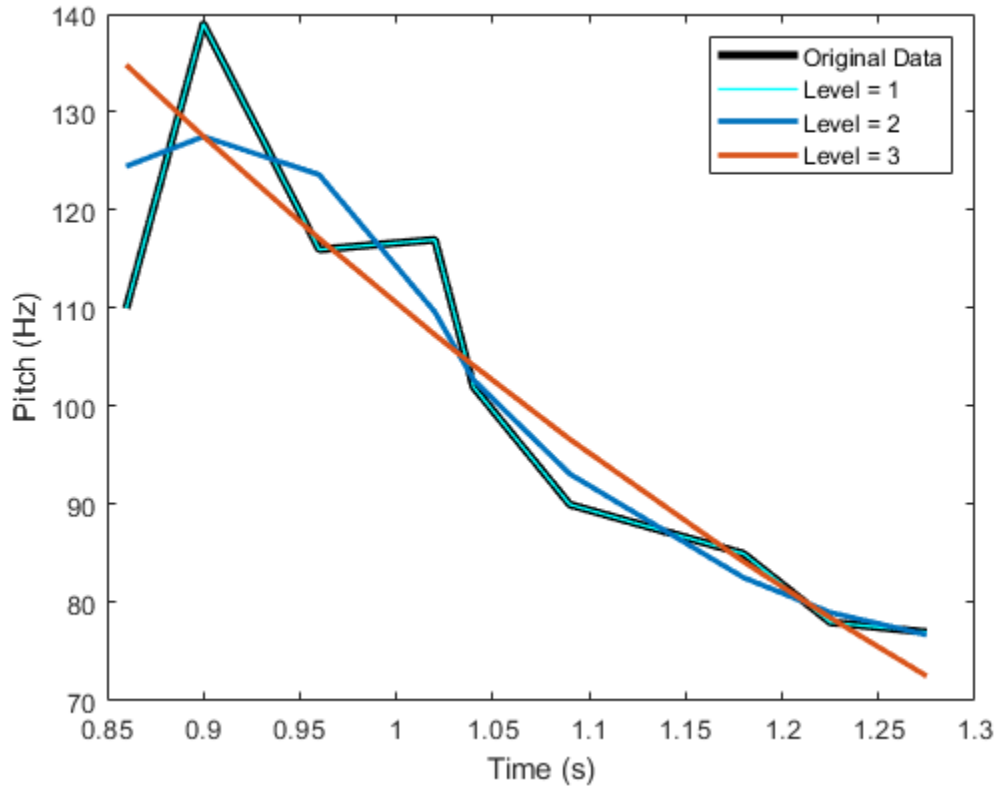
```
[w,t,nj,scalingMoments] = mlpt(pitchContour,time, ...  
    'DualMoments',3, ...  
    'PrimalMoments',4, ...  
    'PreFilter','none');
```

Use `mlptrecon` to reconstruct the signal using the approximation coefficients at different levels.

```
y = zeros(numel(t),3);  
for level = 1:3  
    y(:,level) = mlptrecon('a',w,t,nj,scalingMoments,level,'DualMoments',3);  
end
```

Plot the reconstructed signals. Level two obtains the best smoothed estimate.

```
plot(t,y(:,1),'c','linewidth',1)  
plot(t,y(:,2),'linewidth',2)  
plot(t,y(:,3),'linewidth',2)  
legend('Original Data','Level = 1','Level = 2','Level = 3')  
hold off
```



Input Arguments

type — Type of coefficients

'a' | 'd'

Type of coefficients used to reconstruct the signal, specified as 'a' or 'd'.

- 'a' — Approximation coefficients
- 'd' — Detail coefficients

Approximation coefficients are a lowpass representation of the input. At each level, the approximation coefficients are divided into coarser approximation and detail coefficients.

Data Types: `char` | `string`

coefs — MLPT coefficients

vector | matrix

MLPT coefficients, specified as a vector or matrix of MLPT coefficients returned by the `mlpt` function.

Data Types: `double`

T — Sampling instants corresponding to output

vector | duration array

Sampling instants corresponding to `y`, specified as a vector or duration array of increasing values returned by the `mlpt` function.

Data Types: `double` | `duration`

coefsPerLevel — Coefficients per resolution level

vector

Coefficients per resolution level, specified as a vector containing the number of coefficients at each resolution level in `coefs`. `coefsPerLevel` is an output argument of the `mlpt` function.

The elements of `coefsPerLevel` are organized as follows:

- `coefsPerLevel(1)` — Number of approximation coefficients at the coarsest resolution level.
- `coefsPerLevel(i)` — Number of detail coefficients at resolution level `i`, where $i = \text{numLevel} - i + 2$ for $i = 2, \dots, \text{numLevel} + 1$. `numLevel` is the number of resolution levels used to calculate the MLPT. `numLevel` is inferred from `coefsPerLevel`: $\text{numLevel} = \text{length}(\text{coefsPerLevel} - 1)$.

The smaller the index `i`, the lower the resolution. The MLPT is two times redundant in the number of detail coefficients, but not in the number of approximation coefficients.

Data Types: `double`

scalingMoments — Scaling function moments

matrix

Scaling function moments, specified as a `length(coefs)`-by-`P` matrix, where `P` is the number of primal moments specified by the MLPT.

Data Types: `double`

reconstructionLevel — Resolution level used for reconstruction

positive integer

Resolution level used for reconstruction, specified as a positive integer less than or equal to `length(coefsPerLevel - 1)`. `length(coefsPerLevel - 1)` is the number of resolution levels used to calculate the MLPT. Increasing the value of `reconstructionLevel` corresponds to reconstructing your signal with coarser resolution approximations.

Data Types: `double`

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'DualMoments', 3` computes a transform using three dual vanishing moments.

DualMoments — Number of dual vanishing moments

2 (default) | 3 | 4

Number of dual vanishing moments in the lifting scheme, specified as the comma-separated pair consisting of `'DualMoments'` and 2, 3 or 4. The number of dual moments must match the number used by `mlpt`.

Data Types: `double`

Output Arguments

y — Reconstructed approximation or details of signal

vector | matrix

Reconstructed approximation or details of signal, returned as a vector or matrix, depending on the inputs to the `mlpt` function.

Data Types: `double`

Algorithms

Maarten Jansen developed the theoretical foundation of the multiscale local polynomial transform (MLPT) and algorithms for its efficient computation [1][2][3]. The MLPT uses a lifting scheme, wherein a kernel function smooths fine-scale coefficients with a given bandwidth to obtain the coarser resolution coefficients. The `mlpt` function uses only local polynomial interpolation, but the technique developed by Jansen is more general and admits many other kernel types with adjustable bandwidths [2].

References

- [1] Jansen, M. "Multiscale Local Polynomial Smoothing in a Lifted Pyramid for Non-Equispaced Data." *IEEE Transactions on Signal Processing*. Vol. 61, Number 3, 2013, pp. 545–555.
- [2] Jansen, M. and M. Amghar. "Multiscale local polynomial decompositions using bandwidths as scales". *Statistics and Computing* (forthcoming). 2016.
- [3] Jansen, M. and Patrick Oonincx. *Second Generation Wavelets and Applications*. London: Springer, 2005.

See Also

`imlpt` | `mlpt` | `mlptdenoise`

Topics

Smoothing Nonuniformly Sampled Data

Introduced in R2017a

modwpt

Maximal overlap discrete wavelet packet transform

Syntax

```
wpt = modwpt(x)
wpt = modwpt(x,wname)
wpt = modwpt(x,lo,hi)
wpt = modwpt( ____,lev)

[wpt,packetlevs] = modwpt( ____)
[wpt,packetlevs,cfreq] = modwpt( ____)
[wpt,packetlevs,cfreq,energy] = modwpt( ____)
[wpt,packetlevs,cfreq,energy,relenergy] = modwpt( ____)

[ ____ ] = modwpt( ____,Name,Value)
```

Description

`wpt = modwpt(x)` returns the terminal nodes for the maximal overlap discrete wavelet packet transform (MODWPT) for the 1-D real-valued signal, `x`.

Note The output of the MODWPT is time-delayed compared to the input signal. Most filters used to obtain the MODWPT have a nonlinear phase response, which makes compensating for the time delay difficult. This is true for all orthogonal scaling and wavelet filters, except the Haar wavelet. It is possible to time-align the coefficients with the signal features, but the result is an approximation, not an exact alignment with the original signal. The MODWPT partitions the energy among the wavelet packets at each level. The sum of the energy over all the packets equals the total energy of the input signal. The output of MODWPT is useful for applications where you want to analyze the energy levels in different packets.

The MODWPT details (`modwptdetails`) are the result of zero-phase filtering of the signal. The features in the MODWPT details align exactly with features in the input signal. For a given level, summing the details for each sample returns the exact original signal.

The output of the MODWPT details is useful for applications that require time-alignment, such as nonparametric regression analysis.

`wpt = modwpt(x, wname)` returns the MODWPT using the orthogonal wavelet filter specified by the `wname`.

`wpt = modwpt(x, lo, hi)` returns the MODWPT using the orthogonal scaling filter, `lo`, and wavelet filter, `hi`.

`wpt = modwpt(____, lev)` returns the terminal nodes of the wavelet packet tree at positive integer level `lev`.

`[wpt, packetlevs] = modwpt(____,)` returns a vector of transform levels corresponding to the rows of `wpt`.

`[wpt, packetlevs, cfreq] = modwpt(____,)` returns the center frequencies of the approximate passbands corresponding to the rows of `wpt`.

`[wpt, packetlevs, cfreq, energy] = modwpt(____,)` returns the energy (squared L2 norm) of the wavelet packet coefficients for the nodes in `wpt`.

`[wpt, packetlevs, cfreq, energy, relenergy] = modwpt(____,)` returns the relative energy for the wavelet packets in `wpt`.

`[____,] = modwpt(____, Name, Value)` returns the MODWPT with additional options specified by one or more `Name, Value` pair arguments.

Examples

MODWPT Using Default Wavelet

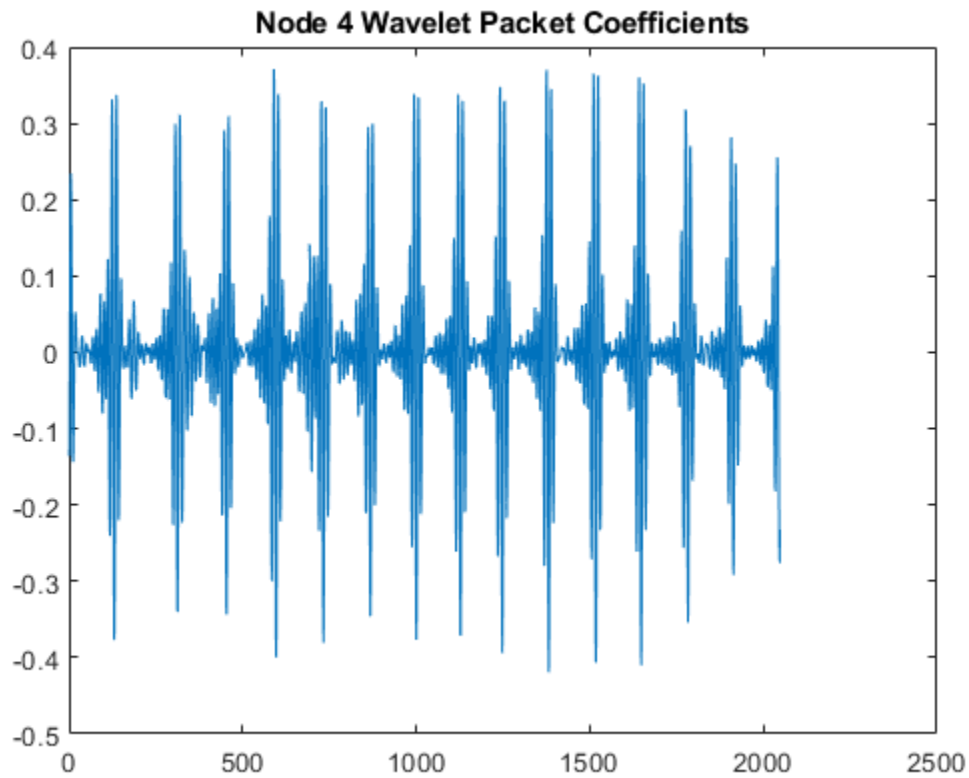
Obtain the MODWPT of an electrocardiogram (ECG) signal using the default length 18 Fejer-Korovkin ('fk18') wavelet.

```
load wecg;  
wpt = modwpt(wecg);
```

`wpt` is a 16-by-2048 matrix containing the sequency-ordered wavelet packet coefficients for the wavelet packet transform nodes. In this case, the nodes are at level 4. Each node

corresponds to an approximate passband filtering of $[nf_s/2^5, (n+1)f_s/2^5)$, where $n = 0, \dots, 15$, and f_s is the sampling frequency. Plot the wavelet packet coefficients at node (4,2), which is level 4, node 2.

```
plot(wpt(3,:))  
title('Node 4 Wavelet Packet Coefficients')
```



MODWPT Using Daubechies Extremal Phase Wavelet with Two Vanishing Moments

Obtain the MODWPT of Southern Oscillation Index data with the Daubechies extremal phase wavelet with two vanishing moments ('db2').

```
load soi;
wsoi = modwpt(soi, 'db2');
```

Verify that the size of the resulting transform contains 16 nodes. Each node is in a separate row.

```
size(wsoi)
ans = 1×2
      16      12998
```

MODWPT Using Scaling and Wavelet Filters

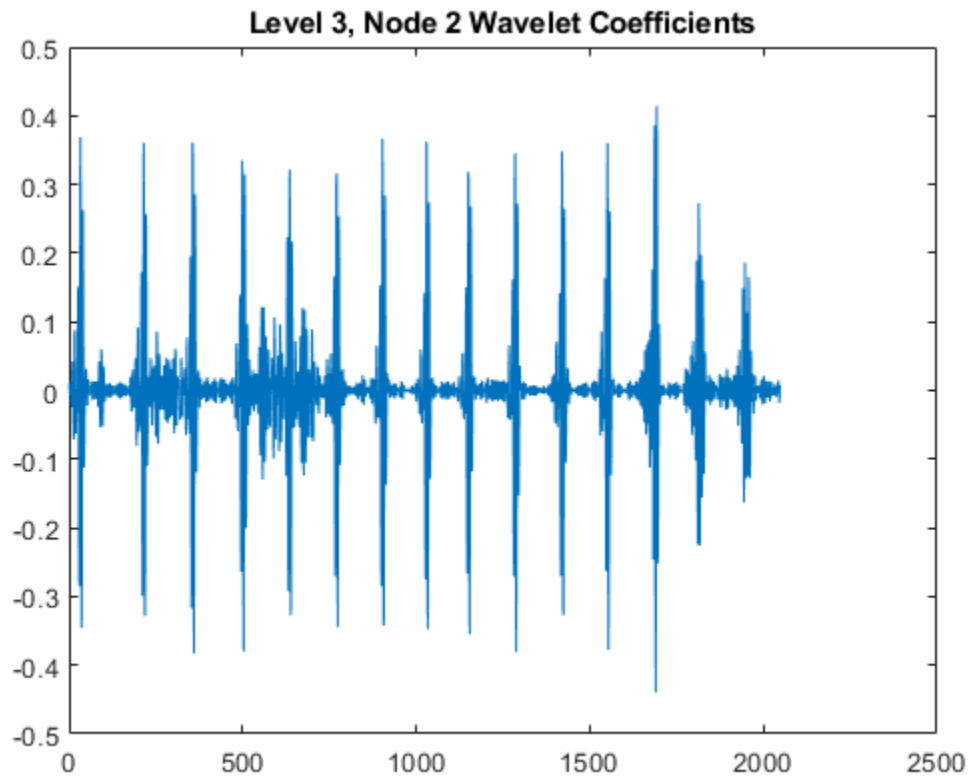
Obtain the MODWPT of an ECG waveform using the Fejer-Korovkin length 18 scaling and wavelet filters.

```
load wecg;
[lo,hi] = wfilters('fk18');
wpt = modwpt(wecg,lo,hi);
```

MODWPT Full Packet Tree and Passband Center Frequencies

Obtain the MODWPT and full wavelet packet tree of an ECG waveform using the default length 18 Fejer-Korovkin ('fk18') wavelet. Extract and plot the node coefficients at level 3, node 2.

```
load wecg;
[wpt,packetlevels,cfreq] = modwpt(wecg, 'FullTree', true);
p3 = wpt(packetlevels==3,:);
plot(p3(3,:))
title('Level 3, Node 2 Wavelet Coefficients')
```



Display the center frequencies at level 3.

```
cfreq(packetlevels==3,:)
```

```
ans = 8x1
```

```
0.0313  
0.0938  
0.1563  
0.2188  
0.2813  
0.3438  
0.4063
```


0.4688

MODWPT Energy and Relative Energy

Obtain and plot the MODWPT energy and relative energy of an ECG waveform.

```
load wecg
[wpt,~,cfreq,energy,relenergy] = modwpt(wecg);
```

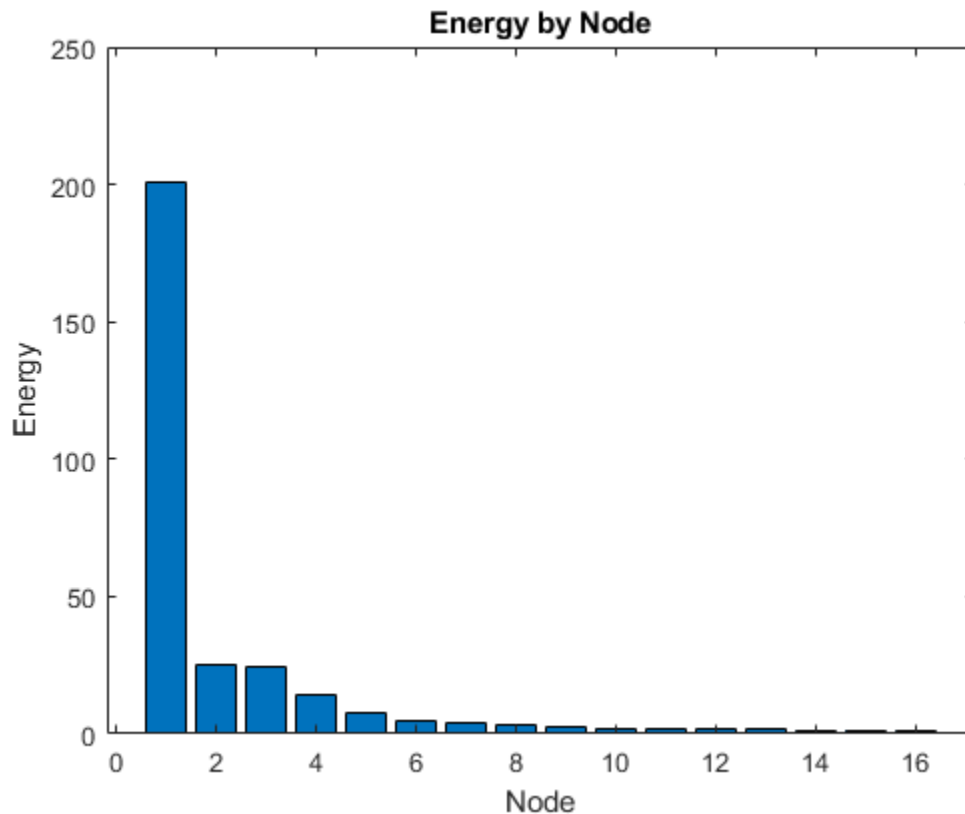
Show that the sum of the MODWPT energies is equal to the sum of the energy in the original signal. The difference between the total MODWPT energy and the signal energy is small enough to be considered insignificant.

```
disp(['Difference between MODWPT energy and signal energy: ',num2str(sum(energy)-sum(w
```

```
Difference between MODWPT energy and signal energy: 3.612e-09
```

Plot the MODWPT energy by node.

```
figure
bar(1:16,energy)
xlabel('Node')
ylabel('Energy')
title('Energy by Node')
```

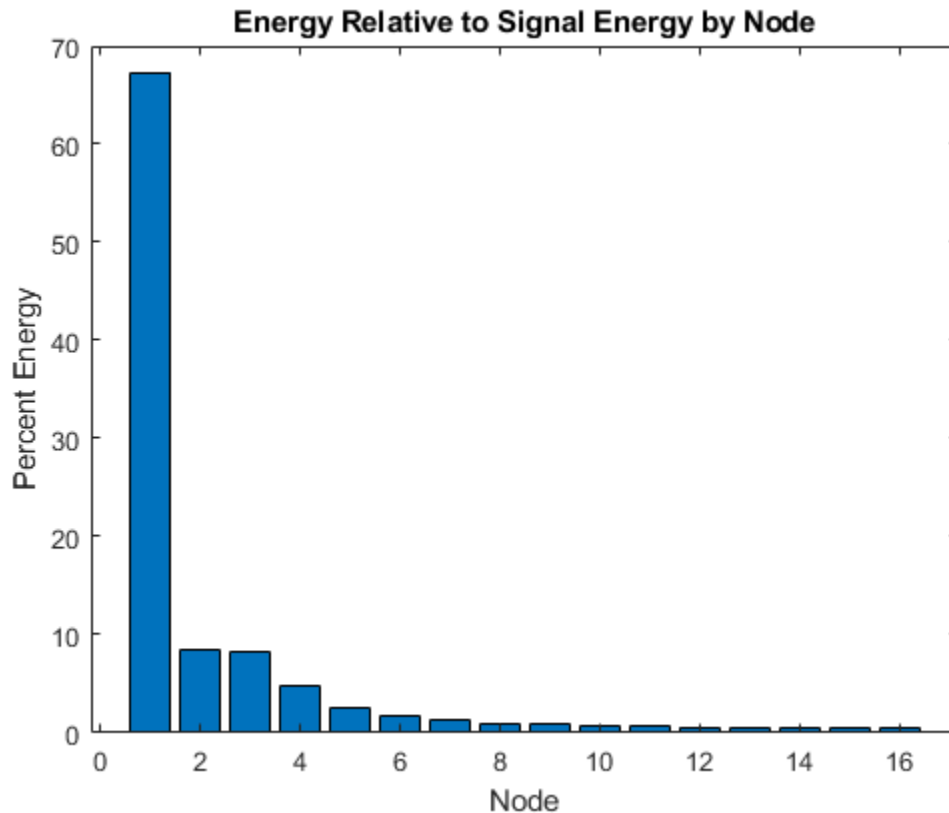


```
disp(['Total power in passband: ', num2str(energy(1))])
```

```
Total power in passband: 200.8446
```

Plot the relative energy and show the percentage of signal energy in the first passband [0,5.6250].

```
figure  
bar(1:16, relenergy*100)  
xlabel('Node')  
ylabel('Percent Energy')  
title('Energy Relative to Signal Energy by Node')
```



```
disp(['Percentage of signal power in passband: ',num2str(releenergy(1)*100)])
```

```
Percentage of signal power in passband: 67.3352
```

Time-Aligned MODWPT

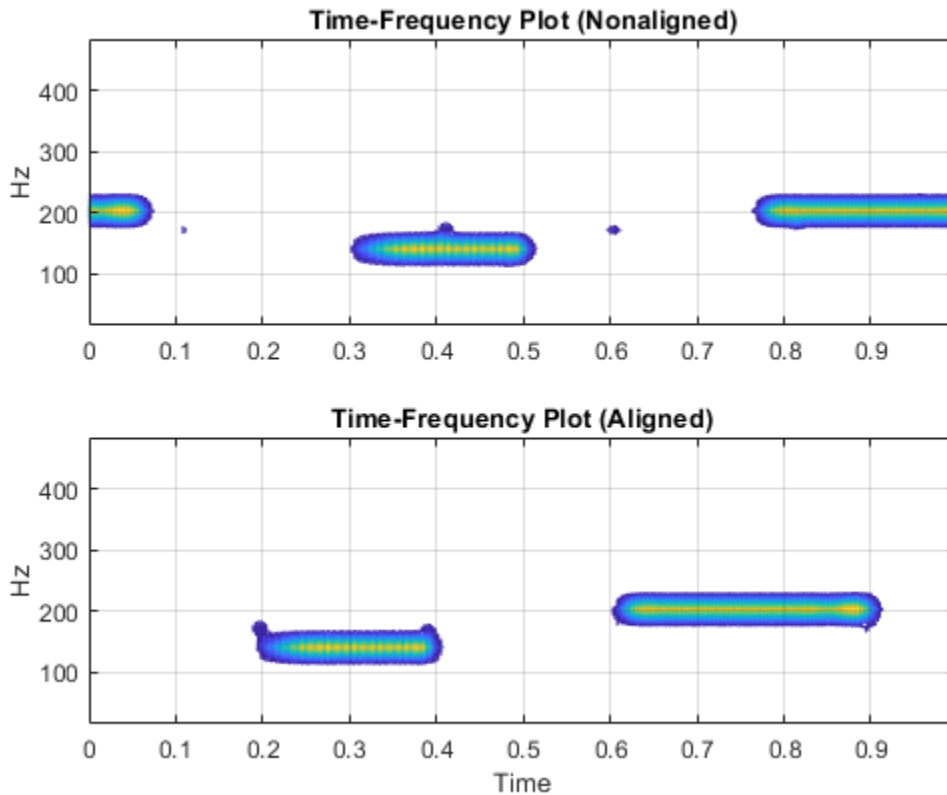
Obtain the time-aligned MODWPT of two intermittent sine waves in noise. The sine wave frequencies are 150 Hz and 200 Hz. The data is sampled at 1000 Hz.

```
dt = 0.001;
t = 0:dt:1-dt;
x = cos(2*pi*150*t).*(t>=0.2 & t<0.4)+ sin(2*pi*200*t).*(t>0.6 & t<0.9);
```

```
y = x+0.05*randn(size(t));  
[wpta,~,Falign] = modwpt(x, 'TimeAlign', true);  
[wptn,~,Fnon] = modwpt(x);
```

Compare the nonaligned and time-aligned time-frequency plots.

```
subplot(2,1,1);  
contour(t,Fnon.*(1/dt),abs(wptn).^2);  
grid on;  
ylabel('Hz');  
title('Time-Frequency Plot (Nonaligned)');  
subplot(2,1,2);  
contour(t,Falign.*(1/dt),abs(wpta).^2);  
grid on;  
xlabel('Time');  
ylabel('Hz');  
title('Time-Frequency Plot (Aligned)');
```



Input Arguments

x — Input signal

real-valued vector

Input signal, specified as a real-valued row or column vector. `x` must have at least two elements.

Data Types: double

wname — Analyzing wavelet filter

'fk18' (default) | 'haar' | 'db2' | ...

Analyzing wavelet filter specified as a that corresponds to an orthogonal wavelet. If you specify the scaling (`lo`) and wavelet (`hi`) filters, `modwpt` ignores the `wname` input.

Valid orthogonal wavelet families begin with one of the following, followed by an integer, N , for example, `sym4`. Note, however, that `'haar'` is not followed by an integer.

- `'haar'` — Haar wavelet, which is the same as Daubechies wavelet with one vanishing moment, `'db1'`.
- `'dbN'` — Daubechies wavelet with N vanishing moments
- `'symN'` — Symlets wavelet with N vanishing moments
- `'coifN'` — Coiflets wavelet with N vanishing moments
- `'fkN'` — Fejer-Korovkin wavelet with N coefficients

To check if your wavelet is orthogonal, use `wavemngr('type',wname)` and verify that it returns 1 as the wavelet type. To determine valid values for N , use `waveinfo`, for example, `waveinfo('fk')`.

lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. `lo` must satisfy the conditions necessary to generate an orthogonal scaling function. You cannot specify both the scaling-wavelet filters and the `wname` input.

hi — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. `hi` must satisfy the conditions necessary to generate an orthogonal wavelet. You cannot specify both the scaling-wavelet filters and the `wname` input.

lev — Transform level

positive integer

Transform level, specified as a positive integer less than or equal to `floor(log2(numel(x)))`.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name, Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes.

You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

Example: `'Fulltree', true` returns the full wavelet packet tree

FullTree — Full packet tree

`false` (default) | `true`

Option to return the full wavelet packet tree, specified as the comma-separated pair consisting of `'FullTree'` and either `false` or `true`. If you specify `false`, then `modwpt` returns only the terminal (final-level) wavelet packet nodes. If you specify `true`, then `modwpt` returns the full wavelet packet tree down to the specified level.

Example: `'Fulltree', true`

TimeAlign — Signal time alignment

`false` (default) | `true`

Option to time align wavelet packet coefficients with signal features, specified as the comma-separated pair consisting of `'TimeAlign'` and either `true` to time align or `false` to not align.

The scaling and wavelet filters have a time delay. Circularly shifting the wavelet packet coefficients in all nodes aligns the signal and wavelet coefficients in time. If you want to reconstruct the signal, such as by using `imodwpt`, do not shift the coefficients because time alignment is done during the inversion process.

Example: `'TimeAlign', true`

Output Arguments

wpt — Wavelet packet transform

matrix

Wavelet packet tree, returned as a matrix with each row containing the sequency-ordered wavelet packet coefficients. By default, `wpt` contains only the terminal level for the MODWPT. The default terminal level is either level 4 or `floor(log2(numel(x)))`, whichever is smaller. At level 4, `wpt` is a 16-by-`numel(x)` matrix. For the full tree, at level j , `wpt` is a 2^{j+2} -by-`numel(x)` matrix, with each row containing the packet coefficients by level and index. The approximate passband for the n th row of `wpt` at level j is

$$\left[\frac{n-1}{2^{j+1}}, \frac{n}{2^{j+1}} \right) \text{ cycles/sample, where } n = 1, 2, \dots, 2^j.$$

packetlevs — Transform levels

vector

Transform levels, returned as a vector. The levels correspond to the rows of `wpt`. If `wpt` contains only the terminal level coefficients, `packetlevs` is a vector of constants equal to the terminal level. If `wpt` contains the full wavelet packet table, `packetlevs` is a vector with 2^j elements for each level, j . To select all the wavelet packet nodes at a particular level, use `packetlevs` with logical indexing.

cfreq — Center frequencies of passbands

vector

Center frequencies of the approximate passbands in the `wpt` rows, returned as a vector. The center frequencies are in cycles/sample. To convert the units to cycles/unit time, multiply `cfreq` by the sampling frequency.

energy — Energy of the wavelet packet coefficients

vector

Energy of the wavelet packet coefficients for the `wpt` nodes, returned as a vector. The sum of the energies (squared L2 norms) for the wavelet packets at each level equals the energy in the signal.

relenergy — Relative energy

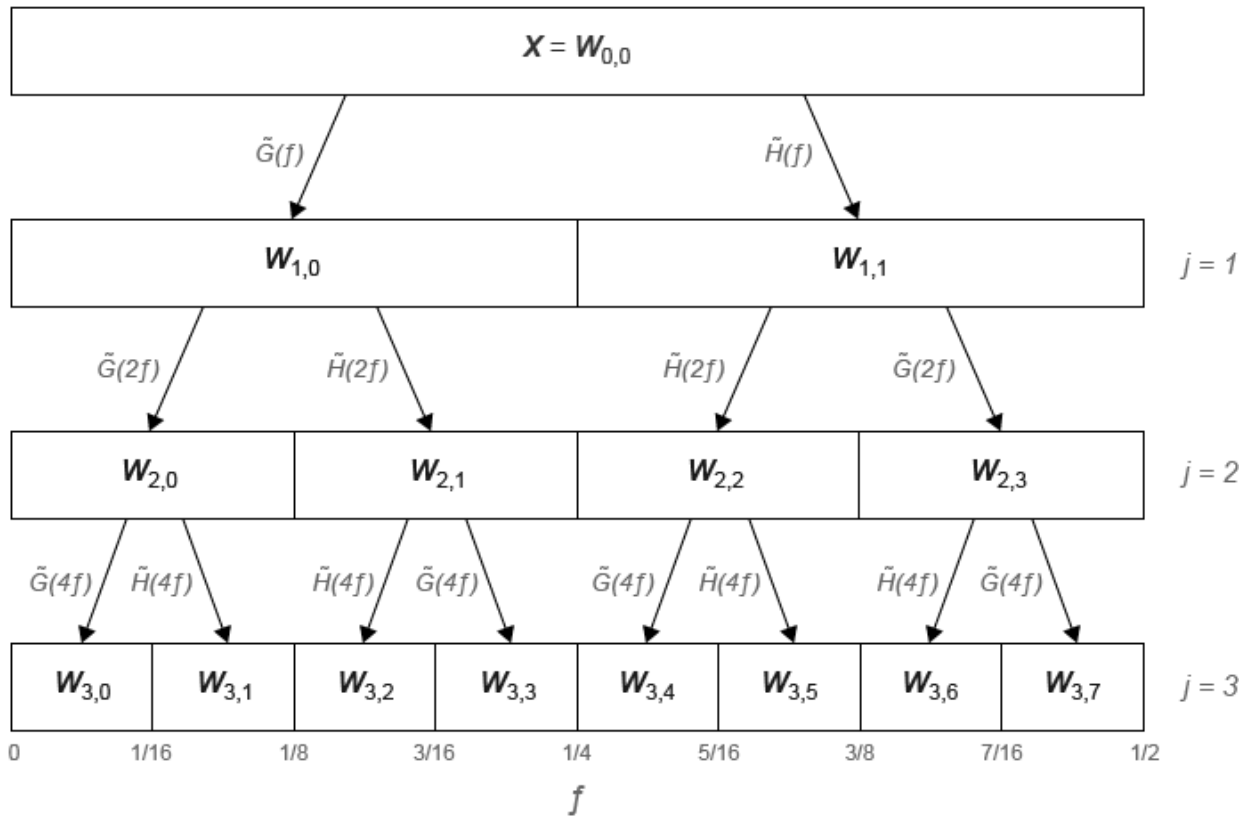
vector

Relative energy for each level, returned as a vector. The relative energy is the proportion of energy in each wavelet packet by level, relative to the total energy of that level. The sum of relative energies in all packets at each level equals 1.

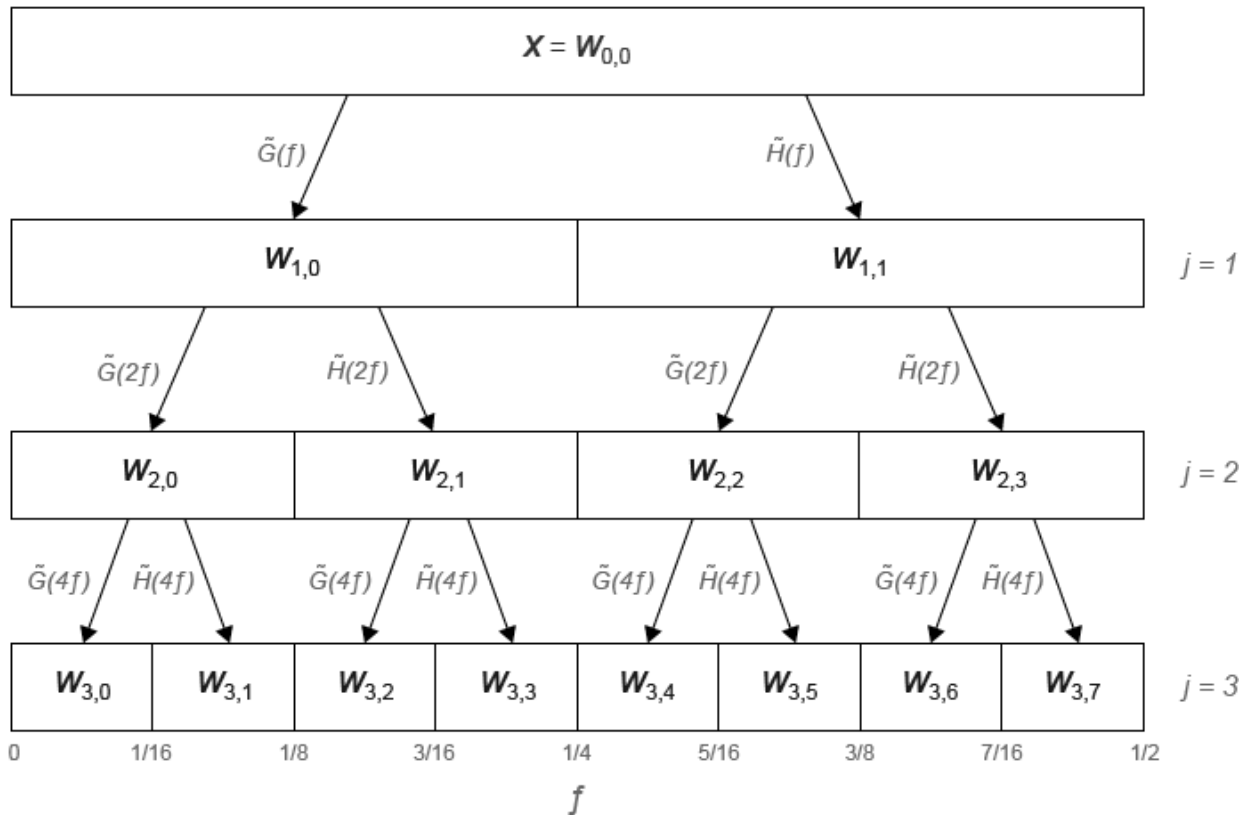
Algorithms

The `modwpt` performs a discrete wavelet packet transform and produces a sequency-ordered wavelet packet tree. Compare the sequency-ordered and normal (Paley)-ordered trees.

Sequency-Ordered Wavelet Packet Tree



Natural-Ordered Wavelet Packet Tree



References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.
- [2] Walden, A.T., and A. Contreras Cristan. "The phase-corrected undecimated discrete wavelet packet transform and its application to interpreting the timing of events." *Proceedings of the Royal Society of London A*. Vol. 454, Issue 1976, 1998, pp. 2243-2266.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`imodwpt` | `modwptdetails`

Introduced in R2016a

modwptdetails

Maximal overlap discrete wavelet packet transform details

Syntax

```
w = modwptdetails(x)
w = modwptdetails(x,wname)
w = modwptdetails(x,lo,hi)
w = modwptdetails( ____,lev)

[w,packetlevs] = modwptdetails( ____)
[w,packetlevs,cfreq] = modwptdetails( ____)

[ ____ ] = modwptdetails( ____, 'FullTree', tf)
```

Description

`w = modwptdetails(x)` returns the maximal overlap discrete wavelet packet transform (MODWPT) details for the 1-D real-valued signal, `x`. The MODWPT details provide zero-phase filtering of the signal. By default, `modwptdetails` returns only the terminal nodes, which are at level 4 or at level `floor(log2(numel(x)))`, whichever is smaller.

Note To decide whether to use `modwptdetails` or `modwpt`, consider the type of data analysis you need to perform. For applications that require time alignment, such as nonparametric regression analysis, use `modwptdetails`. For applications where you want to analyze the energy levels in different packets, use `modwpt`. For more information, see “Algorithms” on page 1-841

`w = modwptdetails(x,wname)` uses the orthogonal wavelet filter specified by `wname`.

`w = modwptdetails(x,lo,hi)` uses the orthogonal scaling filter, `lo`, and wavelet filter, `hi`.

`w = modwptdetails(____,lev)` returns the terminal nodes of the wavelet packet tree at positive integer level `lev`.

`[w,packetlevs] = modwptdetails(___)` returns a vector of transform levels corresponding to the rows of `w`.

`[w,packetlevs,cfreq] = modwptdetails(___)` returns `cfreq`, the center frequencies of the approximate passbands corresponding to the MODWPT details in `w`.

`[___] = modwptdetails(___ , 'FullTree', tf)`, where `tf` is `false`, returns details about only the terminal (final-level) wavelet packet nodes. If you specify `true`, then `modwptdetails` returns details about the full wavelet packet tree down to the default or specified level. The default for `tf` is `false`.

Examples

MODWPT Details Using Default Wavelet

Obtain the MODWPT of an electrocardiogram (ECG) signal using the default length 18 Fejer-Korovkin ('fk18') wavelet and the default level, 4.

```
load wecg;
wptdetails = modwptdetails(wecg);
```

Demonstrate that summing the MODWPT details over each sample reconstructs the signal. The largest absolute difference between the original signal and the reconstruction is on the order of 10^{-11} , which demonstrates perfect reconstruction.

```
xrec = sum(wptdetails);
max(abs(wecg-xrec'))
```

```
ans = 1.7903e-11
```

MODWPT Details for Two Sine Waves

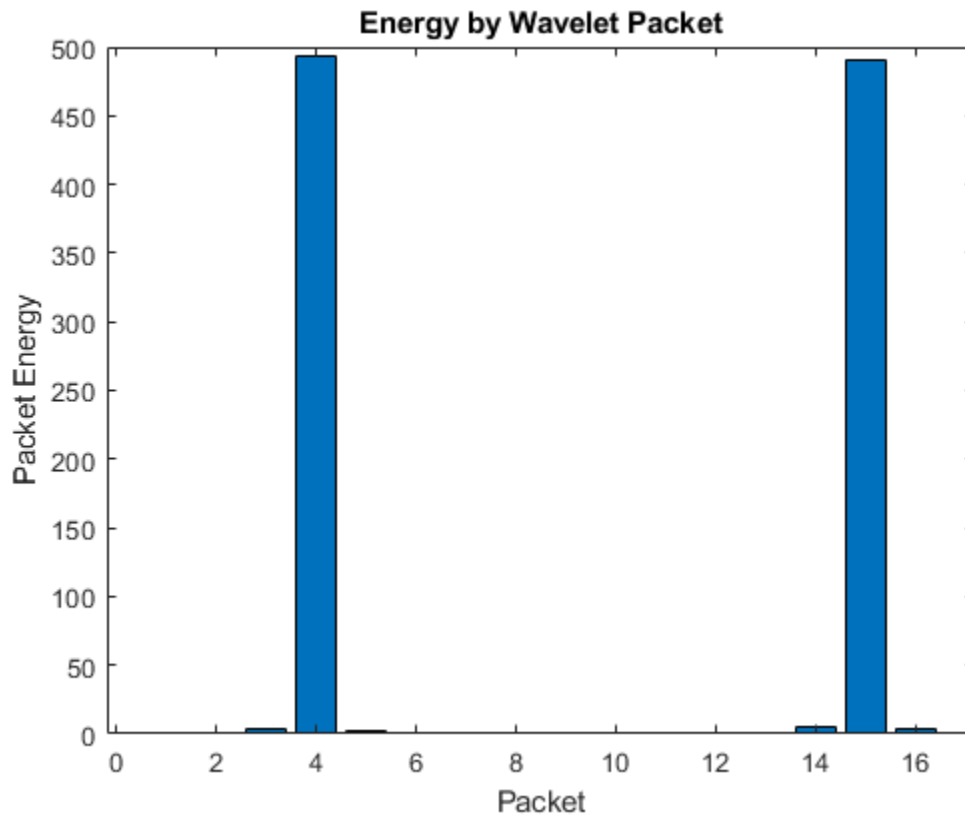
Obtain the MODWPT details for a signal containing 100 Hz and 450 Hz sine waves. Each row of the `modwptdetails` output corresponds to a separate frequency band.

```
dt = 0.001;
fs = 1/dt;
t = 0:dt:1;
```

```
x = (sin(2*pi*100*t)+sin(2*pi*450*t));  
[lo,hi] = wfilters('fk22');  
wptdetails = modwptdetails(x,lo,hi);
```

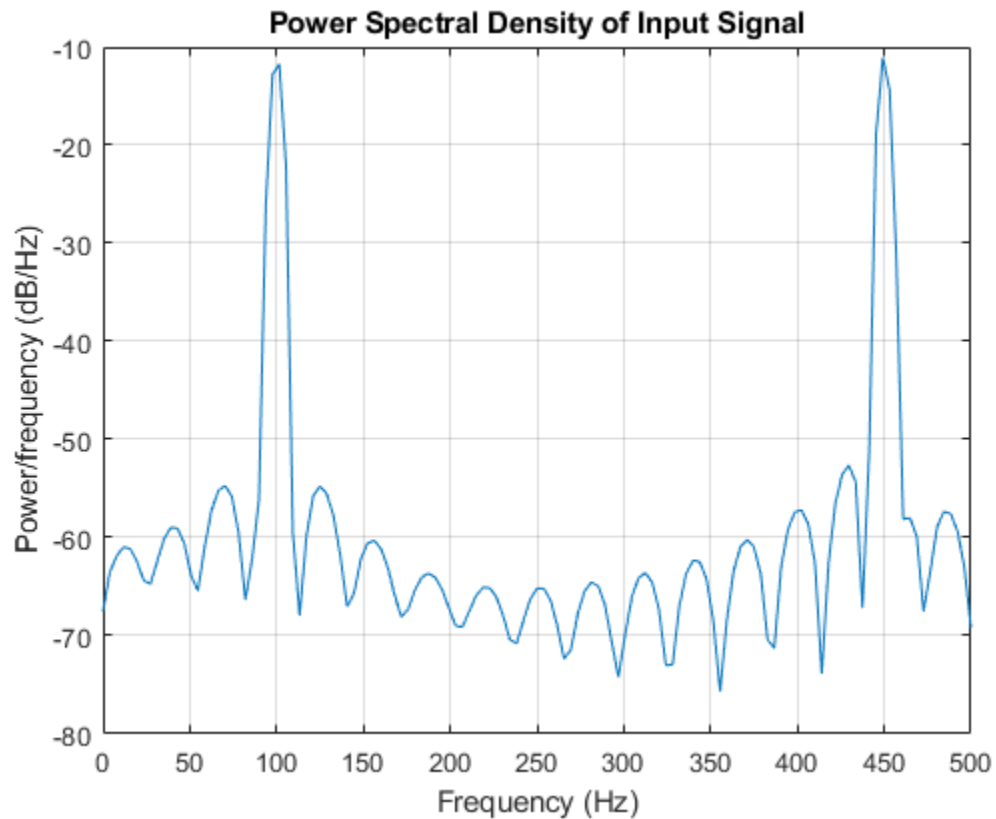
Use `modwpt` to obtain the energy and center frequencies of the signal. Plot the energy in the wavelet packets. The fourth and fifteenth frequency bands contain most of the energy. Other frequency bands have significantly less energy. The frequency ranges of fourth and fifteenth bands are approximately 94-125 Hz and 438-469 Hz, respectively.

```
[wpt,~,cfreqs,energy] = modwpt(x,lo,hi);  
figure  
bar(1:16,energy);  
xlabel('Packet')  
ylabel('Packet Energy')  
title('Energy by Wavelet Packet')
```



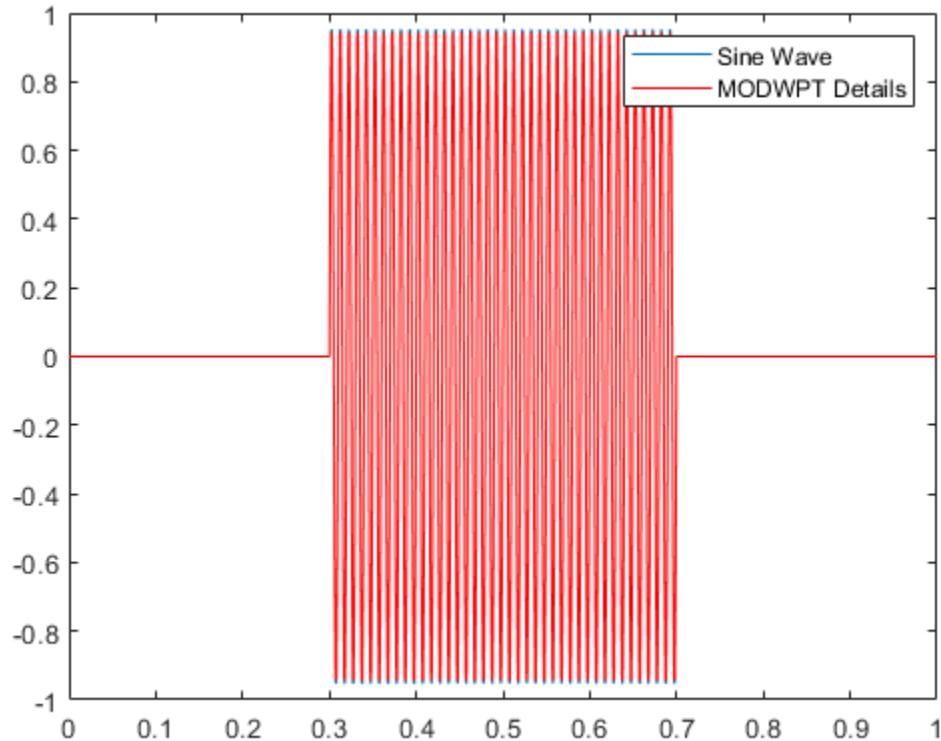
Plot the power spectral density of the input signal.

```
pwelch(x,[],[],[],fs,'onesided');  
title('Power Spectral Density of Input Signal')
```



Show that the MODWPT details have zero-phase shift from the 100 Hz input sine.

```
p4 = wptdetails(4,:);  
plot(t,sin(2*pi*100*t).*(t>0.3 & t<0.7))  
hold on  
plot(t,p4.*(t>0.3 & t<0.7),'r')  
legend('Sine Wave','MODWPT Details')
```

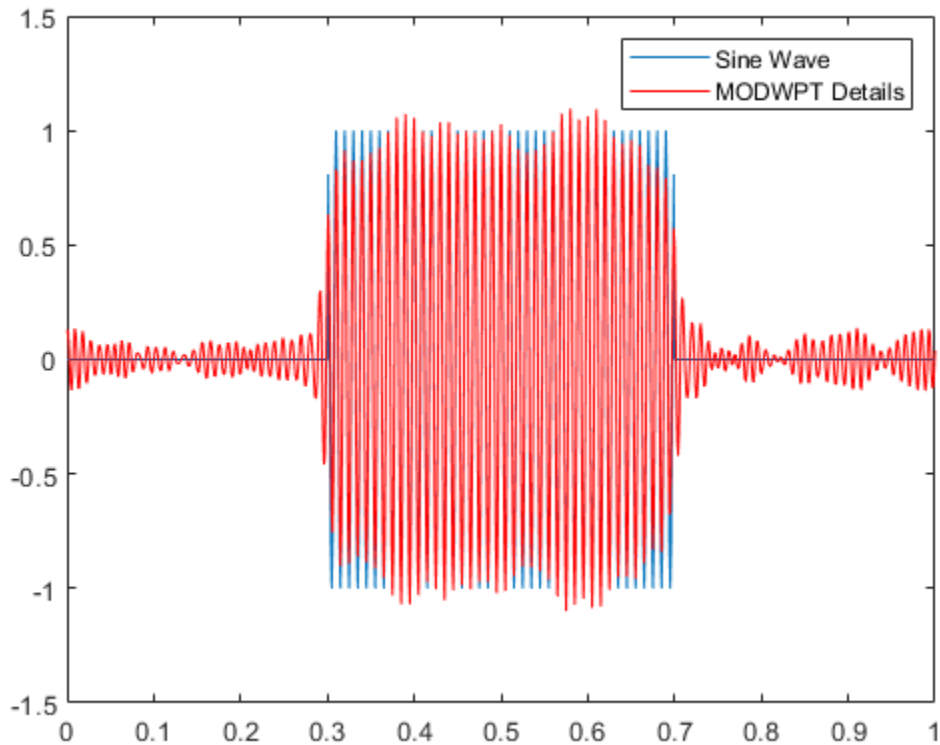
MODWPT Details for Noisy Sine Wave

Obtain the MODWPT details for a 100 Hz time-localized sine wave in noise. The sampling rate is 1000 Hz. Obtain the MODWPT at level 4 using the length 22 Fejer-Korovkin ('fk22') wavelet.

```
dt = 0.001;
t = 0:dt:1;
x = cos(2*pi*100*t).*(t>0.3 & t<0.7)+0.25*randn(size(t));
wptdetails = modwptdetails(x,'fk22');
p4 = wptdetails(4,:);
```

Plot the MODWPT details for level 4, packet number 4. The MODWPT details represent zero-phase filtering of the input signal with an approximate passband of $[3F_s/2^5, 4F_s/2^5)$, where F_s is the sampling frequency.

```
plot(t,cos(2*pi*100*t).*(t>0.3 & t<0.7));  
hold on  
plot(t,p4,'r')  
legend('Sine Wave','MODWPT Details')
```



MODWPT Details Using Scaling and Wavelet Filters

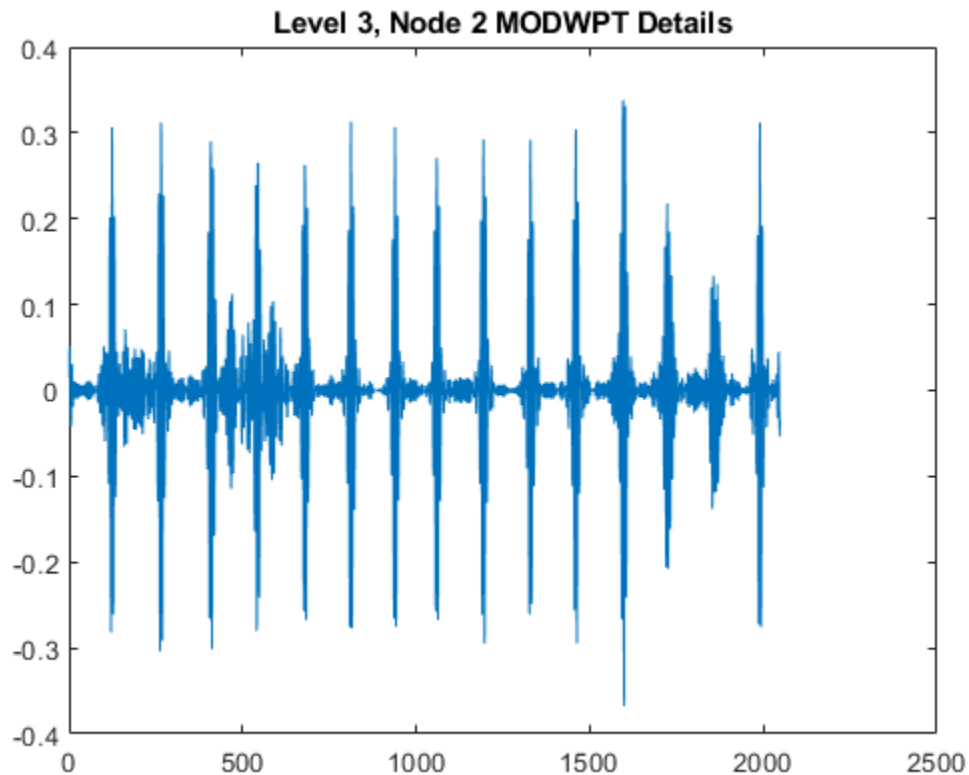
Obtain the MODWPT details of an ECG waveform using the length 18 Fejer-Korovkin scaling and wavelet filters.

```
load wecg;  
[lo,hi] = wfilters('fk18');  
wpt = modwptdetails(wecg,lo,hi);
```

MODWPT Details for Full Packet Tree

Obtain the MODWPT details for the full wavelet packet tree of an ECG waveform. Use the default length 18 Fejer-Korovkin ('fk18') wavelet. Extract and plot the node coefficients at level 3, node 2.

```
load wecg;  
[w,packetlevels] = modwptdetails(wecg,'FullTree',true);  
p3 = w(packetlevels==3,:);  
plot(p3(3,:))  
title('Level 3, Node 2 MODWPT Details')
```



Input Arguments

x — Input signal

real-valued vector

Input signal, specified as a real-valued row or column vector. `x` must have at least two elements.

Data Types: double

wname — Analyzing wavelet filter

'fk18' (default) | 'dbN' | 'coifN' | 'haar' | 'fkN' | 'symN'

Analyzing wavelet, specified as one of the following:

- 'haar' — Haar wavelet
- 'dbN' — Extremal phase Daubechies wavelet with N vanishing moments, where N is a positive integer from 1 to 45.
- 'symN' — Symlets wavelet with N vanishing moments, where N is a positive integer from 2 to 45.
- 'coifN' — Coiflets wavelet with N vanishing moments, where N is a positive integer from 1 to 5.
- 'fkN' — Fejér-Korovkin wavelet with N coefficients, where $N = 4, 6, 8, 14, 18$ and 22.

lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. `lo` must satisfy the conditions necessary to generate an orthogonal scaling function. You can specify the `lo` and `hi` scaling-wavelet filter pair only if you do not specify `wname`.

hi — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. `hi` must satisfy the conditions necessary to generate an orthogonal wavelet. You can specify the `lo` and `hi` scaling-wavelet filter pair only if you do not specify `wname`.

lev — Transform level

positive integer

Transform level, specified as a positive integer less than or equal to $\text{floor}(\log_2(\text{numel}(x)))$.

tf — Return tree option

false (default) | true

Return tree option, specified as `false` or `true`. If `tf` is `false`, then `modwptdetails` returns details about only the terminal (final-level) wavelet packet nodes. If you specify `true`, then `modwptdetails` returns details about the full wavelet packet tree down to the default or specified level.

For the full wavelet packet tree, w is a $2^{j+1} \times 2$ -by- $\text{numel}(x)$ matrix. Each level j has 2^j wavelet packet details.

Output Arguments

w — Wavelet packet tree details

matrix

Wavelet packet tree details, returned as a matrix with each row containing the sequentially ordered wavelet packet details for the terminal nodes. The terminal nodes are at level 4 or at level $\text{floor}(\log_2(\text{numel}(x)))$, whichever is smaller. The MODWPT details are zero-phase-filtered projections of the signal onto the subspaces corresponding to the wavelet packet nodes. The sum of the MODWPT details over each sample reconstructs the original signal.

For the default terminal nodes, w is a 2^j -by- $\text{numel}(x)$ matrix. For the full packet table, at level j , w is a $2^{j+1} \times 2$ -by- $\text{numel}(x)$ matrix of sequentially ordered wavelet packet coefficients by level and index. The approximate passband for the n th row of w at level j is

$\left[\frac{n-1}{2^{(j+1)}}, \frac{n}{2^{(j+1)}} \right)$ cycles per sample, where $n = 1, 2, \dots, 2^j$.

packetLevs — Transform levels

vector

Transform levels, returned as a vector. The levels correspond to the rows of w . If w contains only the terminal level coefficients, **packetLevs** is a vector of constants equal to the terminal level. If w contains the full wavelet packet tree of details, **packetLevs** is a vector with 2^{j-1} elements for each level, j . To select all the MODWPT details at a particular level, use **packetLevs** with logical indexing.

cfreq — Center frequencies of passbands

vector

Center frequencies of the approximate passbands in the w rows, returned as a vector. The center frequencies are in cycles per sample. To convert the units to cycles per unit time, multiply **cfreq** by the sampling frequency.

Algorithms

The MODWPT details (`modwptdetails`) are the result of zero-phase filtering of the signal. The features in the MODWPT details align exactly with features in the input signal. For a given level, summing the details for each sample returns the exact original signal.

The output of the MODWPT (`modwpt`) is time delayed compared to the input signal. Most filters used to obtain the MODWPT have a nonlinear phase response, which makes compensating for the time delay difficult. All orthogonal scaling and wavelet filters have this response, except the Haar wavelet. It is possible to time align the coefficients with the signal features, but the result is an approximation, not an exact alignment with the original signal. The MODWPT partitions the energy among the wavelet packets at each level. The sum of the energy over all the packets equals the total energy of the input signal.

References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.
- [2] Walden, A.T., and A. Contreras Cristan. "The phase-corrected undecimated discrete wavelet packet transform and its application to interpreting the timing of events." *Proceedings of the Royal Society of London A*. Vol. 454, Issue 1976, 1998, pp. 2243-2266.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`imodwpt` | `modwpt` | `waveinfo` | `wavemngr`

Introduced in R2016a

modwt

Maximal overlap discrete wavelet transform

Syntax

```
w = modwt(x)
w = modwt(x,wname)
w = modwt(x,Lo,Hi)
w = modwt( ____,lev)
w = modwt( ____, 'reflection')
```

Description

`w = modwt(x)` returns the maximal overlap discrete wavelet transform (MODWT) of the 1-D real-valued signal, `x`.

`w = modwt(x,wname)` uses the orthogonal wavelet, `wname`, for the MODWT.

`w = modwt(x,Lo,Hi)` uses the scaling filter, `Lo`, and wavelet filter, `Hi`, to compute the MODWT. These filters must satisfy the conditions for an orthogonal wavelet. You cannot specify `Lo` and `Hi` if you specify `wname`.

`w = modwt(____,lev)` computes the MODWT down to the specified level, `lev`, using any of the arguments from previous syntaxes.

`w = modwt(____, 'reflection')` computes the MODWT using reflection boundary handling. Other inputs can be any of the arguments from previous syntaxes. Before computing the wavelet transform, `modwt` extends the signal symmetrically at the right boundary to twice the signal length, [`x flip(x)`]. The number of wavelet and scaling coefficients that `modwt` returns is equal to twice the length of the input signal. By default, the signal is extended periodically.

Examples

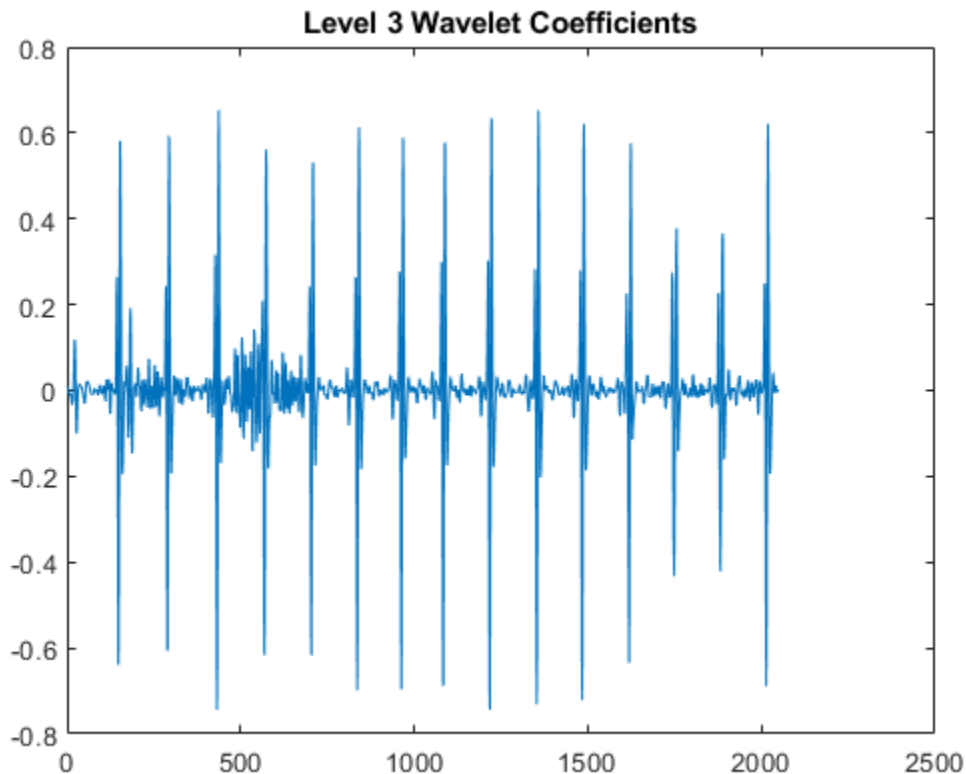
MODWT Using Default Wavelet

Obtain the MODWT of an electrocardiogram (ECG) signal using the default sym4 wavelet down to the maximum level.

```
load wecg;  
wtecg = modwt(wecg);
```

wtecg is 12-by-2048 matrix. The first eleven rows are the wavelet coefficients for scales 2^1 to 2^{11} . The final row contains the scaling coefficients at scale 2^{11} . Plot the detail (wavelet) coefficients for scale 2^3 .

```
plot(wtecg(3,:))  
title('Level 3 Wavelet Coefficients')
```



MODWT Using Daubechies Extremal Phase Wavelet with Two Vanishing Moments

Obtain the MODWT of Southern Oscillation Index data with the 'db2' wavelet down to the maximum level.

```
load soi;
wsoi = modwt(soi, 'db2');
```

MODWT Using Scaling and Wavelet Filters

Obtain the MODWT of the Deutsche Mark - U.S. Dollar exchange rate data using the Fejer-Korovkin length 8 scaling and wavelet filters.

```
load DM_USD;
[Lo,Hi] = wfilters('fk8');
wdm = modwt(DM_USD,Lo,Hi);
```

MODWT to a Specified Level

Obtain the MODWT of an ECG signal down to scale 2^4 , which corresponds to level four. Use the default 'sym4' wavelet.

```
load wecg;
wtecg = modwt(wecg,4);
```

wtecg is a 5-by-2048 matrix. The row size is L+1, where, in this case, the level (L) is 4. The column size matched the number of input samples.

MODWT with Reflection Boundary

Obtain the MODWT of an ECG signal using reflection boundary handling. Use the default 'sym4' wavelet and obtain the transform down to level 4.

```
load wecg;  
wtecg = modwt(wecg,4,'reflection');
```

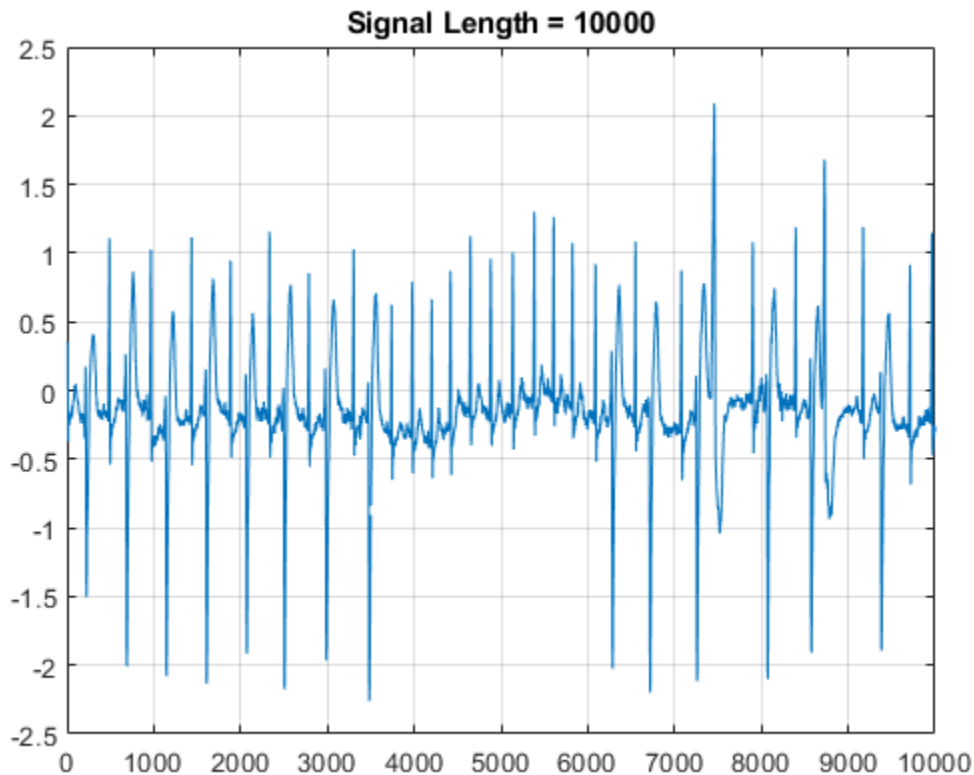
wtecg has 4096 columns, which is twice the length of the input signal, wecg.

Comparing MODWT and MODWTMRA

This example demonstrates the differences between the functions MODWT and MODWTMRA. The MODWT partitions a signal's energy across detail coefficients and scaling coefficients. The MODWTMRA projects a signal onto wavelet subspaces and a scaling subspace.

Choose the 'sym6' wavelet. Load and plot an ECG waveform. The ECG data is taken from the MIT-BIH Arrhythmia Database.

```
load mit200  
wv = 'sym6';  
plot(ecgsig)  
grid on  
title(['Signal Length = ',num2str(length(ecgsig))])
```



Take the MODWT of the signal.

```
wtecg = modwt(ecgsig,wv);
```

The input data are samples of a function $f(x)$ evaluated at N -many time points. The function can be expressed as a linear combination of the scaling function $\phi(x)$ and wavelet

$\psi(x)$ at varying scales and translations: $f(x) = \sum_{k=0}^{N-1} c_k 2^{-J_0/2} \phi(2^{-J_0} x - k) + \sum_{j=1}^{J_0} f_j(x)$ where

$f_j(x) = \sum_{k=0}^{N-1} d_{j,k} 2^{-j/2} \psi(2^{-j} x - k)$ and J_0 is the number of levels of wavelet decomposition.

The first sum is the coarse scale approximation of the signal, and the $f_j(x)$ are the details

at successive scales. MODWT returns the N -many coefficients $\{c_k\}$ and the $(J_0 \times N)$ -many detail coefficients $\{d_{j,k}\}$ of the expansion. Each row in `wtecg` contains the coefficients at a different scale.

When taking the MODWT of a signal of length N , there are $\text{floor}(\log_2(N))$ -many levels of decomposition (by default). Detail coefficients are produced at each level. Scaling coefficients are returned only for the final level. In this example, since $N = 10000$, $J_0 = \text{floor}(\log_2(10000)) = 13$ and the number of rows in `wtecg` is $J_0 + 1 = 13 + 1 = 14$.

The MODWT partitions the energy across the various scales and scaling coefficients:

$\|X\|^2 = \sum_{j=1}^{J_0} \|W_j\|^2 + \|V_{J_0}\|^2$ where X is the input data, W_j are the detail coefficients at scale j , and V_{J_0} are the final-level scaling coefficients.

Compute the energy at each scale, and evaluate their sum.

```
energy_by_scales = sum(wtecg.^2,2);
Levels = {'D1';'D2';'D3';'D4';'D5';'D6';'D7';'D8';'D9';'D10';'D11';'D12';'D13';'A13'};
energy_table = table(Levels,energy_by_scales);
disp(energy_table)
```

Levels	energy_by_scales
'D1'	0.31592
'D2'	2.6504
'D3'	28.802
'D4'	159.37
'D5'	300.5
'D6'	431.33
'D7'	444.93
'D8'	182.37
'D9'	45.381
'D10'	11.578
'D11'	19.809
'D12'	4.5406
'D13'	3.308
'A13'	192.46

```
energy_total = varfun(@sum,energy_table(:,2))
```

```
energy_total=1x1 table
sum_energy_by_scales
```

1827.3

Confirm the MODWT is energy-preserving by computing the energy of the signal and comparing it with the sum of the energies over all scales.

```
energy_ecg = sum(ecgsig.^2);  
max(abs(energy_total.sum_energy_by_scales-energy_ecg))
```

```
ans = 4.0870e-09
```

Take the MODWTMRA of the signal.

```
mraecg = modwtmra(wtecg,wv);
```

MODWTMRA returns the projections of the function $f(x)$ onto the various wavelet subspaces and final scaling space. That is, MODWTMRA returns

$\sum_{k=0}^{N-1} c_k 2^{-J_0/2} \phi(2^{-J_0} x - k)$ and the J_0 -many $\{f_j(x)\}$ evaluated at N -many time points. Each

row in `mraecg` is a projection of $f(x)$ onto a different subspace. This means the original signal can be recovered by adding all the projections. This is *not* true in the case of the MODWT. Adding the coefficients in `wtecg` will *not* recover the original signal.

Choose a time point, add the projections of $f(x)$ evaluated at that time point and compare with the original signal.

```
time_point = 1000;  
abs(sum(mraecg(:,time_point))-ecgsig(time_point))
```

```
ans = 3.0970e-13
```

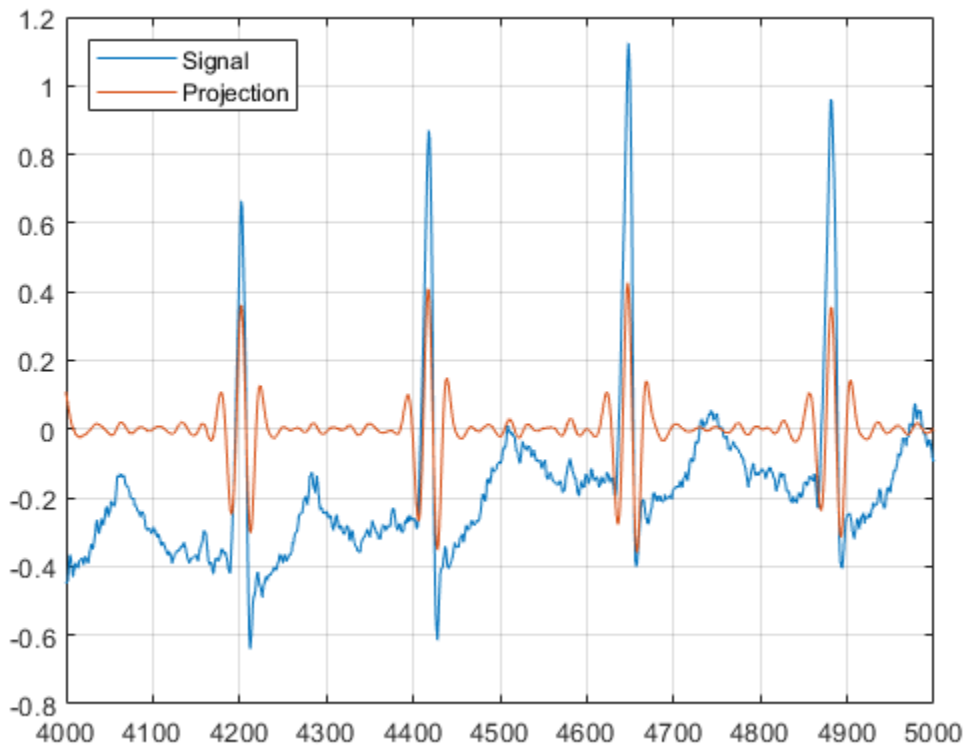
Confirm that, unlike MODWT, MODWTMRA is not an energy-preserving transform.

```
energy_ecg = sum(ecgsig.^2);  
energy_mra_scales = sum(mraecg.^2,2);  
energy_mra = sum(energy_mra_scales);  
max(abs(energy_mra-energy_ecg))
```

```
ans = 534.7949
```

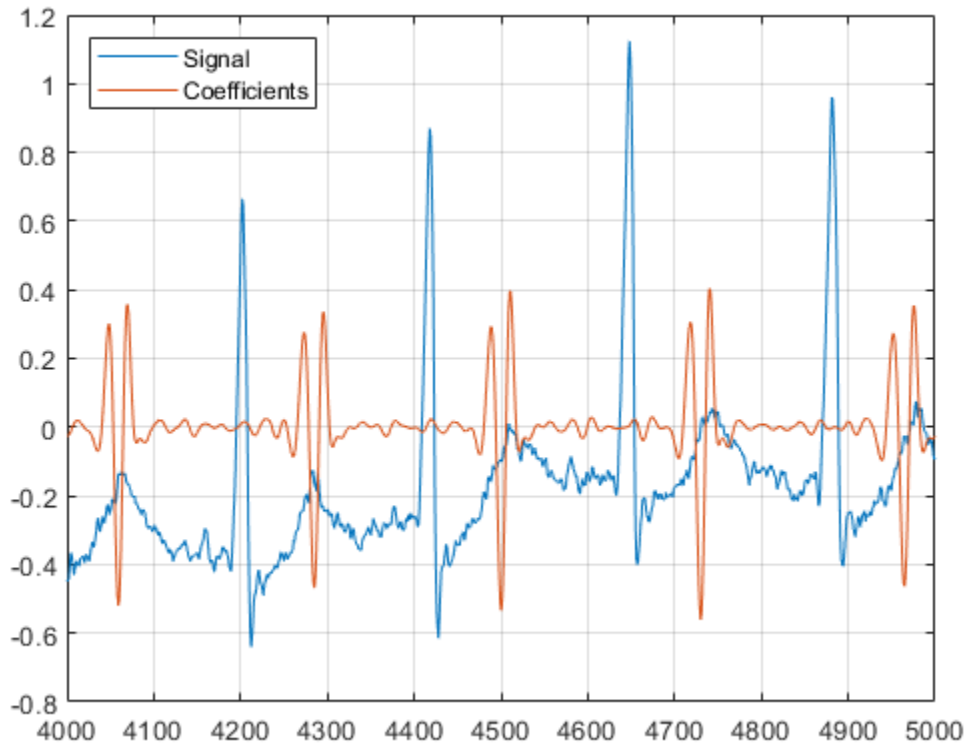
The MODWTMRA is a zero-phase filtering of the signal. Features will be time-aligned. Demonstrate this by plotting the original signal and one of its projections. To better illustrate the alignment, zoom in.

```
figure
plot(ecgsig)
hold on
plot(mraecg(4,:), '-')
grid on
xlim([4000 5000])
legend('Signal', 'Projection', 'Location', 'northwest')
```



Make a similar plot using the MODWT coefficients at the same scale. Note that features will not be time-aligned. The MODWT is *not* a zero-phase filtering of the input.

```
figure
plot(ecgsig)
hold on
plot(wtecg(4,:), '-')
grid on
xlim([4000 5000])
legend('Signal', 'Coefficients', 'Location', 'northwest')
```



References

Goldberger A. L., L. A. N. Amaral, L. Glass, J. M. Hausdorff, P. Ch. Ivanov, R. G. Mark, J. E. Mietus, G. B. Moody, C-K Peng, H. E. Stanley. "PhysioBank, PhysioToolkit, and PhysioNet: Components of a New Research Resource for Complex Physiologic Signals." *Circulation*

101. Vol.23, e215-e220, 2000. <http://circ.ahajournals.org/cgi/content/full/101/23/e215>

Moody, G. B. "Evaluating ECG Analyzers". <http://www.physionet.org/physiotools/wfdb/doc/wag-src/eval0.tex>

Moody G. B., R. G. Mark. "The impact of the MIT-BIH Arrhythmia Database." IEEE Eng in Med and Biol. Vol. 20, Number 3, 2001), pp. 45-50 .

Input Arguments

x — Input signal

real-valued vector

Input signal, specified as a row or column vector. `x` must have at least two elements.

By default, `modwt` computes the wavelet transform down to level `floor(log2(length(x)))` using the Daubechies least-asymmetric wavelet with four vanishing moments ('`sym4`') and periodic boundary handling.

Data Types: `double`

wname — Analyzing wavelet

'`sym4`' (default) | '`haar`' | '`dbN`' | '`symN`' | '`coifN`' | '`fkN`'

Analyzing wavelet, specified as one of the following:

- '`haar`' — Haar wavelet
- '`dbN`' — Extremal phase Daubechies wavelet with `N` vanishing moments, where `N` is a positive integer from 1 to 45.
- '`symN`' — Symlets wavelet with `N` vanishing moments, where `N` is a positive integer from 2 to 45.
- '`coifN`' — Coiflets wavelet with `N` vanishing moments, where `N` is a positive integer from 1 to 5.
- '`fkN`' — Fejér-Korovkin wavelet with `N` coefficients, where `N` = 4, 6, 8, 14, 18 and 22.

Lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. L_0 must satisfy the conditions necessary to generate an orthogonal scaling function. You can specify L_0 only if you do not specify $wname$.

 H_i — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. H_i must satisfy the conditions necessary to generate an orthogonal wavelet. You can specify H_i only if you do not specify $wname$.

 L_{ev} — Transform level

positive integer

Transform level, specified as a positive integer less than or equal to $\text{floor}(\log_2(\text{length}(x)))$.

Output Arguments

 w — Wavelet transform

matrix

Wavelet transform w , returned as an $L+1$ -by- N matrix containing wavelet coefficients and final-level scaling coefficients. L is the level of the MODWT. N is equal to the input signal length unless you specify 'reflection' boundary handling, in which case N is twice the length of the input signal. The k th row of w contains the wavelet coefficients for scale 2^k (wavelet scale $2^{(k-1)}$). The final, $(L+1)$ th, row of w contains the scaling coefficients for scale 2^L .

Algorithms

The standard algorithm for the MODWT implements the circular convolution directly in the time domain. This implementation of the MODWT performs the circular convolution in the Fourier domain. The wavelet and scaling filter coefficients at level j are computed by taking the inverse discrete Fourier transform (DFT) of a product of DFTs. The DFTs in the product are the signal's DFT and the DFT of the j^{th} level wavelet or scaling filter.

Let H_k and G_k denote the length N DFTs of the MODWT wavelet and scaling filters, respectively. Let j denote the level and N denote the sample size.

The j^{th} level wavelet filter is defined by

$$\frac{1}{N} \sum_{k=0}^{N-1} H_{j,k} e^{i2\pi nk/N}$$

where

$$H_{j,k} = H_{2^{j-1}k \bmod N} \prod_{m=0}^{j-2} G_{2^m k \bmod N}$$

The j^{th} level scaling filter is

$$\frac{1}{N} \sum_{k=0}^{N-1} G_{j,k} e^{i2\pi nk/N}$$

where

$$G_{j,k} = \prod_{m=0}^{j-1} G_{2^m k \bmod N}$$

References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.
- [2] Percival, D. B., and H. O. Mofjeld. "Analysis of subtidal coastal sea level fluctuations using wavelets." *Journal of the American Statistical Association*. Vol. 92, pp 868-880.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`imodwt` | `modwtcorr` | `modwtmra` | `modwtvar` | `modwtxcorr` | `waveinfo` | `wavemngr`

Topics

“Wavelet Analysis of Financial Data”

Introduced in R2015b

modwtcorr

Multiscale correlation using the maximal overlap discrete wavelet transform

Syntax

```
wcorr = modwtcorr(w1,w2)
wcrr = modwtcorr(w1,w2,wav)

[wcrr,wcrrci] = modwtcorr( ___ )
[wcrr,wcrrci] = modwtcorr( ___ ,conflvl)
[wcrr,wcrrci,pval] = modwtcorr( ___ )
[wcrr,wcrrci,pval,nj] = modwtcorr( ___ )

wcrrtbl = modwtcorr( ___ ,'table')

[ ___ ] = modwt( ___ ,'reflection')

modwtcorr( ___ )
```

Description

`wcorr = modwtcorr(w1,w2)` returns the wavelet correlation by scale for the maximal overlap discrete wavelet transforms (MODWTs) specified in `w1` and `w2`. `wcorr` is an M -by-1 vector of correlation coefficients, where M is the number of levels with nonboundary wavelet coefficients. If the final level has enough nonboundary coefficients, `modwtcorr` returns the scaling correlation in the final row of `wcorr`.

`wcorr = modwtcorr(w1,w2,wav)` uses the wavelet `wav` to determine the number of boundary coefficients by level.

`[wcrr,wcrrci] = modwtcorr(___)` returns in `wcrrci` the lower and upper 95% confidence bounds for the correlation coefficients of `wcorr`, using any arguments from the previous syntaxes.

`[wcrr,wcrrci] = modwtcorr(___ ,conflvl)` uses `conflvl` for the coverage probability of the confidence interval. `conflvl` is a real scalar strictly

greater than 0 and less than 1. If `conflevel` is unspecified or specified as empty, the coverage probability defaults to 0.95.

`[wcorr,wcorrci,pval] = modwtcorr(___)` returns the p-values for the null hypothesis test that the correlation coefficient in `wcorr` is equal to zero. `pval` is an M -by-2 matrix, where M is the number of levels with nonboundary wavelet coefficients. `T`

`[wcorr,wcorrci,pval,nj] = modwtcorr(___)` returns the number of nonboundary coefficients used in the computation of the correlation estimates by level, `nj`.

`wcorrtable = modwtcorr(___, 'table')` returns an M -by-6 table with the correlation, confidence bounds, p-value, and adjusted p-value. The table also lists the number of nonboundary coefficients by level. The row names of the table `wcorrtable` designate the type and level of each estimate. For example, `D1` designates that the row corresponds to a wavelet or detail estimate at level 1 and `S6` designates that the row corresponds to the scaling estimate at level 6. The scaling correlation is only computed for the final level of the MODWT and only when there are nonboundary scaling coefficients. You can specify the 'table' flag anywhere after the input transforms `w1` and `w2`. You must enter the entire character vector 'table'. If you specify 'table', `modwtcorr` only outputs one argument.

`[___] = modwt(___, 'reflection')` reduces the number of wavelet and scaling coefficients at each scale by half before computing the correlation. Use this option only when you obtain the MODWT of `w1` and `w2` were obtained using the 'reflection' boundary condition. You must enter the entire character vector 'reflection'. If you added a wavelet named 'reflection' using the wavelet manager, you must rename that wavelet prior to using this option.

`modwtcorr` supports only unbiased estimates of the wavelet correlation. For these estimates, the algorithm must removed the extra coefficients obtained using the 'reflection' boundary condition. Specifying the 'reflection' option in `modwtcorr` is identical to first obtaining the MODWT of `w1` and `w2` using the default 'periodic' boundary handling and then computing the wavelet correlation estimates.

`modwtcorr(___)` with no output arguments plots the wavelet correlations by scale with lower and upper confidence bounds. By default, the coverage probability is 0.95. Scales with NaNs for the confidence bounds and the scaling correlation are excluded.

Examples

Correlation by Scale

Find the correlation by scale for monthly DM-USD exchange rate returns from 1970 to 1998. The return data are log transformed. Use the Daubechies wavelet with two vanishing moments ('db2') to obtain the MODWT down to level 6. Then obtain the correlation data.

```
load DM_USD;
load JY_USD;
wdm = modwt(DM_USD, 'db2', 6);
wjy = modwt(JY_USD, 'db2', 6);
wcorr = modwtcorr(wdm, wjy, 'db2')
```

```
wcorr = 7×1
```

```
    0.5854
    0.5748
    0.6264
    0.4948
    0.3787
    0.9072
    0.7976
```

wcorr contains seven elements. The first six elements are the correlation coefficients for the wavelet (detail) levels one to six. The final element is the correlation for the scaling (lowpass) level six.

Multiscale Correlation

Obtain the MODWT of the Southern Oscillation Index and Truk Island daily pressure data sets. Tabulate the correlation between the two data sets by level.

```
load soi;
load truk;
wsoi = modwt(soi);
wtruk = modwt(truk);
wcorr = modwtcorr(wsoi, wtruk)
```

```
wcorr = 10×1
```

```
    0.1749
```

```
0.2936
0.0914
0.0883
0.2667
0.0894
-0.0415
0.4825
0.4394
0.7433
```

Show that the number of nonboundary coefficients, in this case, is less than the maximal length of the input. The MODWT is computed down to level thirteen, which is the maximal level for the length of the input. Level thirteen contains thirteen wavelet coefficient vectors and one scaling coefficient vector.

```
size(wsoi,1)
```

```
ans = 14
```

The multiscale correlations are computed only down to level ten because the levels after that do not contain nonboundary coefficients. For unbiased estimates, you must use nonboundary coefficients only.

```
numel(wcorr)
```

```
ans = 10
```

Confidence Intervals for Correlation

Obtain the MODWT of the monthly US-DM and US-JPY exchange return data from 1970 to 1998. The return data are log transformed. Use the Daubechies wavelet with two vanishing moments ('db2') and obtain the MODWT of each series down to level six. Obtain the correlation estimates by scale and the 95% confidence intervals.

```
load DM_USD
load JY_USD
wdm = modwt(DM_USD, 'db2', 6);
wjy = modwt(JY_USD, 'db2', 6);
[wcorr, wcorrci] = modwtcorr(wdm, wjy, 'db2');
[wcorr wcorrci]
```



```
ans = 7×3

    0.5854    0.4780    0.6756
    0.5748    0.4133    0.7013
    0.6264    0.4016    0.7800
    0.4948    0.0803    0.7634
    0.3787   -0.3295    0.8142
    0.9072    0.1247    0.9939
    0.7976   -0.2857    0.9860
```

The width of the confidence interval increases as you go down in level.

Confidence Intervals with 0.99 Coverage Probability

Specify the coverage probability for the confidence intervals. Obtain the 99% confidence intervals for the US-DM and US-JY exchange returns.

```
load DM_USD;
load JY_USD;
wdm = modwt(DM_USD, 'db2', 6);
wjy = modwt(JY_USD, 'db2', 6);
[wcorr, wcorrci] = modwtcorr(wdm, wjy, 'db2', 0.99);
[wcorr wcorrci]
```

```
ans = 7×3

    0.5854    0.4407    0.7005
    0.5748    0.3557    0.7340
    0.6264    0.3169    0.8153
    0.4948   -0.0646    0.8176
    0.3787   -0.5191    0.8792
    0.9072   -0.3006    0.9975
    0.7976   -0.6227    0.9941
```

P-values for Correlation

Return p -values for the test of zero correlation by scale. Obtain the MODWT of the DM-USD and JY-USD exchange return data down to level six using the Daubechies wavelet

with two vanishing moments ('db2') wavelet. Compute the correlation by scale and return the p -values.

```
load DM_USD;
load JY_USD;
wdm = modwt(DM_USD, 'db2', 6);
wjy = modwt(JY_USD, 'db2', 6);
[wcorr, wcorrci, pval] = modwtcorr(wdm, wjy, 'db2');
format longE
pval

pval = 7x2

    2.694174887029593e-17    4.889927419958712e-16
    7.125460513473891e-09    6.466355415977556e-08
    7.012389783536512e-06    4.242495819039590e-05
    2.258540027996925e-02    1.024812537703605e-01
    2.805930327935258e-01    7.275376493146417e-01
    3.348079529469842e-02    1.215352869197552e-01
    1.059217509938027e-01    3.204132967562530e-01
```

format

The first column contains the p -value and the second column contains the adjusted p -value based on the false discovery rate.

Multiscale Correlation in Tabular Form

Output results from `modwtcorr` in tabular form. Obtain the MODWT of the DM-USD and JY-USD exchange returns down to level six using the Daubechies wavelet with two vanishing moments ('db2'). Output the results in a table.

```
load DM_USD;
load JY_USD;
wdm = modwt(DM_USD, 'db2', 6);
wjy = modwt(JY_USD, 'db2', 6);
corrtable = modwtcorr(wdm, wjy, 'db2', 'table')

corrtable=7x6 table
      NJ      Lower      Rho      Upper      Pvalue      AdjustedPvalue
      _____
```

D1	344	0.47797	0.58542	0.67561	2.6942e-17	4.8899e-16
D2	338	0.41329	0.57483	0.70129	7.1255e-09	6.4664e-08
D3	326	0.40163	0.62641	0.78001	7.0124e-06	4.2425e-05
D4	302	0.080255	0.4948	0.76342	0.022585	0.10248
D5	254	-0.32954	0.37865	0.81417	0.28059	0.72754
D6	158	0.12469	0.90716	0.99393	0.033481	0.12154
S6	158	-0.28573	0.79761	0.98601	0.10592	0.32041

Correlation with Reflection Boundary Conditions

Obtain multiscale correlation estimates when using 'reflection' boundary handling. Obtain the MODWT of the Southern Oscillation Index and Truk Islands pressure data sets using 'reflection' boundary handling for both data sets.

```
load soi
load truk
wsoi = modwt(soi,'fk4',6,'reflection');
wtruk = modwt(truk,'fk4',6,'reflection');
corrtable = modwtcorr(wsoi,wtruk,'fk4',0.95,'reflection','table')
```

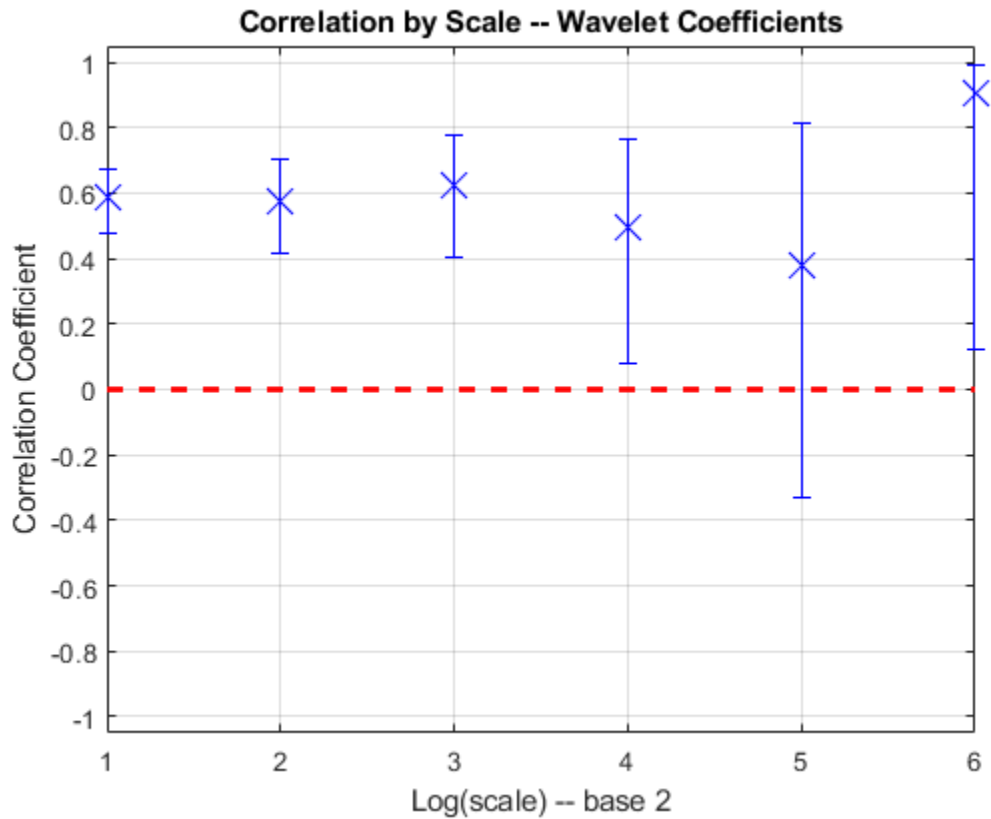
corrtable=7×6 table

	NJ	Lower	Rho	Upper	Pvalue	AdjustedPvalue
D1	12995	0.16942	0.19294	0.21624	1.5466e-55	2.8071e-54
D2	12989	0.21426	0.24683	0.27885	2.7037e-46	2.4536e-45
D3	12977	0.057885	0.10623	0.15407	1.789e-05	6.494e-05
D4	12953	0.048034	0.11645	0.18378	0.00088579	0.0026795
D5	12905	0.13281	0.2272	0.3175	3.7566e-06	1.7046e-05
D6	12809	-0.019835	0.1182	0.25181	0.093044	0.24125
S6	12809	0.26664	0.39003	0.50084	8.8066e-09	5.328e-08

Plot Correlation with Confidence Intervals

Plot the multiscale correlation of the DM-USD and JY-USD exchange returns down to level six. Use modwtcorr with no output arguments.

```
load DM_USD;  
load JY_USD;  
wdm = modwt(DM_USD, 'db2', 6);  
wjy = modwt(JY_USD, 'db2', 6);  
modwtcorr(wdm, wjy, 'db2')
```



Input Arguments

w1 — MODWT transform of signal 1
matrix

MODWT transform of signal 1, specified as a matrix. `w1` is the output of `modwt`. `w1` and `w2` must be the same size and both must have been obtained using the same analyzing wavelet.

Data Types: double

w2 — MODWT transform of signal 2

matrix

MODWT transform of signal 2, specified as a matrix. `w2` is the output of `modwt`. `w1` and `w2` must be the same size and both must have been obtained using the same analyzing wavelet.

wav — Wavelet

'sym4' (default) | character vector | string scalar | positive even scalar

Wavelet, specified as a character vector or string scalar indicating a valid wavelet name, or as a positive even scalar indicating the length of the wavelet and scaling filters. `wav` must be the same wavelet and length used to obtain the MODWTs of `w1` and `w2`. For a list of valid wavelets, see `modwt`. If unspecified or specified as an empty, `[]`, `wav` defaults to the symlet wavelet with four vanishing moments, 'sym4'.

conflevel — Confidence level

0.95 (default) | positive scalar less than 1

Confidence level, specified as a positive scalar less than 1. `conflevel` determines the coverage probability of the confidence intervals in `wcorr` and in the table, if you specify 'table' as an input. If unspecified, or if specified as empty, `[]`, `conflevel` defaults to 0.95.

Output Arguments

wcorr — Correlation coefficients by scale

vector

Correlation coefficients by scale, returned as a vector. `wcorr` is an M -by-1 vector of correlation coefficients, where M is the number of levels with nonboundary wavelet coefficients. `modwtcorr` returns correlation estimates only where there are nonboundary coefficients. This condition is satisfied when the transform level is not greater than $\text{floor}(\log_2(N/(L-1)+1))$, where N is the length of the original signal and L is the filter length. If the final level has enough nonboundary coefficients, `modwtcorr` returns

the scaling correlation in the final row of `wcorr`. By default, `modwtcorr` uses the `symlet` wavelet with four vanishing moments, `'sym4'` to determine the boundary coefficients.

wcorrci — Confidence intervals by scale

matrix

Confidence intervals by scale, returned as a matrix. The matrix is of size M -by-2, where M is the number of levels with nonboundary wavelet coefficients. The first column contains the lower confidence bound and the second column contains the upper confidence bound. The `conlevel` determines the coverage probability.

Confidence bounds are computed using Fisher's Z-transformation. The standard error of Fisher's Z statistic is the square root of $(N - 3)$. In this case, N is the equivalent number of coefficients in the critically sampled discrete wavelet transform (DWT), `floor(size(w1,2)/2^LEV)`, where `LEV` is the level of the wavelet transform. `modwtcorr` returns NaNs for the confidence bounds when $(N - 3)$ is less than or equal to zero.

pval — P-values for null hypothesis test

matrix

P -values for null hypothesis test, returned as a matrix. `pval` is an M -by-2 matrix.

- The first column of `pval` is the p -value computed using the standard t -statistic test for a correlation coefficient of zero.
- The second column of `pval` contains the adjusted p -value using the false discovery procedure of Benjamini & Yekutieli under arbitrary dependence assumptions.

The degrees of freedom, $(N - 2)$, for the t -statistic are determined by the equivalent number of coefficients N in the critically sampled DWT, `floor(size(w1,2)/2^LEV)`, where `LEV` is the level of the wavelet transform. `modwtcorr` returns NaNs when $(N - 2)$ is less than or equal to zero.

nj — Number of nonboundary coefficients

vector

Number of nonboundary coefficients by scale, returned as a vector.

wcorrtable — Correlation table

table

Correlation table, returned as a MATLAB table. The table contains six variables:

- **NJ** — Number of nonboundary coefficients by level.
- **Lower** — Lower confidence bound for the coverage probability specified by `conlevel`.
- **Rho** — Correlation coefficient.
- **Upper** — Upper confidence bound for the coverage probability specified by `conlevel`.
- **Pvalue** — P-value for hypothesis test. The null hypothesis is that the correlation coefficient is equal to zero.
- **AdjustedPvalue** — P-value adjusted for multiple comparisons. The p-values are adjusted using false discovery rate under dependency assumptions.

References

- [1] Percival, D. B., and Walden, A. T. *Wavelet Methods for Time Series Analysis*. Cambridge, U.K: Cambridge University Press, 2000.
- [2] Whitcher, B., P. Guttorp, and D. B. Percival. "Wavelet analysis of covariance with application to atmospheric time series." *Journal of Geophysical Research*, Vol. 105, pp. 14941-14962, 2000.
- [3] Benjamini, Y., and Yekutieli, D. "The Control of the False Discovery Rate in Multiple Testing Under Dependency." *Annals of Statistics*, Vol. 29, Number 4, pp. 1165-1188, 2001.

See Also

`imodwt` | `modwt` | `modwtmra` | `modwtvar` | `modwtxcorr`

Topics

"Wavelet Cross-Correlation for Lead-Lag Analysis"

"Wavelet Analysis of Financial Data"

Introduced in R2015b

modwtmra

Multiresolution analysis based on MODWT

Syntax

```
mra = modwtmra(w)
mra = modwtmra(w,wname)
mra = modwtmra(w,Lo,Hi)
mra = modwtmra( ____, 'reflection')
```

Description

`mra = modwtmra(w)` returns the multiresolution analysis (MRA) of the maximal overlap discrete wavelet transform (MODWT) matrix, `w`. The MODWT matrix, `w`, is the output of the `modwt` function.

`mra = modwtmra(w,wname)` constructs the MRA using the wavelet corresponding to `wname`. The `wname` wavelet must be the same wavelet used to obtain the MODWT.

`mra = modwtmra(w,Lo,Hi)` constructs the MRA using the scaling filter `Lo` and wavelet filter `Hi`. The `Lo` and `Hi` filters must be the same filters used to obtain the MODWT.

`mra = modwtmra(____, 'reflection')` uses the reflection boundary condition in the construction of the MRA using any of the arguments from previous syntaxes. If you specify `'reflection'`, `modwtmra` assumes that the length of the original signal is one half the number of columns in the input coefficient matrix.

Examples

Perfect Reconstruction with the MODWTMRA

Obtain the MODWTMRA of a simple time-series signal and demonstrate perfect reconstruction.

Create a time-series signal

```
t = 1:10;  
x = sin(2*pi*200*t);
```

Obtain the MODWT and the MODWTMRA and sum the MODWTMRA rows.

```
m = modwt(x);  
mra = modwtmra(m);  
xrec = sum(mra);
```

Use the maximum of the absolute values to show that the difference between the original signal and the reconstruction is extremely small. The largest absolute value is on the order of 10^{-25} , which demonstrates perfect reconstruction.

```
max(abs(x-xrec))  
ans = 5.5738e-25
```

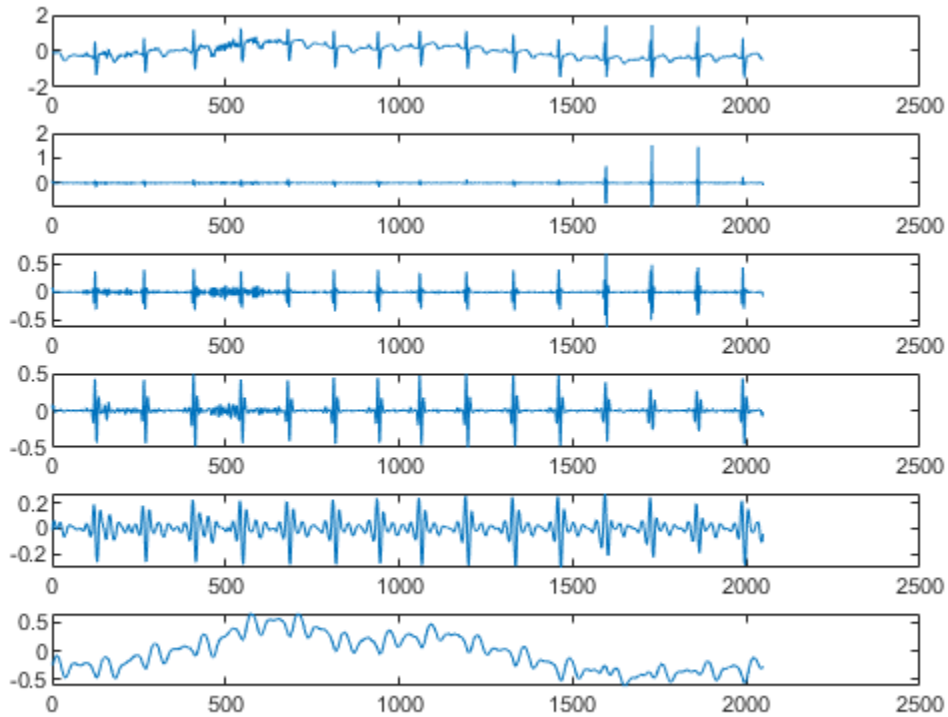
MRA Using Non-Default Wavelet

Construct an MRA of an ECG signal down to level four using the 'db2' wavelet.

```
load wecg;  
lev = 4;  
wtecg = modwt(wecg, 'db2', lev);  
mra = modwtmra(wtecg, 'db2');
```

Plot the ECG waveform and the MRA.

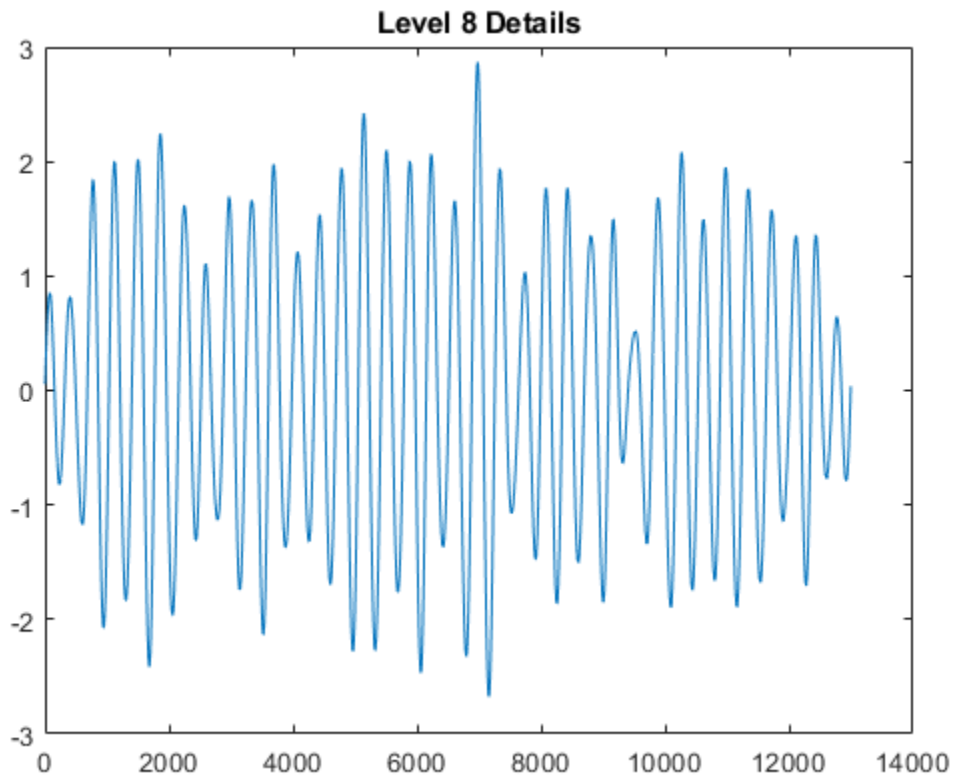
```
subplot(6,1,1)  
plot(wecg)  
for kk = 2:lev+2  
    subplot(6,1, kk)  
    plot(mra(kk-1, :))  
end
```



MRA Using the Default Wavelet

Construct a multiresolution analysis for the Southern Oscillation Index data. The sampling period is one day. Plot the level eight details corresponding to a scale of 2^8 days. The details at this scale capture oscillations on a scale of approximately one year.

```
load soi
wtsoi = modwt(soi);
mrasoi = modwtmra(wtsoi);
plot(mrasoi(8,:))
title('Level 8 Details')
```



MRA Using Minimum Bandwidth Scaling and Wavelet Filters

Obtain the MRA for the Deutsch Mark - U.S. Dollar exchange rate data using the minimum bandwidth scaling and wavelet filters with four coefficients.

```
load DM_USD;  
Lo = [0.4801755, 0.8372545, 0.2269312, -0.1301477];  
Hi = qmf(Lo);  
wdm = modwt(DM_USD,Lo,Hi);  
mra = modwtmra(wdm,Lo,Hi);
```

MRA Using Reflection Boundary

Obtain the MRA for an ECG signal using 'reflection' boundary handling.

```
load wecg;  
wtecg = modwt(wecg, 'reflection');  
mra = modwtmra(wtecg, 'reflection');
```

Show that the number of columns in the MRA is equal to the number of elements in the original signal.

```
isequal(size(mra,2), numel(wecg))
```

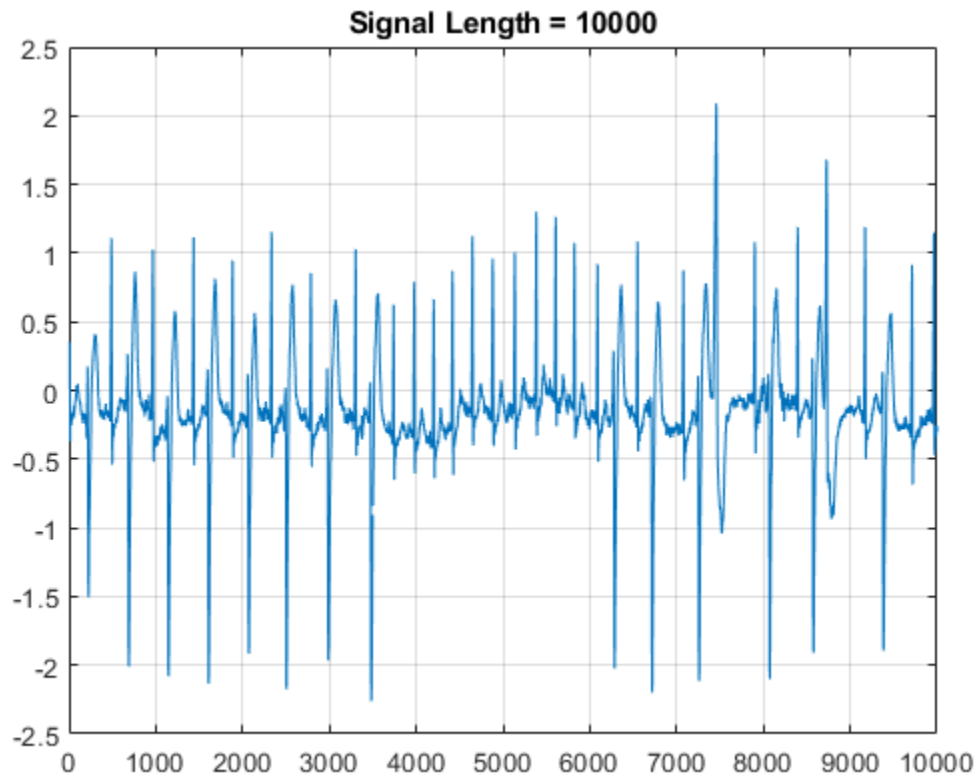
```
ans = logical  
     1
```

Comparing MODWT and MODWTMRA

This example demonstrates the differences between the functions MODWT and MODWTMRA. The MODWT partitions a signal's energy across detail coefficients and scaling coefficients. The MODWTMRA projects a signal onto wavelet subspaces and a scaling subspace.

Choose the 'sym6' wavelet. Load and plot an ECG waveform. The ECG data is taken from the MIT-BIH Arrhythmia Database.

```
load mit200  
wv = 'sym6';  
plot(ecgsig)  
grid on  
title(['Signal Length = ', num2str(length(ecgsig))])
```



Take the MODWT of the signal.

```
wtecg = modwt(ecgsig,wv);
```

The input data are samples of a function $f(x)$ evaluated at N -many time points. The function can be expressed as a linear combination of the scaling function $\phi(x)$ and wavelet

$\psi(x)$ at varying scales and translations: $f(x) = \sum_{k=0}^{N-1} c_k 2^{-J_0/2} \phi(2^{-J_0} x - k) + \sum_{j=1}^{J_0} f_j(x)$ where

$f_j(x) = \sum_{k=0}^{N-1} d_{j,k} 2^{-j/2} \psi(2^{-j} x - k)$ and J_0 is the number of levels of wavelet decomposition.

The first sum is the coarse scale approximation of the signal, and the $f_j(x)$ are the details

at successive scales. MODWT returns the N -many coefficients $\{c_k\}$ and the $(J_0 \times N)$ -many detail coefficients $\{d_{j,k}\}$ of the expansion. Each row in `wtecg` contains the coefficients at a different scale.

When taking the MODWT of a signal of length N , there are $\text{floor}(\log_2(N))$ -many levels of decomposition (by default). Detail coefficients are produced at each level. Scaling coefficients are returned only for the final level. In this example, since $N = 10000$, $J_0 = \text{floor}(\log_2(10000)) = 13$ and the number of rows in `wtecg` is $J_0 + 1 = 13 + 1 = 14$.

The MODWT partitions the energy across the various scales and scaling coefficients:

$\|X\|^2 = \sum_{j=1}^{J_0} \|W_j\|^2 + \|V_{J_0}\|^2$ where X is the input data, W_j are the detail coefficients at scale j , and V_{J_0} are the final-level scaling coefficients.

Compute the energy at each scale, and evaluate their sum.

```
energy_by_scales = sum(wtecg.^2,2);
Levels = {'D1'; 'D2'; 'D3'; 'D4'; 'D5'; 'D6'; 'D7'; 'D8'; 'D9'; 'D10'; 'D11'; 'D12'; 'D13'; 'A13'};
energy_table = table(Levels,energy_by_scales);
disp(energy_table)
```

Levels	energy_by_scales
'D1'	0.31592
'D2'	2.6504
'D3'	28.802
'D4'	159.37
'D5'	300.5
'D6'	431.33
'D7'	444.93
'D8'	182.37
'D9'	45.381
'D10'	11.578
'D11'	19.809
'D12'	4.5406
'D13'	3.308
'A13'	192.46

```
energy_total = varfun(@sum,energy_table(:,2))
energy_total=1x1 table
    sum_energy_by_scales
```

1827.3

Confirm the MODWT is energy-preserving by computing the energy of the signal and comparing it with the sum of the energies over all scales.

```
energy_ecg = sum(ecgsig.^2);
max(abs(energy_total.sum_energy_by_scales-energy_ecg))
```

```
ans = 4.0870e-09
```

Take the MODWTMRA of the signal.

```
mraecg = modwtmra(wtecg,wv);
```

MODWTMRA returns the projections of the function $f(x)$ onto the various wavelet subspaces and final scaling space. That is, MODWTMRA returns

$\sum_{k=0}^{N-1} c_k 2^{-J_0/2} \phi(2^{-J_0} x - k)$ and the J_0 -many $\{f_j(x)\}$ evaluated at N -many time points. Each

row in `mraecg` is a projection of $f(x)$ onto a different subspace. This means the original signal can be recovered by adding all the projections. This is *not* true in the case of the MODWT. Adding the coefficients in `wtecg` will *not* recover the original signal.

Choose a time point, add the projections of $f(x)$ evaluated at that time point and compare with the original signal.

```
time_point = 1000;
abs(sum(mraecg(:,time_point))-ecgsig(time_point))
```

```
ans = 3.0970e-13
```

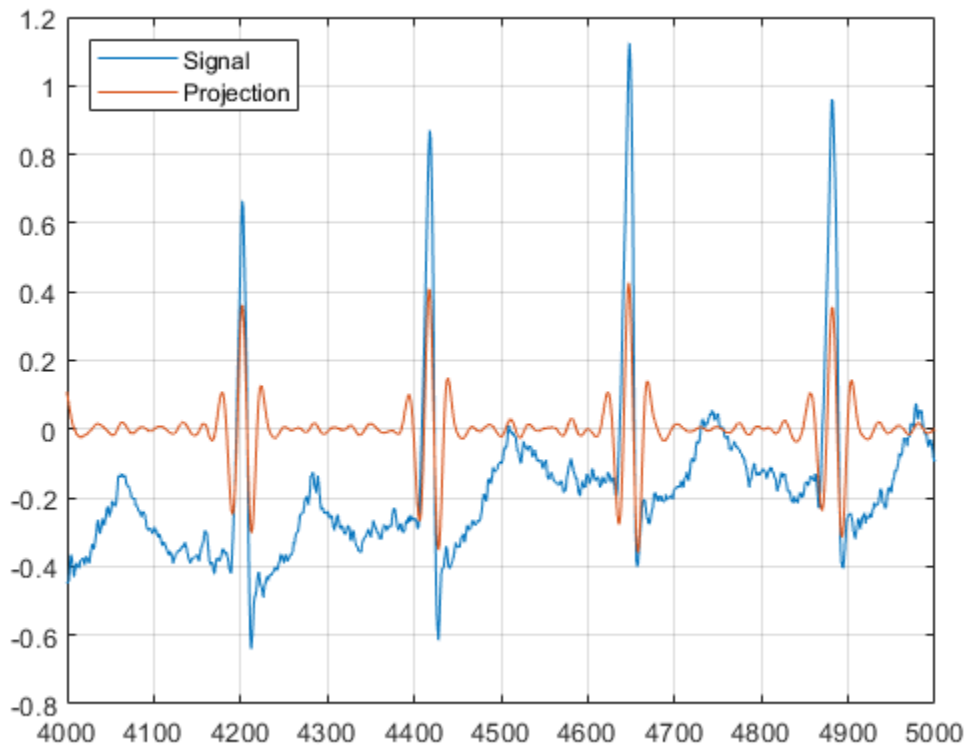
Confirm that, unlike MODWT, MODWTMRA is not an energy-preserving transform.

```
energy_ecg = sum(ecgsig.^2);
energy_mra_scales = sum(mraecg.^2,2);
energy_mra = sum(energy_mra_scales);
max(abs(energy_mra-energy_ecg))
```

```
ans = 534.7949
```

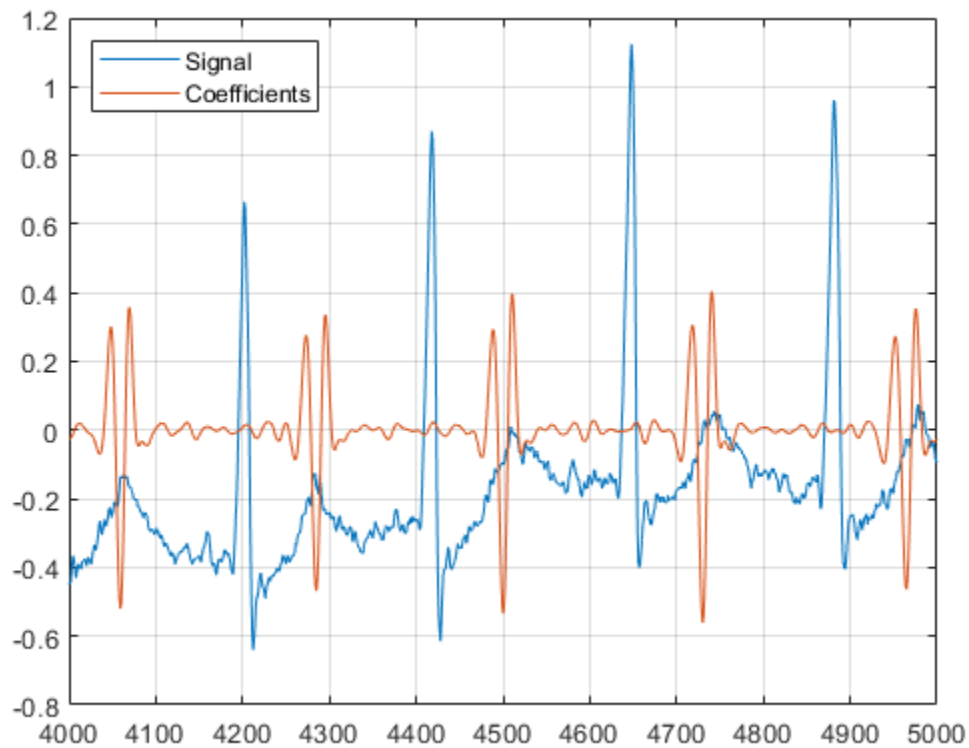
The MODWTMRA is a zero-phase filtering of the signal. Features will be time-aligned. Demonstrate this by plotting the original signal and one of its projections. To better illustrate the alignment, zoom in.

```
figure
plot(ecgsig)
hold on
plot(mraecg(4,:), '-')
grid on
xlim([4000 5000])
legend('Signal', 'Projection', 'Location', 'northwest')
```



Make a similar plot using the MODWT coefficients at the same scale. Note that features will not be time-aligned. The MODWT is *not* a zero-phase filtering of the input.


```
figure
plot(ecgsig)
hold on
plot(wtecg(4,:), '-')
grid on
xlim([4000 5000])
legend('Signal', 'Coefficients', 'Location', 'northwest')
```



References

Goldberger A. L., L. A. N. Amaral, L. Glass, J. M. Hausdorff, P. Ch. Ivanov, R. G. Mark, J. E. Mietus, G. B. Moody, C-K Peng, H. E. Stanley. "PhysioBank, PhysioToolkit, and PhysioNet: Components of a New Research Resource for Complex Physiologic Signals." *Circulation*

101. Vol.23, e215-e220, 2000. <http://circ.ahajournals.org/cgi/content/full/101/23/e215>

Moody, G. B. "Evaluating ECG Analyzers". <http://www.physionet.org/physiotools/wfdb/doc/wag-src/eval0.tex>

Moody G. B., R. G. Mark. "The impact of the MIT-BIH Arrhythmia Database." IEEE Eng in Med and Biol. Vol. 20, Number 3, 2001), pp. 45-50 .

Input Arguments

w — MODWT transform

matrix

Maximal overlap discrete wavelet transform, specified as a matrix. *w* is the output of `modwt`.

The input *w* is an $L+1$ -by- N matrix containing the MODWT of an N -point input signal down to level L . By default, `modwtmra` assumes that you obtained the MODWT using the symlet wavelet with four vanishing moments, 'sym4', and using periodic boundary handling.

Data Types: double

wname — Synthesis wavelet

'sym4' (default) | character vector | string scalar

Synthesis wavelet, specified as a character vector or string scalar. The synthesis wavelet must be the same wavelet used to obtain the MODWT with the `modwt` function.

Lo — Scaling filter

even-length real-valued vector

Scaling filter, specified as an even-length real-valued vector. You can specify *Lo* only if you do not specify *wname*. *Lo* must be the same scaling filter used to obtain the MODWT with the `modwt` function.

Hi — Wavelet filter

even-length real-valued vector

Wavelet filter, specified as an even-length real-valued vector. You can specify H_i only if you do not specify $wname$. H_i must be the same wavelet filter used to obtain the MODWT with the `modwt` function.

Output Arguments

mra — Multiresolution analysis

matrix

Multiresolution analysis, returned as a matrix. By default, the `mra` is the same size as the input transform matrix w . If you specify reflection boundary handling, then `mra` has one half the number of columns as the input matrix w .

The output `mra` is an $L+1$ -by- N matrix. The k^{th} row of `mra` contains the details for the k^{th} level. The $(L+1)^{\text{th}}$ row of `mra` contains the L^{th} level smooth.

References

- [1] Percival, D. B., and Walden, A. T. *Wavelet Methods for Time Series Analysis*. Cambridge, U.K: Cambridge University Press, 2000.
- [2] Whitcher, B., P. Guttorp, and D. B. Percival. "Wavelet analysis of covariance with application to atmospheric time series." *Journal of Geophysical Research*, Vol. 105, 2000, pp. 14941-14962.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

`imodwt` | `modwt`

Introduced in R2015b

modwtvar

Multiscale variance of maximal overlap discrete wavelet transform

Syntax

```
wvar = modwtvar(w)
wvar = modwtvar(w,wname)
[wvar,wvarci] = modwtvar( ___ )

[ ___ ] = modwtvar(w,wname, ___ ,conflevel)
[ ___ ] = modwtvar(w,wname, ___ ,Name,Value,)
[wvar,wvarci,nj] = modwtvar(w,wname, ___ )

wvartable = modwtvar(w,wname,'table')

modwtvar( ___ )
```

Description

`wvar = modwtvar(w)` returns unbiased estimates of the wavelet variance by scale for the maximal overlap discrete wavelet transform (MODWT). The default wavelet type is `sym4`.

`wvar = modwtvar(w,wname)` uses the wavelet `wname` to determine the number of boundary coefficients by level for unbiased estimates.

`[wvar,wvarci] = modwtvar(___)` returns the 95% confidence intervals for the variance estimates by scale.

`[___] = modwtvar(w,wname, ___ ,conflevel)` uses `conflevel` for the coverage probability of the confidence interval.

`[___] = modwtvar(w,wname, ___ ,Name,Value,)` returns wavelet variance with additional options specified by one or more `Name,Value` pair arguments.

`[wvar,wvarci,nj] = modwtvar(w,wname, ___)` returns the number of coefficients used to form the variance and confidence intervals by level.

`wvartable = modwtvar(w,wname,'table')`, where 'table' returns a MATLAB table, `wvartable`, containing the number of MODWT coefficients by level, the confidence boundaries, and the variance estimates. You can place 'table' anywhere after input `w`, except between the name and value of another Name, Value pair.

`modwtvar(___)` with no output arguments plots the wavelet variances by scale with lower and upper confidence bounds. The scaling variance is not included in the plot because the scaling variance can be much larger than the wavelet variances.

Examples

Wavelet Variance Using Default Wavelet

Obtain the MODWT of the Southern Oscillation Index data using the default symlets wavelet with 4 vanishing moments. Compute the unbiased estimates of the wavelet variance by scale.

```
load soi
wsoi = modwt(soi);
wvar = modwtvar(wsoi)
```

```
wvar = 10×1
```

```
0.3568
0.9026
1.1576
1.0952
0.9678
0.5478
0.6353
1.9570
0.8398
0.8247
```

Wavelet Variance Using Specified Wavelet

Obtain the MODWT of the Southern Oscillation Index data using the Daubechies wavelet with 2 vanishing moments ('db2'). Compute the unbiased estimates of the wavelet variance by scale.

```
load soi
wsoi = modwt(soi, 'db2');
wvar = modwtvar(wsoi, 'db2')
```

```
wvar = 12×1
```

```
    0.4296
    0.9204
    1.1370
    1.0847
    0.9255
    0.5932
    0.7630
    1.6672
    0.8048
    0.7555
    :
```

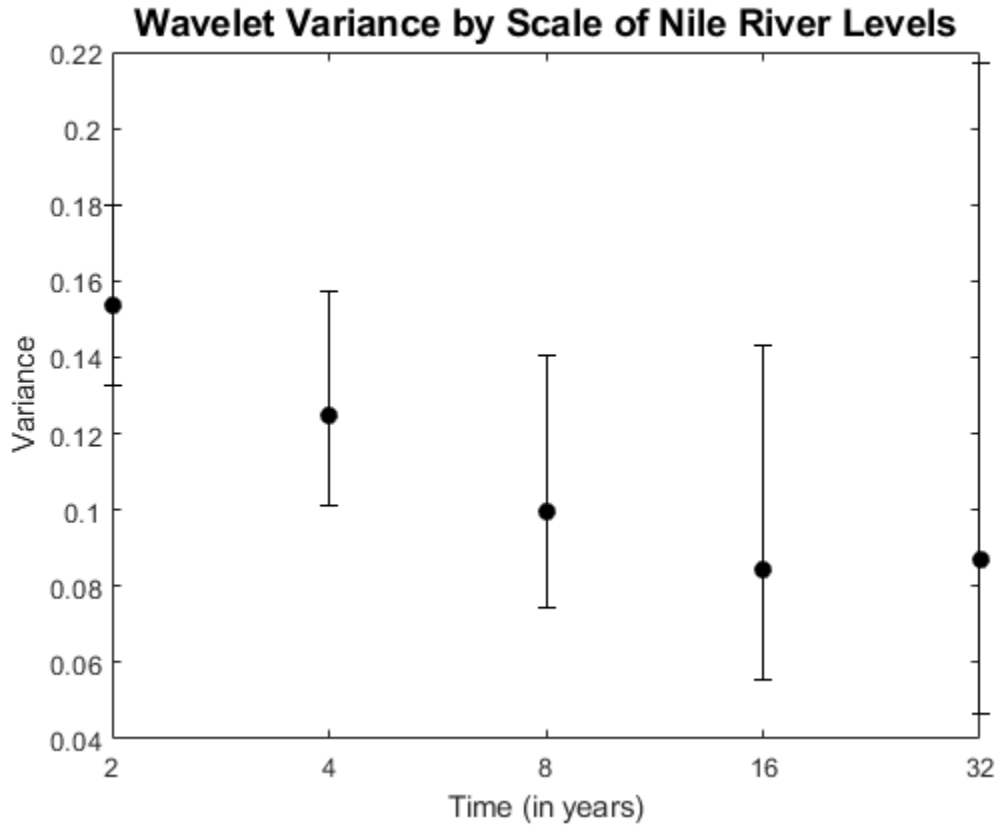
Variance Estimates and Confidence Intervals Using MODWTVAR

Obtain the MODWT of the Nile River minimum level data using the Fejer- Korovkin wavelet with eight coefficients down to level five. Use modwtvar to obtain and plot the variance estimates and 95% confidence intervals.

```
load nileriverminima;
wtnile = modwt(nileriverminima, 'fk8', 5);
[wnilevar, wvarci] = modwtvar(wtnile, 'fk8');

errlower = (wnilevar - wvarci(:, 1));
errupper = (wvarci(:, 2) - wnilevar);
errorbar(1:5, wnilevar(1:5), errlower(1:5), ...
         errupper(1:5), 'ko', 'markerfacecolor', 'k')
hold on
title('Wavelet Variance by Scale of Nile River Levels', 'fontsize', 14);
ylabel('Variance');
```

```
xlabel('Time (in years)');  
ax = gca;  
ax.XTick = [1:5];  
ax.XTickLabel = {'2', '4', '8', '16', '32'};  
hold off
```



Wavelet Confidence Intervals

Show how different confidence level values affect the width of the confidence intervals. An increased confidence level value increases the confidence interval width.

Obtain the MODWT of the Southern Oscillation Index data using the Fejer-Korovkin wavelet with eight coefficients.

```
load soi;
wsoi = modwt(soi, 'fk8');
```

Obtain the width of the .90, .95, and .99 confidence intervals for each level.

```
[~,wvarci90] = modwtvar(wsoi, 'fk8', 0.90);
w90 = wvarci90(:,2)-wvarci90(:,1);
 [~,,wvarci95] = modwtvar(wsoi, 'fk8', 0.95);
w95 = wvarci95(:,2)-wvarci95(:,1);
[~,wvarci99] = modwtvar(wsoi, 'fk8', 0.99);
w99 = wvarci99(:,2)-wvarci99(:,1);
```

Compare the three columns. The first column shows the .90 confidence level values, the second the .95 values, and the third the .99 values. Each row is the width of the interval at each wavelet scale. You can see that the width of the confidence interval increases with larger confidence level values.

```
[w90,w95,w99]
```

```
ans = 10x3
```

```
    0.0195    0.0233    0.0306
    0.0739    0.0880    0.1158
    0.1347    0.1606    0.2113
    0.1798    0.2145    0.2826
    0.2304    0.2751    0.3634
    0.1825    0.2184    0.2900
    0.2858    0.3435    0.4613
    1.5445    1.8757    2.5837
    1.0625    1.3262    1.9551
    2.8460    3.9883    7.8724
```

Compare Chi2Eta2 and Gaussian Confidence Intervals

Specify non-default confidence methods using name-value pairs to compare the width of their confidence levels. Note that for Gaussian confidence level intervals, it is possible to obtain negative lower confidence bounds.

Obtain the MODWT of the Southern Oscillation Index data using the Fejer-Korovkin wavelet with eight coefficients.

```
load soi;
wsoi = modwt(soi, 'fk8');
```

Use the Chi2Eta and Gaussian confidence methods to obtain the variances and confidence interval bounds for each method.

```
[wvar_c, wvarci_c] = modwtvar(wsoi, 'fk8', [], 'ConfidenceMethod', 'chi2eta1');
[wvar_g, wvarci_g] = modwtvar(wsoi, 'fk8', [], 'ConfidenceMethod', 'gaussian');
```

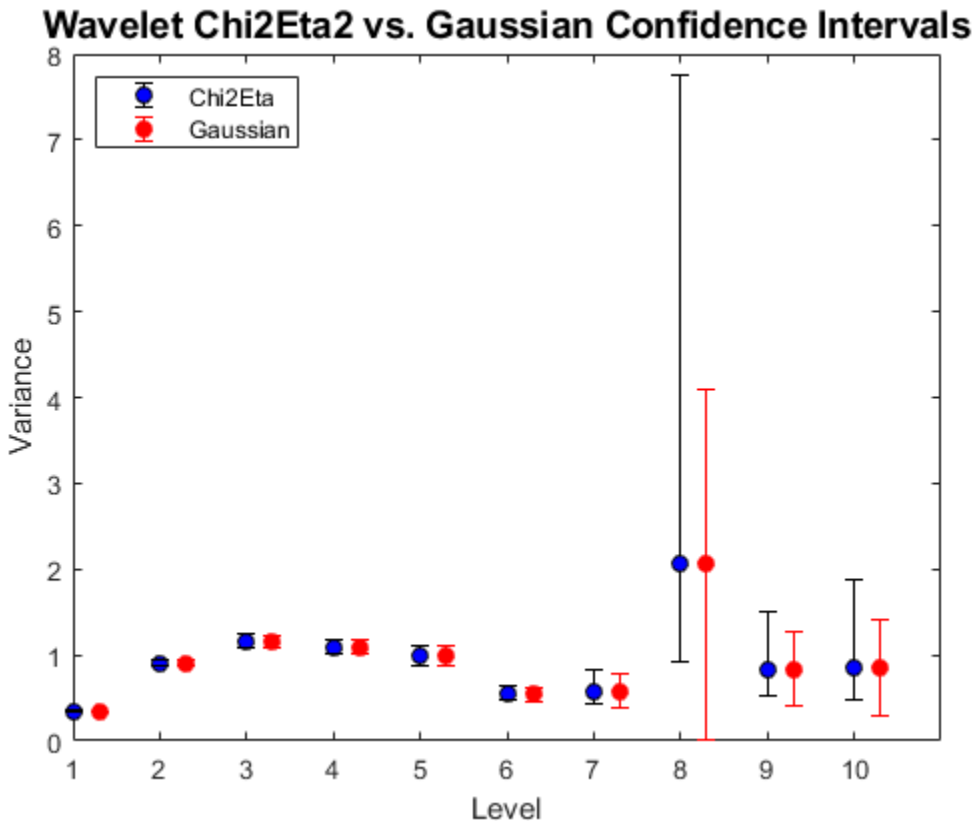
Compute the upper and lower errors for each confidence interval and plot the results. Note that the Gaussian intervals are slightly shifted to enable better visualization.

```
errlower_c = wvar_c - wvarci_c(:, 1);
errupper_c = wvarci_c(:, 2) - wvar_c;

errlower_g = wvar_g(:, 1) - wvarci_g(:, 1);
errupper_g = wvarci_g(:, 2) - wvar_g;

errorbar(1:10, wvar_c(1:10), errlower_c(1:10), ...
         errupper_c(1:10), 'ko', 'markerfacecolor', 'b')
hold on;
xoffset = (1.3:10.3);
errorbar(xoffset, wvar_g(1:10), errlower_g(1:10), ...
         errupper_g(1:10), 'ro', 'markerfacecolor', 'r')

title('Wavelet Chi2Eta2 vs. Gaussian Confidence Intervals', 'fontsize', 14);
ylabel('Variance');
xlabel('Level')
ax = gca;
ax.XTick = [1:10];
legend('Chi2Eta', 'Gaussian', 'Location', 'northwest');
hold off
```



Compare Number of Coefficients for Unbiased and Biased Variance Estimates

Compare the number of coefficients for unbiased and biased wavelet variance estimates. For the unbiased (default) estimates, the number of nonboundary coefficients decreases by scale. For biased estimates, the number of coefficients matches the number of input rows and is constant for every scale.

Obtain the MODWT of the Southern Oscillation Index data using the Fejer-Korovkin wavelet with eight coefficients. Compute the unbiased and biased estimates of the wavelet variance down to level ten. The number of coefficients used in the unbiased estimates decrease by scale.

```
load soi
wsoi = modwt(soi, 'fk8');
[wvar_unb, wvarci_unb, nj_unb] = modwtvar(wsoi, 'fk8');
[wvar_b, wvarci_b, nj_b] = modwtvar(wsoi, 'fk8', [], 'EstimatorType', 'biased');
[nj_unb(1:10), nj_b(1:10)]
```

```
ans = 10x2
```

```
    12991    12998
    12977    12998
    12949    12998
    12893    12998
    12781    12998
    12557    12998
    12109    12998
    11213    12998
     9421    12998
     5837    12998
```

Table of Wavelet Variance Estimates Using Gaussian Confidence Intervals

Compute the MODWT of the Southern Oscillation Index data using the Fejer- Korovkin wavelet with eight coefficients. Compute a variance table for the data.

```
load soi;
wsoi = modwt(soi, 'fk8');
[wvartable] = modwtvar(wsoi, 'fk8', 0.90, 'ConfidenceMethod', 'Gaussian', ...
    'table')
```

```
wvartable=10x4 table
```

	NJ	Lower	Variance	Upper
	-----	-----	-----	-----
D1	12991	0.3291	0.33848	0.34786
D2	12977	0.87172	0.9034	0.93508
D3	12949	1.1041	1.1628	1.2216
D4	12893	1.0204	1.0933	1.1662
D5	12781	0.8833	0.98255	1.0818
D6	12557	0.47178	0.54152	0.61125
D7	12109	0.41916	0.57934	0.73951
D8	11213	0.33639	2.055	3.7736

D9	9421	0.4752	0.83369	1.1922
D10	5837	0.37485	0.84386	1.3129

The resulting table contains the number of nonboundary coefficients, the lower and upper confidence level bounds, and the variance estimate for each level.

Input Arguments

w — MODWT transform matrix

matrix

MODWT transform, specified as a matrix. `w` is the output of `modwt`.

Data Types: double

wname — Wavelet

'sym4' (default) | character vector | string scalar | positive even scalar

Wavelet, specified as a character vector or string scalar corresponding to a valid wavelet, or as a positive even scalar indicating the length of the wavelet and scaling filters. The wavelet filter length must match the length used in the MODWT of the input.

If you use `Name, Value` pairs arguments or the 'table' syntax and you did not specify a `wname`, you must use `[]` as the second argument.

conflevel — Confidence level

0.95 (default) | real scalar greater than 0 and less than 1

Confidence level, specified as a real scalar value greater than 0 and less than 1. The confidence level determines the coverage probability of the confidence intervals. The confidence levels are also shown in `wvartable`, if you specify 'table' as an input.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name, Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: 'EstimatorType', 'biased' specifies a biased estimator.

EstimatorType — Estimator

'unbiased' (default) | 'biased'

Type of estimator used for variance estimates and confidence bounds, specified as the comma-separated pair consisting of 'EstimatorType' and one of these values.

- 'unbiased' — Unbiased estimator, which identifies and removes boundary coefficients prior to computing the variance estimates and confidence bounds. Unbiased estimates are used more frequently for wavelet variance computations.
- 'biased' — Biased estimator, which uses all coefficients to compute the variance estimates and confidence bounds.

ConfidenceMethod — Confidence method

'chi2eta3' (default) | 'chi2eta1' | 'gaussian'

Confidence method used to compute the confidence intervals, specified as the comma-separated pair consisting of 'ConfidenceMethod' and one of these values:

'chi2eta3'	Chi-square probability density method three, which determines the degrees of freedom.[1].
'chi2eta1'	Chi-square probability density method one, which determines the degrees of freedom [1].
'gaussian'	Gaussian method [1] . This method can result in negative lower bounds.

See “Algorithm” on page 1-891 for information on each of these confidence methods.

Boundary — Boundary condition

'periodic' (default) | 'reflection'

Boundary condition used to compute the variance estimates and confidence bounds, specified as the comma-separated pair consisting of 'Boundary' and one of these values:

'periodic'

Periodic boundary handling, which does not change the original signal before computing the MODWT. If `modwt` uses periodic boundary handling, you must specify 'Boundary', 'periodic' for `modwtvar` to obtain a correct estimate.

'reflection'

Reflection boundary handling. If the MODWT uses reflection boundary handling, you must also specify 'Boundary', 'reflection' for `modwtvar` to obtain a correct unbiased estimate. The MODWT, with reflection boundary handling, extends the original signal symmetrically at the right boundary to twice the signal length. The MODWTVAR algorithm has to know about this extended signal to calculate the correct unbiased estimate.

For biased estimators, all the coefficients are used to form the variance estimates and confidence intervals regardless of the boundary handling.

Output Arguments

wvar — Wavelet variance estimates

matrix

Wavelet variance estimates, returned as vector. The number of elements in `wvar` depends on the number of scales in the input matrix and, for unbiased estimates, on the wavelet length. For the unbiased case, `modwtvar` returns estimates only where nonboundary coefficients exist. This condition is satisfied when the transform level is not greater than $\text{floor}(\log_2(N/(L-1)+1))$, where N is the input signal length and L is the length of the wavelet filter. The number of biased estimates equals the input signal length. If the final level has sufficient nonboundary coefficients, `modwtvar` returns the scaling variance in the final element of `wvar`.

wvarci — Confidence intervals for each variance estimate

matrix

Confidence bounds, expressed as upper and lower confidence bounds, for the variance estimates in `wvar`, returned as a matrix. The default is 95% confidence bounds, but you can use a different value using the `conflevel` input argument. The confidence bounds matrix is M -by-2, where M is the number of levels. For unbiased estimates, the number of levels is limited by the number of nonboundary coefficients. For biased estimates, all levels are used. The first column of the confidence interval matrix contains the lower confidence bound and the second column contains the upper confidence bound. By default, `modwtvar` calculates the confidence intervals using the chi-square probability density, with the equivalent degrees of freedom estimated using the 'Chi2Eta3' confidence method.

nj — Number of coefficients by level

vector

Number of nonboundary coefficients by scale, returned as a vector. For unbiased estimates, `nj` is the number of nonboundary coefficients and decreases by level. For biased estimates, `nj` is a vector of constants equal to the number of columns in the input matrix.

wvartable — Variance table

table

Variance table, returned as a MATLAB table. The four variables in the table are:

- **NJ** — Number of MODWT coefficients by level. For biased estimates, `NJ` is the number of coefficients in the MODWT. For unbiased estimates, `NJ` is the number of nonboundary coefficients.
- **Lower** — Lower confidence bound for the variance estimate.
- **Variance** — Variance estimate by level.
- **Upper** — Upper confidence bound for the variance estimate.

The row names of `wvartable` indicate the type and level of each estimate. For example, `D1` indicates that the row corresponds to a wavelet or detail estimate at level 1. `S6` indicates that the row corresponds to the scaling estimate at level 6. The scaling variance is computed for the final level of the MODWT. For unbiased estimates, `modwtvar` computes the scaling variance only when nonboundary scaling coefficients exist.

Algorithms

The following expressions define the variance and confidence methods used in the MODWTVAR. The variables are

- N_j — Number of coefficients at level j
- v^2 — Variance
- j — Level
- $W_{j,t}$ — Wavelet coefficients

The variance estimate is

$$\widehat{v}_j^2 = \frac{1}{N_j} \sum_{t=0}^{N_j-1} W_{j,t}^2$$

The degrees of freedom for the Chi2Eta1 (chi2eta1) method are defined as

$$\eta_1 = \frac{N_j \widehat{v}_j^4}{\widehat{A}_j}$$

where

$$\widehat{A}_j = \frac{1}{2} \int_{-1/2}^{1/2} \left[\widehat{S}_j^{(p)}(f) \right]^2 df$$

In this equation, $\widehat{S}_j^{(p)}$ is the spectral density function estimate of the wavelet coefficients at level j .

The chi-square statistic is

$$\frac{\eta_1 N_j \widehat{v}_j^2}{v_j^2} \sim X_{\eta_1}^2$$

The degrees of freedom for the Chi2Eta3 (chi2eta3) method are defined as

$$\eta_3 = \max\left(\frac{N_j}{2^j}, 1\right)$$

The chi-square statistic is

$$\frac{\eta_3 N_j \widehat{v}_j^2}{v_j^2} \sim X_{\eta_3}^2$$

For the Gaussian method, the statistic

$$\frac{N_j^{1/2} (\widehat{v}_j^2 - v_j^2)}{(2\widehat{A}_j)^{1/2}}$$

is distributed as $N(0, 1)$. The variable \widehat{A}_j is as described for `chi2eta1`.

References

- [1] Percival, D. B., and A. T. Walden. *Wavelet Methods for Time Series Analysis*. Cambridge, UK: Cambridge University Press, 2000.
- [2] Percival, D. B., D. Mondal, "A Wavelet Variance Primer." *Handbook of Statistics, Volume. 300, Time Series Analysis: Methods and Applications*, (T. S. Rao, S. S. Rao, and C. R. Rao, eds.). Oxford, UK: Elsevier, 2012, pp. 623-658.
- [3] Cornish, C. R., C. S. Bretherton, and D. B. Percival. "Maximal Overlap Wavelet Statistical Analysis with Application to Atmospheric Turbulence." *Boundary-Layer Meteorology*. Vol. 119, Number 2, 2005, pp. 339-374.

See Also

`imodwt` | `modwt` | `modwtcorr` | `modwtmra` | `modwtxcorr`

Topics

"Wavelet Analysis of Financial Data"

Introduced in R2015b

modwtcorr

Wavelet cross-correlation sequence estimates using the maximal overlap discrete wavelet transform (MODWT)

Syntax

```
xcseq = modwtcorr(w1,w2)
xcseq = modwtcorr(w1,w2,wav)

[xcseq,xcseqci] = modwtcorr( ___ )
[xcseq,xcseqci] = modwtcorr(w1,w2,wav,confllevel)
[xcseq,xcseqci,lags] = modwtcorr( ___ )

[ ___ ] = modwtcorr( ___ , 'reflection')
```

Description

`xcseq = modwtcorr(w1,w2)` returns the wavelet cross-correlation sequence estimates for the maximal overlap discrete wavelet transform (MODWT) transforms specified in `w1` and `w2`. `xcseq` is a cell array of vectors where the elements in each cell correspond to cross-correlation sequence estimates. If there are enough nonboundary coefficients at the final level, `modwtcorr` returns the scaling cross-correlation sequence estimate in the final cell of `xcseq`.

`xcseq = modwtcorr(w1,w2,wav)` uses the wavelet `wav` to determine the number of boundary coefficients by level.

`[xcseq,xcseqci] = modwtcorr(___)` returns in `xcseqci` the 95% confidence intervals for the cross-correlation sequence estimates in `xcseq`, using any arguments from the previous syntaxes.

`[xcseq,xcseqci] = modwtcorr(w1,w2,wav,confllevel)` uses `confllevel` for the coverage probability of the confidence interval. `confllevel` is a real scalar strictly greater than 0 and less than 1. If `confllevel` is unspecified or specified as empty, the coverage probability defaults to 0.95.

`[xcseq,xcseqci,lags] = modwtxcorr(___)` returns the lags for the wavelet cross-correlation sequence estimates in a cell array of column vectors.

`[___] = modwtxcorr(___ , 'reflection')` reduces the number of wavelet and scaling coefficients at each scale by half before computing the cross-correlation sequences. Specifying the 'reflection' option in `modwtxcorr` is equivalent to first obtaining the MODWT of `w1 w2` with 'periodic' boundary handling and then computing the wavelet cross-correlation sequence estimates. Use this option only when you obtain the MODWT of `w1` and `w2` using the 'reflection' boundary condition. You must enter the entire character vector 'reflection'. If you added a wavelet named 'reflection' using the wavelet manager, you must rename that wavelet prior to using this option. 'reflection' may be placed in any position in the input argument list after the input transforms `w1 w2`.

Examples

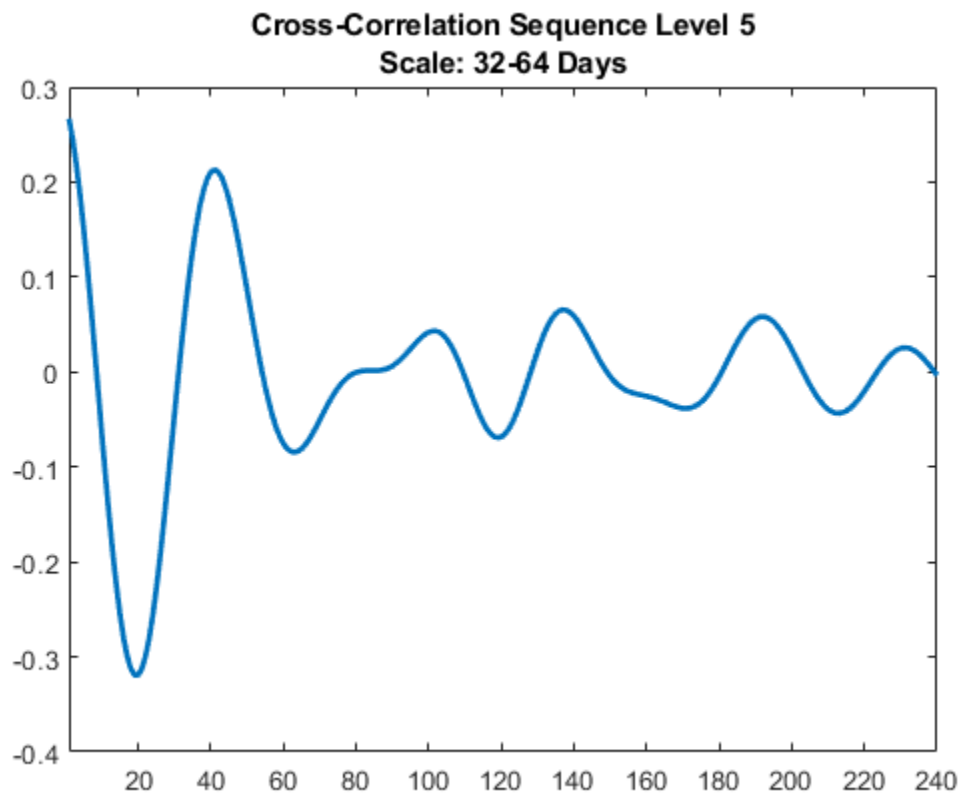
Cross-Correlation Sequence

Obtain the MODWT of the Southern Oscillation Index and Truk Islands pressure data. The sampling period is one day.

```
load soi
load truk
wsoi = modwt(soi);
wtruk = modwt(truk);
```

Compute the wavelet cross-correlation sequences. Examine the level-five cross-correlation sequence corresponding to a scale of 32-64 days. Determine the index corresponding to a lag of zero and plot out to 240 lags.

```
xcseq = modwtxcorr(wsoi,wtruk);
zerolag = floor(numel(xcseq{5})/2)+1;
plot(xcseq{5}(zerolag:zerolag+240), 'linewidth', 2)
set(gca, 'xlim', [1 240]);
title({'Cross-Correlation Sequence Level 5'; 'Scale: 32-64 Days'});
hold off
```



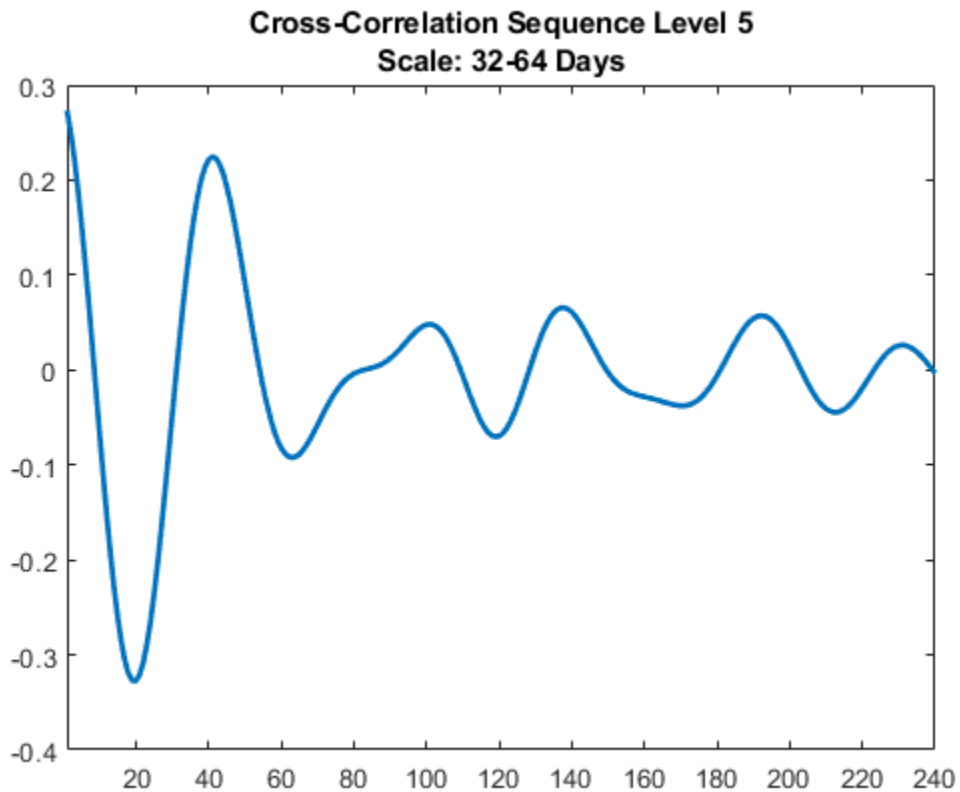
Cross-Correlation Sequence with Fejér-Korovkin Wavelet

Obtain the MODWT of the Southern Oscillation Index and Truk Islands pressure data using the Fejér-Korovkin wavelet filter with 8 coefficients. The sampling period of the data is one day.

```
load soi
load truk
wsoi = modwt(soi, 'fk8');
wtruk = modwt(truk, 'fk8');
```

Compute the wavelet cross-correlation sequences. Examine the level-five cross-correlation sequence corresponding to a scale of 32-64 days. Determine the index corresponding to a lag of zero and plot out to 240 lags.

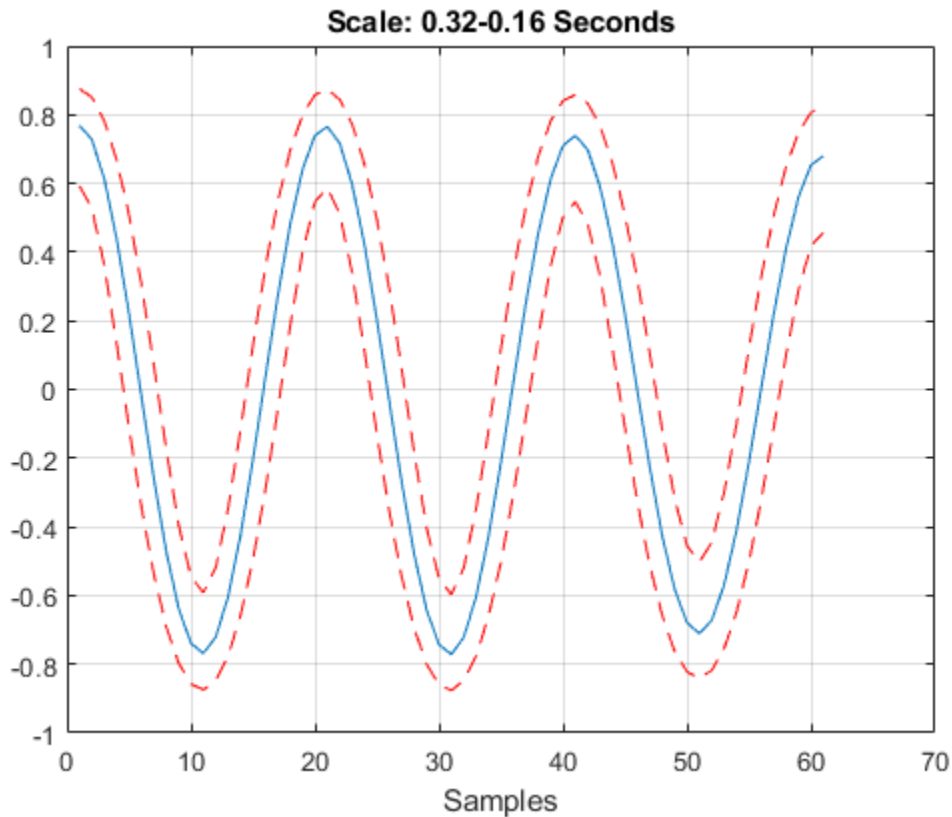
```
xcseq = modwtcorr(wsoi,wtruk,'fk8');  
zerolag = floor(numel(xcseq{5})/2)+1;  
plot(xcseq{5}(zerolag:zerolag+240),'linewidth',2)  
set(gca,'xlim',[1 240]);  
title({'Cross-Correlation Sequence Level 5'; 'Scale: 32-64 Days'});  
hold off
```



Cross-Correlation Confidence Intervals by Scale

Plot the wavelet cross-correlation with 95% confidence intervals at scale 4 for two 5-Hz sine wave signals with additive noise.

```
dt = 0.01;
t = 0:dt:6;
x = cos(2*pi*5*t)+1.5*randn(size(t));
y = cos(2*pi*5*t-pi)+2*randn(size(t));
wx = modwt(x, 'fk14', 5);
wy = modwt(y, 'fk14', 5);
modwtcorr(wx, wy, 'fk14')
j = 4;
[xcseq, xcseqci] = modwtcorr(wx, wy, 'fk14');
zerolag = floor(numel(xcseq{j})/2)+1;
lagidx = zerolag-30:zerolag+30;
plot(xcseq{j}(lagidx));
hold on;
grid
plot(xcseqci{j}(lagidx,:), 'r--');
xlabel('Samples');
title('Scale: 0.32-0.16 Seconds');
```



Cross-Correlation .90 and .95 Confidence Intervals Comparison

Compare the .90 and .95 (default) confidence intervals for the wavelet cross-correlation at level four.

Obtain the MODWT for two noisy sine waves using the Fejér-Korovkin with 14 coefficients, and specify the level to use.

```
dt = 0.01;  
t = 0:dt:6;  
x = cos(2*pi*5*t)+1.5*randn(size(t));  
y = cos(2*pi*5*t-pi)+2*randn(size(t));
```



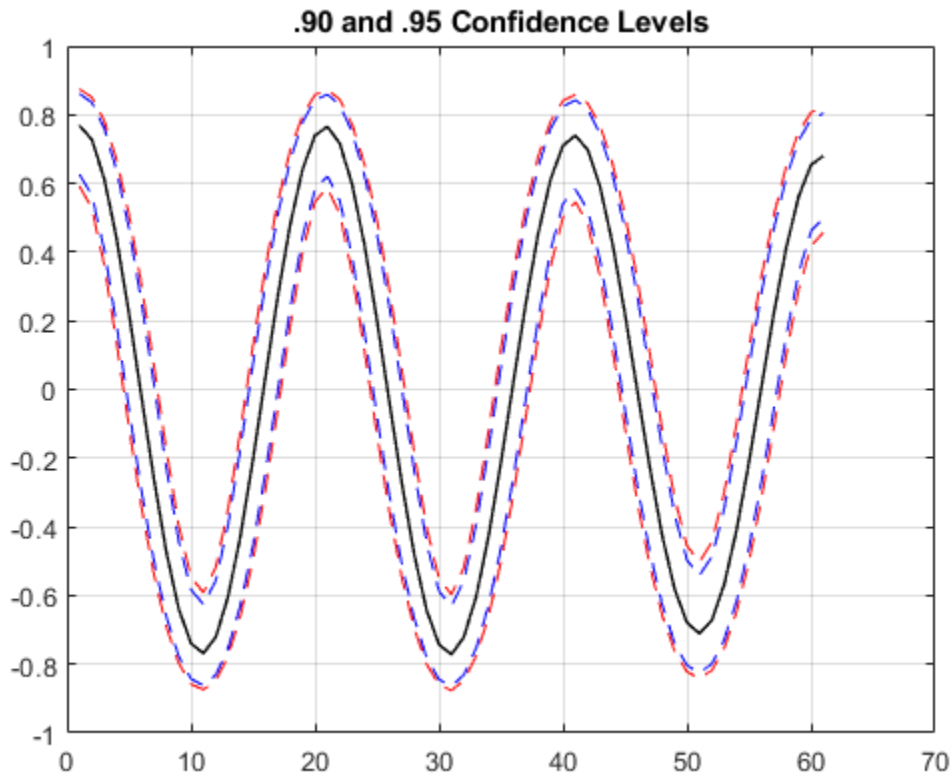
```
wx = modwt(x, 'fk14', 4);
wy = modwt(y, 'fk14', 4);
lev = 4;

[xcseq,xcseqci] = modwtcorr(wx,wy, 'fk14');
[xcseq90,xcseqci90] = modwtcorr(wx,wy, 'fk14',0.90);

zerolag = floor(numel(xcseq{lev})/2)+1;
zerolag90 = floor(numel(xcseq90{lev})/2)+1;

lagidx = zerolag-30:zerolag+30;
lagidx90 = zerolag90-30:zerolag90+30;

plot(xcseqci{lev}(lagidx,:), '--r');
hold on
plot(xcseqci90{lev}(lagidx90,:), '--b');
plot(xcseq{lev}(lagidx), '-k', 'LineWidth', 1);
grid
title('.90 and .95 Confidence Levels')
```



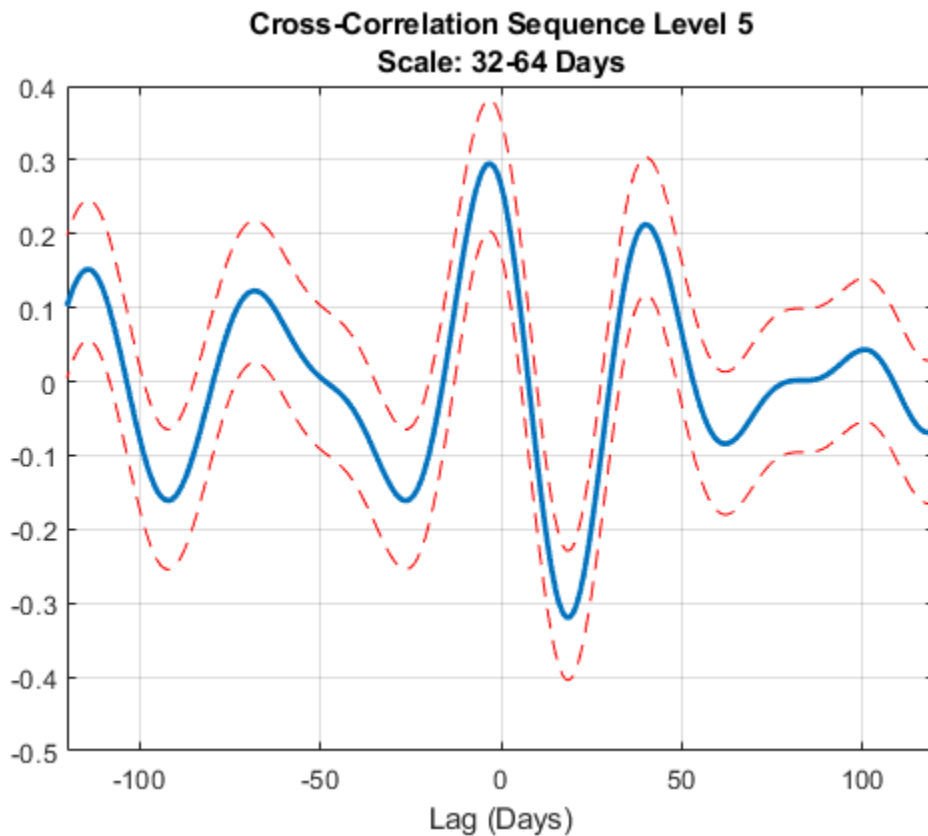
Notice that the .95 confidence interval width (in red) is larger than the .90 confidence interval width (in blue).

Plot Cross-Correlation Sequences by Lag

Plot the cross-correlation sequence estimate for the Southern Oscillation Index and Truk Island pressure data. Estimate 95% confidence intervals for scale of 2^5 days.

```
load soi
load truk
wsoi = modwt(soi);
```

```
wtruk = modwt(truk);  
[xcseq,xcseqci,lags] = modwtcorr(wsoi,wtruk);  
plot(lags{5},xcseq{5},'linewidth',2)  
hold on  
plot(lags{5},xcseqci{5},'r--')  
set(gca,'xlim',[-120 120]);  
xlabel('Lag (Days)');  
grid  
title({'Cross-Correlation Sequence Level 5'; 'Scale: 32-64 Days'});  
hold off
```



Cross-Correlation with Reflection Boundary

Obtain the MODWT of 36 years of Southern Oscillation Index and Truk Islands pressure data with both periodic and reflection boundary conditions. The `modwt` function with the `'reflection'` option extends the input signal symmetrically at the right boundary. The input signal is then twice its original length. MODWTXCORR with the reflection boundary handling reduces the number of wavelet and scaling coefficients at each half before computing the cross-correlation sequences. The size of the cross-correlation sequences is the same as acquiring the MODWT with the default periodic boundary condition.

```
load soi
load truk
```

Obtain the MODWT with the default periodic boundary condition.

```
wsoi_default = modwt(soi);
wtruk_default = modwt(truk);
```

Obtain the MODWT with the reflection boundary condition.

```
wsoi_reflect = modwt(soi, 'reflection');
wtruk_reflect = modwt(truk, 'reflection');
```

Obtain the cross-correlation sequences.

```
xcseq_default = modwtxcorr(wsoi_default, wtruk_default);
xcseq_reflect = modwtxcorr(wsoi_reflect, wtruk_reflect, 'reflection');
```

Compare the number of elements in the cell array output for both boundary conditions.

```
cellfun(@numel, xcseq_reflect)
```

```
ans = 10×1
    25981
    25953
    25897
    25785
    25561
    25113
    24217
    22425
    18841
    11673
```

```
cellfun(@numel,xcseq_default)
```

```
ans = 10×1
```

```

25981
25953
25897
25785
25561
25113
24217
22425
18841
11673

```

Input Arguments

w1 — MODWT transform of signal 1

matrix

MODWT transform of signal 1, specified as a matrix of doubles. The input w1 must be the same size as w2 and must have been obtained with the same wavelet. By default, `modwtcorr` assumes that you obtained the MODWT using the symlet wavelet with four vanishing moments, 'sym4').

w2 — MODWT transform of signal 2

matrix

MODWT transform of signal 2, specified as a matrix of doubles. The input w2 must be the same size as w1 and must have been obtained with the same wavelet. By default, `modwtcorr` assumes that you obtained the MODWT using the symlet wavelet with four vanishing moments ('sym4').

wav — Wavelet

'sym4' (default) | character vector | string scalar | positive even integer

Wavelet, specified as a character vector or string scalar, indicating a valid wavelet, or as a positive even integer indicating the length of the wavelet and scaling filters. If `wav` is unspecified or specified as an empty, [], `wav` defaults to 'sym4'.

conflevel — Confidence level

0.95 (default) | positive scalar less than 1

Confidence level, specified as a positive scalar less than 1. `conflevel` determines the coverage probability of the confidence intervals in `xcseqci`. If unspecified, or specified as empty, [], `conflevel` defaults to 0.95.

Output Arguments

xcseq — Cross-correlation sequences by scale

cell array of vectors

Cross-correlation sequences by scale, returned as a cell array of vectors. The vectors are of size $2NJ$ -by-1, where NJ is the number of nonboundary coefficients by level (scale). This level is the minimum of `size(w1,1)` and `floor(log2(N/(L-1)+1))` where N is the length of the data and L is the filter length. If there are enough nonboundary coefficients at the final level, `modwtcorr` returns the scaling cross-correlation sequence estimate in the final cell of `xcseq`.

xcseqci — Confidence intervals by scale

cell array of matrices

Confidence intervals by scale, returned as a cell array of matrices. The size of each matrix is $(2NJ-1)$ -by-2, where NJ is the number of nonboundary coefficients by level (scale).

- For the .95 default value, the first column of the i^{th} element of `xcseqci` contains the lower 95% confidence bound for the cross-correlation coefficient at each lag.
- For the .95 default value, the second column contains the upper 95% confidence bound.

Confidence bounds are computed using Fisher's Z-transformation. The standard error of Fisher's Z statistic is the square root of $N-3$. In this case, N is the equivalent number of coefficients in the critically sampled discrete wavelet transform (DWT), `floor(size(w1,2)/2^LEV)`, where LEV is the level of the wavelet transform. `modwtcorr` returns NaNs for the confidence bounds when N^{-3} is less than or equal to zero.

lags — Lags for the cross-correlation sequences

cell array of vectors

Lags for the cross-correlation sequences, returned as a cell array of vectors. `lags` is a cell array of column vectors the same length as `xcseq`. The elements in each cell of `xcseq` correspond to the cross-correlation sequence estimates at lags from $-(NJ-1)$ to $(NJ-1)$, where NJ is the number of nonboundary coefficients at level J . The 0th lag element is located at the index $\text{floor}((2*NJ-1)/2)+1$.

References

- [1] Percival, D. B., and Walden, A. T. *Wavelet Methods for Time Series Analysis*. Cambridge, U.K: Cambridge University Press, 2000.
- [2] Whitcher, B., P. Guttorp, and D. B. Percival. "Wavelet analysis of covariance with application to atmospheric time series." *Journal of Geophysical Research*, Vol. 105, 2000, pp. 14941-14962.

See Also

`imodwt` | `modwt` | `modwtcorr` | `modwtmra` | `modwtvar`

Topics

"Wavelet Analysis of Financial Data"

Introduced in R2015b

morlet

Morlet wavelet

Syntax

```
[PSI,X] = morlet(LB,UB,N)
```

Description

[PSI,X] = morlet(LB,UB,N) returns values of the Morlet wavelet on an N point regular grid in the interval [LB,UB].

Output arguments are the wavelet function PSI computed on the grid X, and the grid X.

This wavelet has [-4 4] as effective support. Although [-4 4] is the correct theoretical effective support, a wider effective support, [-8 8], is used in the computation to provide more accurate results.

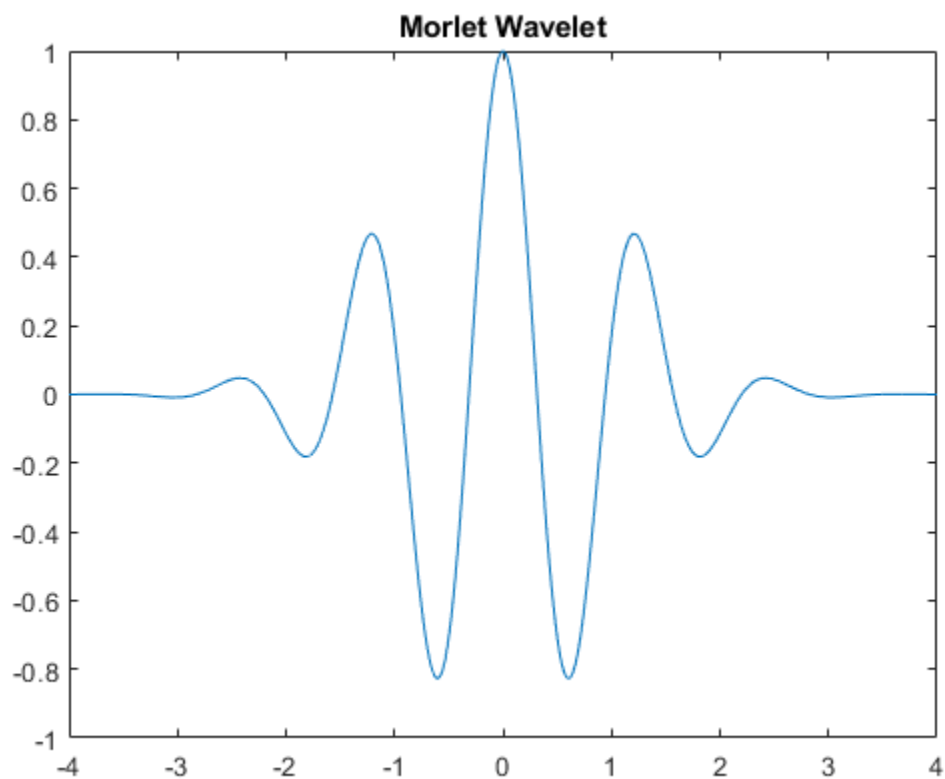
$$\psi(x) = e^{-x^2/2} \cos(5x)$$

Examples

Morlet Wavelet

This example shows how to create a Morlet wavelet. The wavelet has an effective support of [-4,4]. Use 1,000 sample points.

```
lb = -4;  
ub = 4;  
n = 1000;  
[psi,xval] = morlet(lb,ub,n);  
plot(xval,psi)  
title('Morlet Wavelet');
```

See Also

`waveinfo`

Introduced before R2006a

mswcmp

Multisignal 1-D compression using wavelets

Syntax

```
[XC,DECCMP,THRESH] = mswcmp('cmp',DEC,METH)
[XC,DECCMP,THRESH] = mswcmp('cmp',DEC,METH,PARAM)
[XC,THRESH] = mswcmp('cmpsig',...)
[DECCMP,THRESH] = mswcmp('cmpdec',...)
THRESH = mswcmp('thr',...)
[...] = mswcmp(OPTION,DIRDEC,X,WNAME,LEV,METH)
[...] = mswcmp(OPTION,DIRDEC,X,WNAME,LEV,METH,PARAM)
[...] = mswcmp(...,S_OR_H)
[...] = mswcmp(...,S_OR_H,KEEPAPP)
[...] = mswcmp(...,S_OR_H,KEEPAPP,IDXSIG)
```

Description

mswcmp computes thresholds and, depending on the selected option, performs compression of 1-D signals using wavelets.

[XC,DECCMP,THRESH] = mswcmp('cmp',DEC,METH) or
[XC,DECCMP,THRESH] = mswcmp('cmp',DEC,METH,PARAM) returns a compressed (indicated by 'cmp' input) version XC of the original multisignal matrix X, whose wavelet decomposition structure is DEC. The output XC is obtained by thresholding the wavelet coefficients: DECCMP, which is the wavelet decomposition associated with XC (see `mdwtdec`), and THRESH is the matrix of threshold values. The input METH is the name of the compression method and PARAM is the associated parameter, if required.

Valid compression methods METH are shown in the following tables. For methods that use an associated parameter, the range of allowable PARAM values is also shown.

'rem_n0'	Remove near 0
'bal_sn'	Balance sparsity-norm

'sqrtbal_sn'	Balance sparsity-norm (sqrt)
'scarce'	Scarce, PARAM (any number)
'scarcehi'	Scarce high, $2.5 \leq \text{PARAM} \leq 10$
'scarceme'	Scarce medium, $1.5 \leq \text{PARAM} \leq 2.5$
'scarcelo'	Scarce low, $1 \leq \text{PARAM} \leq 2$

PARAM is a sparsity parameter, and it should be such that: $1 \leq \text{PARAM} \leq 10$. For scarce method no control is done.

'L2_perf'	Energy ratio
'N0_perf'	Zero coefficients ratio

PARAM is a real number which represents the required performance:

$$0 \leq \text{PARAM} \leq 100.$$

'glb_thr'	Global threshold
-----------	------------------

PARAM is a real positive number.

'man_thr'	Manual method
-----------	---------------

PARAM is an NbSIG-by-NbLEV matrix or NbSIG-by-(NbLEV+1) matrix such that:

- - $\text{PARAM}(i, j)$ is the threshold for the detail coefficients of level j for the i th signal ($1 \leq j \leq \text{NbLEV}$).
- - $\text{PARAM}(i, \text{NbLEV}+1)$ is the threshold for the approximation coefficients for the i th signal (if KEEPAPP is 0).

Where NbSIG is the number of signals and NbLEV the number of levels of decomposition.

[XC, THRESH] = mswcmp('cmpsig', ...) or
 [DECCMP, THRESH] = mswcmp('cmpdec', ...) or
 THRESH = mswcmp('thr', ...) Instead of the 'cmp' input OPTION, you can use 'cmpsig', 'cmpdec' or 'thr' to select other output arguments. 'thr' returns the computed thresholds, but compression is not performed.

[...] = mswcmp(OPTION, DIRDEC, X, WNAME, LEV, METH)

[...] = mswcmp(OPTION,DIRDEC,X,WNAME,LEV,METH,PARAM) The decomposition structure input argument DEC can be replaced by four arguments: DIRDEC, X, WNAME, and LEV. Before performing a compression or computing thresholds, the multisignal matrix X is decomposed at level LEV using the wavelet WNAME, in the direction DIRDEC.

[...] = mswcmp(...,S_OR_H)

[...] = mswcmp(...,S_OR_H,KEEPAPP)

[...] = mswcmp(...,S_OR_H,KEEPAPP,IDXSIG) Three more optional inputs may be used:

- S_OR_H ('s' or 'h') stands for soft or hard thresholding (see mswthresh for more details). Default is 'h'.
- KEEPAPP (true or false) indicates whether to keep approximation coefficients (true) or not (false). Default is false.
- IDXSIG is a vector which contains the indices of the initial signals, or 'all'. Default is 'all'.

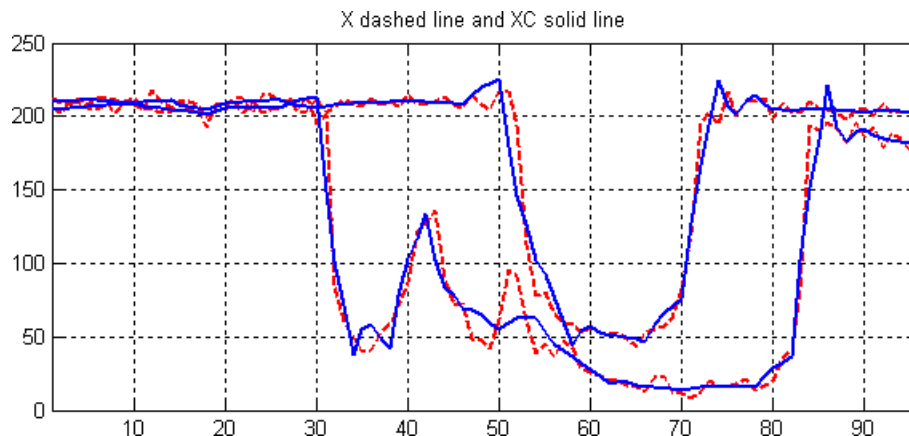
Examples

```
% Load original 1D-multisignal.
load thinker

% Perform a decomposition at level 2 using wavelet db2.
dec = mdwtdec('r',X,2,'db2');

% Compress the signals to obtain a percentage of zeros
% near 95% for the wavelet coefficients.
[XC,decCMP,THRESH] = mswcmp('cmp',dec,'N0_perf',95);
[Ecmp,PECcmp,PECFScmp] = wdecenergy(decCMP);

% Plot the original signals 1 and 31, and
% the corresponding compressed signals.
figure;
plot(X([1 31],:),'r--','linewidth',2); hold on
plot(XC([1 31],:),'b','linewidth',2);
grid; set(gca,'Xlim',[1,96])
title('X dashed line and XC solid line')
```



References

Birgé L.; P. Massart (1997), "From Model Selection to Adaptive Estimation," in D. Pollard (ed), *Festchrift for L. Le Cam*, Springer, pp. 55-88.

DeVore, R.A.; B. Jawerth, B.J. Lucier (1992), "Image Compression Through Wavelet Transform Coding," *IEEE Trans. on Inf. Theory*, vol. 38, No 2, pp. 719-746.

Donoho, D.L. (1993), "Progress in Wavelet Analysis and WVD: a Ten Minute Tour," in *Progress in Wavelet Analysis and Applications*, Y. Meyer, S. Roques, pp. 109-128. Frontières Ed.

Donoho, D.L.; I.M. Johnstone(1994), "Ideal Spatial Adaptation by Wavelet Shrinkage," *Biometrika*, vol. 81, pp. 425-455.

Donoho, D.L.; I.M. Johnstone, G. Kerkyacharian, D. Picard (1995), "Wavelet Shrinkage: Asymptopia?" *Jour. Roy. Stat. Soc., series B*, vol. 57 no. 2, pp. 301-369.

Donoho, D.L.; I.M. Johnstone, "Ideal De-noising in an Orthonormal Basis Chosen from a Library of Bases," *C.R.A.S. Paris*, t. 319, Ser. I, pp. 1317-1322.

Donoho, D.L. (1995), "De-noising by Soft-thresholding," *IEEE Trans. on Inf. Theory*, 41, 3, pp. 613-627.

See Also

[mdwtdec](#) | [mdwtrec](#) | [mswthresh](#) | [wthresh](#)

Introduced in R2007a

mswcmpscr

Multisignal 1-D wavelet compression scores

Syntax

`[THR, L2SCR, N0SCR, IDXSORT] = mswcmpscr(DEC)`

Description

`[THR, L2SCR, N0SCR, IDXSORT] = mswcmpscr(DEC)` computes four matrices: thresholds THR, compression scores L2SCR and N0SCR, and indices IDXSORT. The decomposition DEC corresponds to a matrix of wavelet coefficients CFS obtained by concatenation of detail and (optionally) approximation coefficients, where

$CFS = [cd\{DEC.level\}, \dots, cd\{1\}]$ or $CFS = [ca, cd\{DEC.level\}, \dots, cd\{1\}]$

The concatenation is made row-wise if `DEC.dirDec` is equal to 'r' or column-wise if `DEC.dirDec` is equal to 'c'.

If `NbSIG` is the number of original signals and `NbCFS` the number of coefficients for each signal (all or only the detail coefficients), then CFS is an `NbSIG`-by-`NbCFS` matrix.

Therefore,

- THR, L2SCR, N0SCR are `NbSIG`-by-`(NbCFS+1)` matrices
- IDXSORT is an `NbSIG`-by-`NbCFS` matrix
- `THR(:, 2:end)` is equal to CFS sorted by row in ascending order with respect to the absolute value.
- For each row, IDXSORT contains the order of coefficients and `THR(:, 1)=0`.

For the *i*th signal:

- `L2SCR(i, j)` is the percentage of preserved energy (L2-norm), corresponding to a threshold equal to `CFS(i, j-1)` ($2 \leq j \leq NbCFS$), and `L2SCR(:, 1)=100`.

- $N0SCR(i, j)$ is the percentage of zeros corresponding to a threshold equal to $CFS(i, j-1)$ ($2 \leq j \leq NbCFS$), and $N0SCR(:, 1)=0$.

Three more optional inputs may be used:

```
[...] = mswcmpscr(..., S_OR_H, KEEPAPP, IDXSIG)
```

- `S_OR_H` ('s' or 'h') stands for soft or hard thresholding (see `mswthresh` for more details).
- `KEEPAPP` (true or false) indicates whether to keep approximation coefficients (true) or not (false).
- `IDXSIG` is a vector that contains the indices of the initial signals, or 'all'.

The defaults are, respectively, 'h', false and 'all'.

Examples

```
% Load original 1D-multisignal.  
load thinker
```

```
% Perform a decomposition at level 2 using wavelet db2.  
dec = mdwtdec('r', X, 2, 'db2');
```

```
% Compute compression performances for soft an hard thresholding.  
[THR_S, L2SCR_S, N0SCR_S] = mswcmpscr(dec, 's');  
[THR_H, L2SCR_H, N0SCR_H] = mswcmpscr(dec, 'h');
```

References

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed.

Mallat, S. (1989), "A theory for multiresolution signal decomposition: the wavelet representation," *IEEE Pattern Anal. and Machine Intell.*, vol. 11, no. 7, pp. 674-693.

Meyer, Y. (1990), *Ondelettes et opérateurs*, Tome 1, Hermann Ed. (English translation: *Wavelets and operators*, Cambridge Univ. Press. 1993.)

See Also

`ddencmp` | `mdwtdec` | `mdwtrec` | `wdencmp`

Introduced in R2007a

mswcmptp

Multisignal 1-D compression thresholds and performances

Syntax

```
[THR_VAL,L2_Perf,N0_Perf] = mswcmptp(DEC,METH)
[THR_VAL,L2_Perf,N0_Perf] = mswcmptp(DEC,METH,PARAM)
```

Description

[THR_VAL,L2_Perf,N0_Perf] = mswcmptp(DEC,METH) or [THR_VAL,L2_Perf,N0_Perf] = mswcmptp(DEC,METH,PARAM) computes the vectors THR_VAL, L2_Perf and N0_Perf obtained after a compression using the METH method and, if required, the PARAM parameter (see mswcmp for more information on METH and PARAM).

For the *i*th signal:

- THR_VAL(*i*) is the threshold applied to the wavelet coefficients. For a level dependent method, THR_VAL(*i*, *j*) is the threshold applied to the detail coefficients at level *j*
- L2_Perf(*i*) is the percentage of energy (L2_norm) preserved after compression.
- N0_Perf(*i*) is the percentage of zeros obtained after compression.

You can use three more optional inputs:

```
[...] = mswcmptp(...,S_OR_H,KEEPAPP,IDXSIG)
```

- S_OR_H ('s' or 'h') stands for soft or hard thresholding (see mswthresh for more details).
- KEEPAPP (true or false) indicates whether to keep approximation coefficients (true) or not (false)
- IDXSIG is a vector which contains the indices of the initial signals, or 'all'.

The defaults are, respectively, 'h', false and 'all'.

Examples

```
% Load original 1D-multisignal.  
load thinker  
  
% Perform a decomposition at level 2 using wavelet db2.  
dec = mdwtdec('r',X,2,'db2');  
  
% Compute compression thresholds and exact performances  
% obtained after a compression using the method 'N0_perf' and  
% requiring a percentage of zeros near 95% for the wavelet  
% coefficients.  
[THR_VAL,L2_Perf,N0_Perf] = mswcmptp(dec,'N0_perf',95);
```

References

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics. SIAM Ed.

Mallat, S. (1989), "A theory for multiresolution signal decomposition: the wavelet representation," *IEEE Pattern Anal. and Machine Intell.*, vol. 11, no. 7, pp. 674-693.

Meyer, Y. (1990), *Ondelettes et opérateurs*, Tome 1, Hermann Ed. (English translation: *Wavelets and operators*, Cambridge Univ. Press. 1993.)

See Also

ddencmp | mdwtdec | mdwtrec | wdencmp

Introduced in R2007a

mswden

Multisignal 1-D denoising using wavelets

Syntax

```
[XD,DECDEN,THRESH] = mswden('den',...)  
[XD,THRESH] = mswden('densig',...)  
[DECDEN,THRESH] = mswden('dendec',...)  
THRESH = mswden('thr',...)  
[...] = mswden(OPTION,DIRDEC,X,WNAME,LEV,METH,PARAM)  
[...] = mswden(...,S_OR_H)  
[...] = mswden(...,S_OR_H,KEEPAPP)  
[...] = mswden(...,S_OR_H,KEEPAPP,IDXSIG)
```

Description

`mswden` computes thresholds and, depending on the selected option, performs denoising of 1-D signals using wavelets.

`[XD,DECDEN,THRESH] = mswden('den',...)` returns a denoised version `XD` of the original multisignal matrix `X`, whose wavelet decomposition structure is `DEC`. The output `XD` is obtained by thresholding the wavelet coefficients, `DECDEN` is the wavelet decomposition associated to `XD` (see `mdwtdec`), and `THRESH` is the matrix of threshold values. The input `METH` is the name of the denoising method and `PARAM` is the associated parameter, if required.

Valid denoising methods `METH` and associated parameters `PARAM` are:

'rigrsure'	Principle of Stein's Unbiased Risk
'heursure'	Heuristic variant of the first option
'sqtwolog'	Universal threshold $\sqrt{2 \cdot \log(\cdot)}$
'minimaxi'	Minimax thresholding (see <code>thselect</code>)

For these methods `PARAM` defines the multiplicative threshold rescaling:

'one'	No rescaling
'sln'	Rescaling using a single estimation of level noise based on first level coefficients
'mln'	Rescaling using a level dependent estimation of level noise

Penalization methods

'penal'	Penal
'penalhi'	Penal high, $2.5 \leq \text{PARAM} \leq 10$
'penalme'	Penal medium, $1.5 \leq \text{PARAM} \leq 2.5$
'penallo'	Penal low, $1 \leq \text{PARAM} \leq 2$

PARAM is a sparsity parameter, and it should be such that: $1 \leq \text{PARAM} \leq 10$. For penal method, no control is done.

Manual method

'man_thr'	Manual method
-----------	---------------

PARAM is an NbSIG-by-NbLEV matrix or NbSIG-by-(NbLEV+1) matrix such that:

- $\text{PARAM}(i, j)$ is the threshold for the detail coefficients of level j for the i th signal ($1 \leq j \leq \text{NbLEV}$).
- $\text{PARAM}(i, \text{NbLEV}+1)$ is the threshold for the approximation coefficients for the i th signal (if KEEPAPP is 0).

where NbSIG is the number of signals and NbLEV the number of levels of decomposition.

Instead of the 'den' input OPTION, you can use 'densig', 'dendec' or 'thr' OPTION to select output arguments:

```
[XD, THRESH] = mswden('densig', ...) or [DEC DEN, THRESH] =
mswden('dendec', ...)
```

THRESH = mswden('thr', ...) returns the computed thresholds, but denoising is not performed.

The decomposition structure input argument DEC can be replaced by four arguments: DIRDEC, X, WNAME and LEV.

[...] = mswden(OPTION, DIRDEC, X, WNAME, LEV, METH, PARAM) before performing a denoising or computing thresholds, the multisignal matrix X is decomposed at level LEV using the wavelet WNAME, in the direction DIRDEC.

You can use three more optional inputs:

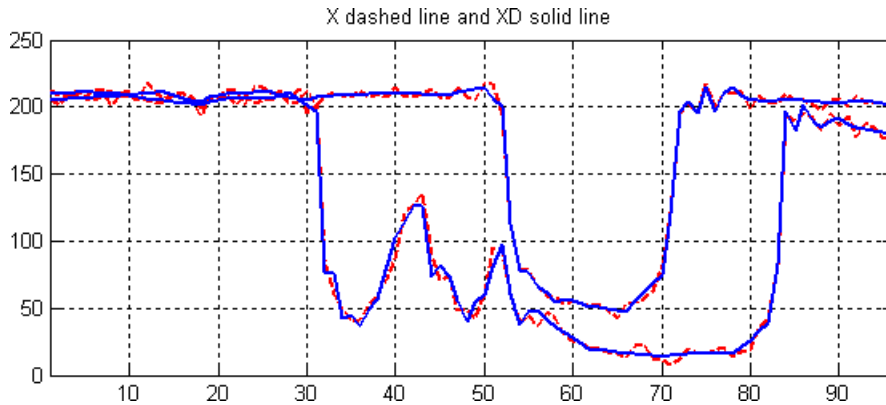
```
[...] = mswden(..., S_OR_H) or  
[...] = mswden(..., S_OR_H, KEEPAPP) or  
[...] = mswden(..., S_OR_H, KEEPAPP, IDXSIG)
```

- S_OR_H ('s' or 'h') stands for soft or hard thresholding (see mswthresh for more details).
- KEEPAPP (true or false) indicates whether to keep approximation coefficients (true) or not (false).
- IDXSIG is a vector that contains the indices of the initial signals, or 'all'.

The defaults are, respectively, 'h', false and 'all'.

Examples

```
% Load original 1D-multisignal.  
load thinker  
  
% Perform a decomposition at level 2 using the wavelet db2.  
dec = mdwtdec('r', X, 2, 'db2');  
  
% Denoise signals using the universal method  
% of thresholding (sqrtwolog) and the 'sln'  
% threshold rescaling (with a single estimation  
% of level noise, based on first level coefficients).  
[XD, decDEN, THRESH] = mswden('den', dec, 'sqrtwolog', 'sln');  
  
% Plot the original signals 1 and 31, and the  
% corresponding denoised signals.  
figure;  
plot(X([1 31],:)), 'r--', 'linewidth', 2); hold on  
plot(XD([1 31],:)), 'b', 'linewidth', 2);  
grid; set(gca, 'Xlim', [1, 96])  
title('X dashed line and XD solid line')
```



References

Birgé, L.; P. Massart (1997), "From model selection to adaptive estimation," in D. Pollard (ed), *Festchrift for L. Le Cam*, Springer, pp. 55-88.

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Donoho, D.L. (1993), "Progress in wavelet analysis and WVD: a ten minute tour," in *Progress in wavelet analysis and applications*, Y. Meyer, S. Roques, pp. 109-128. Frontières Ed.

Donoho, D.L.; I.M. Johnstone(1994), "Ideal spatial adaptation by wavelet shrinkage," *Biometrika*, vol. 81, pp. 425-455.

Donoho, D.L.; I.M. Johnstone, G. Kerkyacharian, D. Picard (1995), "Wavelet shrinkage: asymptopia," *Jour. Roy. Stat. Soc., series B*, vol. 57 no. 2, pp. 301-369.

Donoho, D.L.; I.M. Johnstone, "Ideal de-noising in an orthonormal basis chosen from a library of bases," *C.R.A.S. Paris*, t. 319, Ser. I, pp. 1317-1322.

Donoho, D.L. (1995), "De-noising by soft-thresholding," *IEEE Trans. on Inf. Theory*, 41, 3, pp. 613-627.

See Also

Functions

`mdwtdec` | `mdwtrec` | `mswthresh` | `wdenoise` | `wthresh`

Apps

Wavelet Signal Denoiser

Introduced in R2007a

mswthresh

Perform multisignal 1-D thresholding

Syntax

```
Y = mswthresh(X,sorh,T)
Y = mswthresh(X,sorh,T,'c')
```

Description

`Y = mswthresh(X,sorh,T)` returns the soft or hard T -thresholding of the matrix X . T can be a single value, a matrix the same size as X , or a vector. If T is a vector, thresholding is performed row-wise, and $LT = \text{length}(T)$ must be such that $\text{size}(X,1) \leq LT$. Only the first $\text{size}(X,1)$ values of T are used.

`Y = mswthresh(X,sorh,T,'c')` performs thresholding column-wise, and $LT = \text{length}(T)$ must be such that $\text{size}(X,2) \leq LT$. Only the first $\text{size}(X,2)$ values of T are used.

Examples

Multisignal Thresholding

Create a 3-by-3 matrix and a 1-by-3 vector of threshold values.

```
mat = [1 1 3; 1 1 3; 2 2 3]
```

```
mat = 3×3
```

```
    1    1    3
    1    1    3
    2    2    3
```

```
thr = [1 2 3]
```

```
thr = 1×3  
    1    2    3
```

Apply soft thresholding to the matrix row-wise. The k th threshold in `thr` is applied to the k th row of `mat`.

```
mswthresh(mat, 's', thr)
```

```
ans = 3×3  
    0    0    2  
    0    0    1  
    0    0    0
```

Apply soft thresholding to the matrix column-wise. The k th threshold in `thr` is applied to the k th column of `mat`.

```
mswthresh(mat, 's', thr, 'c')
```

```
ans = 3×3  
    0    0    0  
    0    0    0  
    1    0    0
```

Apply hard thresholding to the matrix row-wise.

```
mswthresh(mat, 'h', thr)
```

```
ans = 3×3  
    0    0    3  
    0    0    3  
    0    0    0
```

Apply hard thresholding to the matrix column-wise.

```
mswthresh(mat, 'h', thr, 'c')
```

```
ans = 3×3
```

```
0    0    0
0    0    0
2    0    0
```

Input Arguments

X — Input data

real-valued matrix

Input data to threshold, specified as a real-valued matrix.

Data Types: double

sorh — Type of thresholding

's' | 'h'

Type of thresholding to perform, specified as:

- 's' — Soft thresholding
- 'h' — Hard thresholding

T — Threshold value

real-valued scalar or vector

Threshold value, specified as a real-valued scalar or vector.

Data Types: double

Output Arguments

Y — Thresholded data

real-valued matrix

Thresholded data, returned as a real-valued matrix. Y has the same dimensions as X.

Algorithms

- If sorh is 's', Y is the soft thresholding of X: $Y = \text{sign}(X) \cdot (|X| - T)_+$ where

$$(x)_+ = \begin{cases} x & \text{if } x \geq 0 \\ 0 & \text{otherwise} \end{cases}$$

Soft thresholding is wavelet shrinkage.

- If sorh is 'h', Y is the hard thresholding of X: $Y = X \cdot \mathbf{1}_{(|X| > T)}$ where

$$\mathbf{1}_{(|X| > T)} = \begin{cases} 1 & \text{if } |X| > T \\ 0 & \text{otherwise} \end{cases}$$

Hard thresholding is cruder than soft thresholding.

See Also

`mswcmp` | `mswden` | `wden` | `wdencomp` | `wdenoise` | `wdencomp` | `wthresh`

Introduced in R2007a

nodeasc

Node ascendants

Syntax

`A = nodeasc(T,N)`

Description

nodeasc is a tree-management utility.

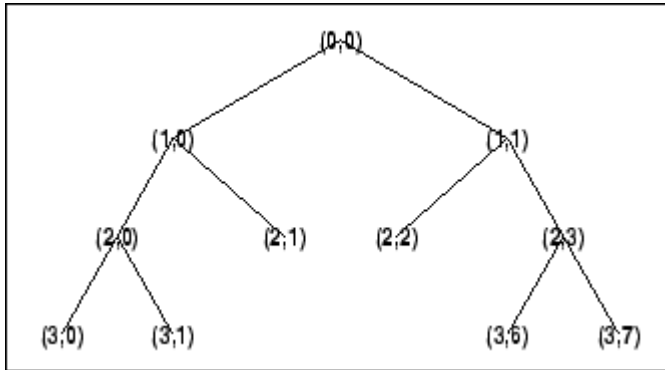
`A = nodeasc(T,N)` returns the indices of all the ascendants of the node N in the tree T where N can be the index node or the depth and position of the node. A is a column vector with $A(1) = \text{index of node } N$.

`A = nodeasc(T,N, 'deppos')` is a matrix, which contains the depths and positions of all ascendants. $A(i, 1)$ is the depth of the i -th ascendant and $A(i, 2)$ is the position of the i -th ascendant.

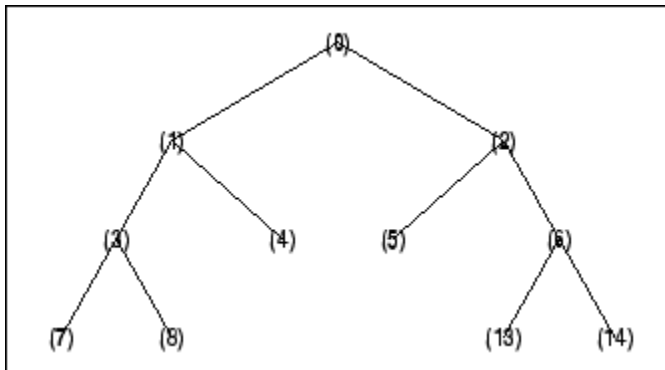
The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

```
% Create binary tree of depth 3.  
t = ntree(2,3);  
t = nodejoin(t,5);  
t = nodejoin(t,4);  
plot(t)
```



% Change Node Label from Depth_Position to Index
 % (see the plot function).



```

nodeasc(t,[2 2])
ans =
     5
     2
     0
  
```

```

nodeasc(t,[2 2], 'depos')
ans =
     2     2
     1     1
     0     0
  
```

See Also

nodedesc | nodepar | wt reemgr

Introduced before R2006a

nodedesc

Node descendants

Syntax

```
D = nodedesc(T,N)
D = nodedesc(T,N, 'deppos')
```

Description

`nodedesc` is a tree-management utility.

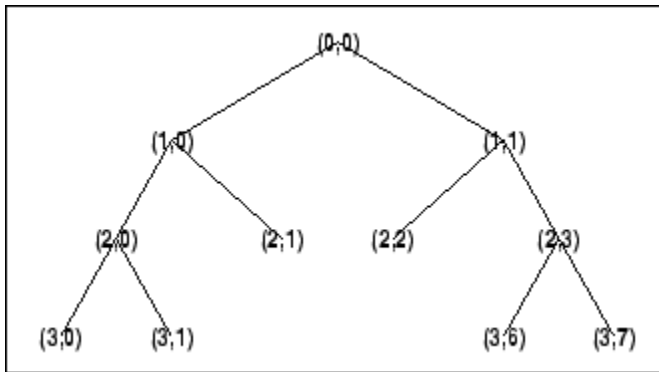
`D = nodedesc(T,N)` returns the indices of all the descendants of the node N in the tree T where N can be the index node or the depth and position of node. D is a column vector with $D(1) = \text{index of node } N$.

`D = nodedesc(T,N, 'deppos')` is a matrix that contains the depths and positions of all descendants. $D(i,1)$ is the depth of the i -th descendant and $D(i,2)$ is the position of the i -th descendant.

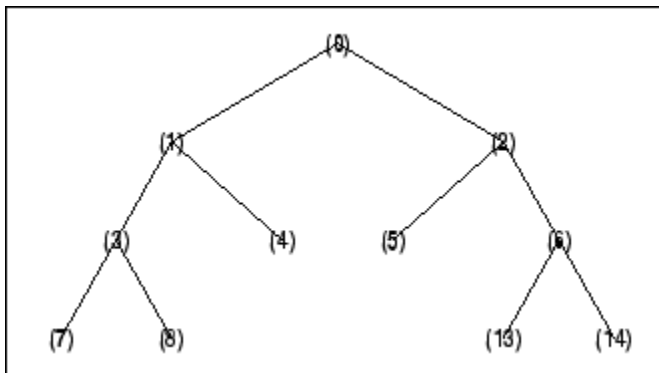
The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

```
% Create binary tree of depth 3.
t = ntree(2,3);
t = nodejoin(t,5);
t = nodejoin(t,4);
plot(t)
```

% Change Node Label from Depth_Position to Index
 % (see the plot function).



% Node descendants.

nodedesc(t,2)

```
ans =
    2
    5
    6
    13
    14
```

nodedesc(t,2, 'depos')

```
ans =
    1    1
    2    2
    2    3
```

```
3    6
3    7
```

```
nodeDESC(t,[1 1], 'depos')
```

```
ans =
     1     1
     2     2
     2     3
     3     6
     3     7
```

```
nodeDESC(t,[1 1])
```

```
ans =
     2
     5
     6
    13
    14
```

See Also

[nodeasc](#) | [nodepar](#) | [wtreemgr](#)

Introduced before R2006a

nodejoin

Recompose node

Syntax

```
T = nodejoin(T,N)
T = nodejoin(T)
T = nodejoin(T,0)
```

Description

nodejoin is a tree-management utility.

$T = \text{nodejoin}(T,N)$ returns the modified tree T corresponding to a recomposition of the node N .

The nodes are numbered from left to right and from top to bottom. The root index is 0.

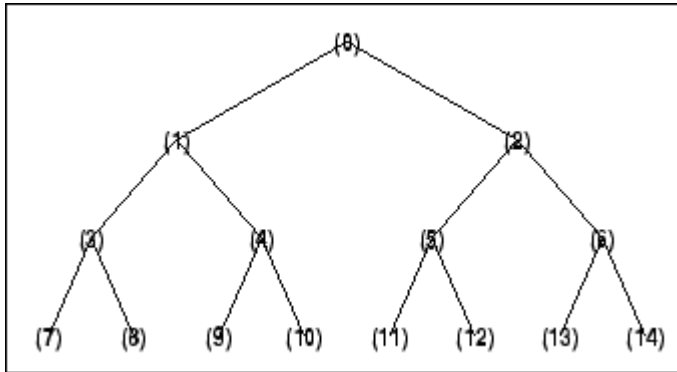
$T = \text{nodejoin}(T)$ is equivalent to $T = \text{nodejoin}(T,0)$.

Examples

```
% Create binary tree of depth 3.
t = ntree(2,3);

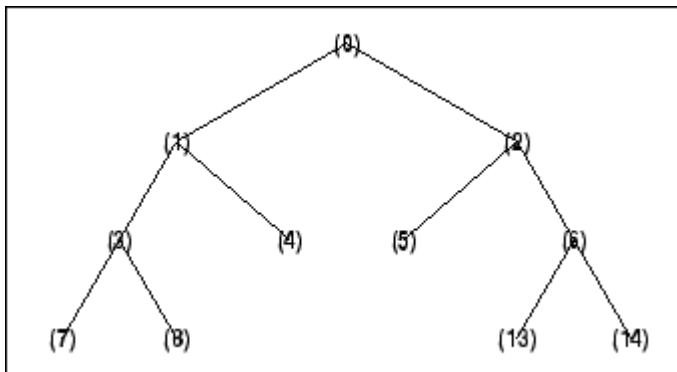
% Plot tree t.
plot(t)

% Change Node Label from Depth_Position to Index
% (see the plot function).
```



```
% Merge nodes of indices 4 and 5.  
t = nodejoin(t,5);  
t = nodejoin(t,4);  
% Plot new tree t.  
plot(t)
```

```
% Change Node Label from Depth_Position to Index  
% (see the plot function).
```



See Also

nodesplt

Introduced before R2006a

nodepar

Node parent

Syntax

```
F = nodepar(T,N)
F = nodepar(T,N, 'deppos')
```

Description

nodepar is a tree-management utility.

`F = nodepar(T,N)` returns the indices of the parent(s) of the nodes N in the tree T where N can be a column vector containing the indices of nodes or a matrix that contains the depths and positions of nodes. In the last case, $N(i,1)$ is the depth of the i -th node and $N(i,2)$ is the position of the i -th node.

`F = nodepar(T,N, 'deppos')` is a matrix that contains the depths and positions of returned nodes. $F(i,1)$ is the depth of the i -th node and $F(i,2)$ is the position of the i -th node.

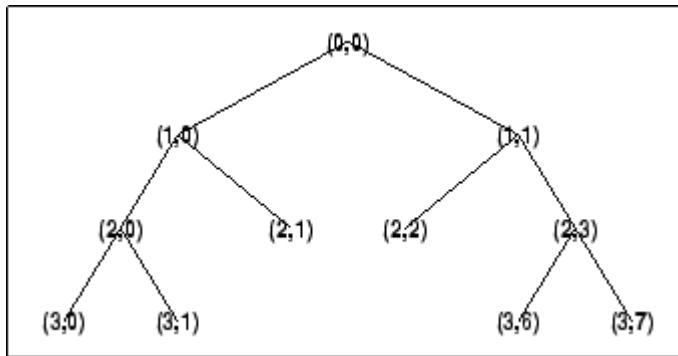
`nodepar(T,0)` or `nodepar(T,[0,0])` returns -1.

`nodepar(T,0, 'deppos')` or `nodepar(T,[0,0], 'deppos')` returns [-1,0].

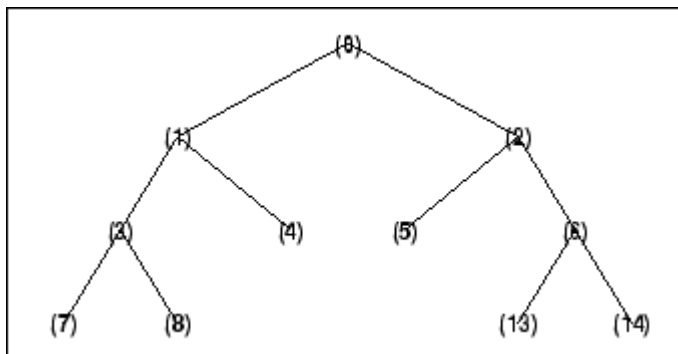
The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

```
% Create binary tree of depth 3.
t = ntree(2,3);
t = nodejoin(t,5);
t = nodejoin(t,4);
plot(t)
```



```
% Change Node Label from Depth_Position to Index
% (see the plot function).
```



```
% Nodes parent.
nodepar(t,[2 2], 'depos')
```

```
ans =
     1     1
```

```
nodepar(t,[1;7;14])
```

```
ans =
     0
     3
     6
```

See Also

nodeasc | nodedesc | wt reemgr

Introduced before R2006a

nodesplt

Split (decompose) node

Syntax

```
T = nodesplt(T,N)
```

Description

nodesplt is a tree-management utility.

`T = nodesplt(T,N)` returns the modified tree T corresponding to the decomposition of the node N .

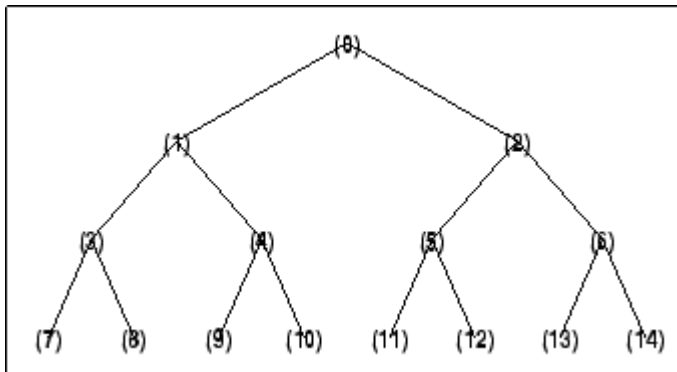
The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

```
% Create binary tree (tree of order 2) of depth 3.  
t = ntree(2,3);
```

```
% Plot tree t.  
plot(t)
```

```
% Change Node Label from Depth_Position to Index  
% (see the plot function).
```

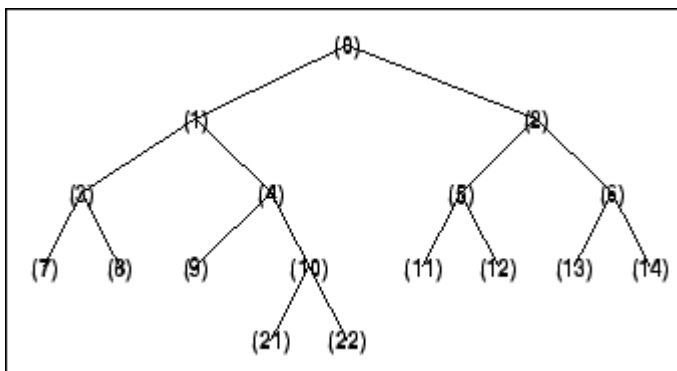



```

% Split node of index 10.
t = nodesplt(t,10);

% Plot new tree t.
plot(t)
% Change Node Label from Depth_Position to Index
% (see the plot function).

```



See Also

nodejoin

Introduced before R2006a

noleaves

Determine nonterminal nodes

Syntax

```
N = noleaves(T)
N = noleaves(T, 'dp')
```

Description

`N = noleaves(T)` returns the indices of nonterminal nodes of the tree T (i.e., nodes that are not leaves). N is a column vector.

The nodes are ordered from left to right as in tree T .

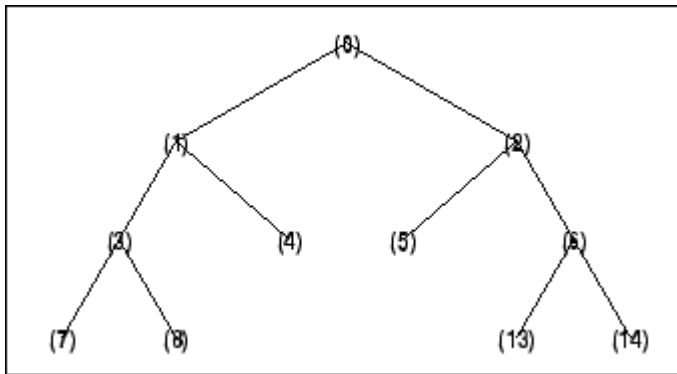
`N = noleaves(T, 'dp')` returns a matrix N , which contains the depths and positions of nonterminal nodes.

$N(i,1)$ is the depth of the i -th nonterminal node and
 $N(i,2)$ is the position of the i -th nonterminal node.

Examples

```
% Create initial tree.
ord = 2;
t = ntree(ord,3);          % binary tree of depth 3.
t=nodejoin(t,5);
t=nodejoin(t,4);
plot(t)

% Change Node Label from Depth_Position to Index
% (see the plot function).
```



```
% List nonterminal nodes (index).
ntnodes_ind = noleaves(t)
```

```
ntnodes_ind =
    0
    1
    2
    3
    6
```

```
% List nonterminal nodes (Depth_Position).
ntnodes_depo = noleaves(t,'dp')
```

```
ntnodes_depo =
    0     0
    1     0
    1     1
    2     0
    2     3
```

See Also

leaves

Introduced before R2006a

ntnode

Number of terminal nodes

Syntax

```
NB = ntnode(T)
```

Description

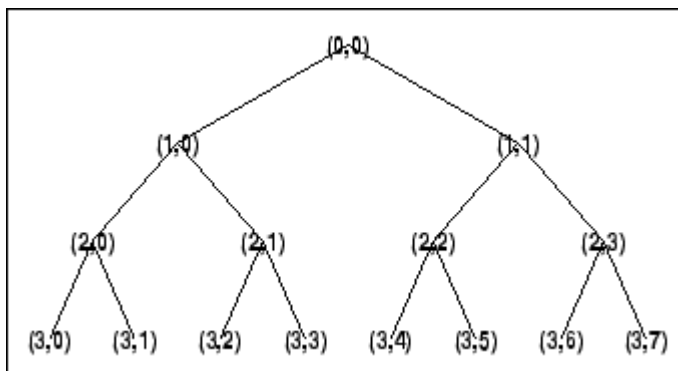
ntnode is a tree-management utility.

NB = ntnode(T) returns the number of terminal nodes in the tree T.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

Examples

```
% Create binary tree (tree of order 2) of depth 3.  
t = ntree(2,3);  
  
% Plot tree t.  
plot(t)
```



```
% Number of terminal nodes.  
ntnode(t)
```

```
ans =  
      8
```

See Also

wtreemgr

Introduced before R2006a

ntree

NTREE constructor

Syntax

```
T = ntree(ORD,D)
T = ntree
T = ntree(2,0)
T = ntree(ORD)
T = ntree(ORD,0)
T = ntree(ORD,D,S)
T = ntree(ORD,D,S,U)
```

Description

`T = ntree(ORD,D)` returns an NTREE object, which is a complete tree of order `ORD` and depth `D`.

`T = ntree` is equivalent to `T = ntree(2,0)`.

`T = ntree(ORD)` is equivalent to `T = ntree(ORD,0)`.

With `T = ntree(ORD,D,S)` you can set a “split scheme” for nodes. The split scheme field `S` is a logical array of size `ORD` by 1.

The root of the tree can be split and it has `ORD` children. You can split the `j`-th child if `S(j) = 1`.

Each node that you can split has the same property as the root node.

With `T = ntree(ORD,D,S,U)` you can, in addition, set a userdata field.

Inputs can be given in another way:

`T = ntree('order',ORD,'depth',D,'spsch',S,'ud',U)`. For “missing” inputs the defaults are `ORD = 2` and `D = 0`, `S = ones([1:ORD])`, `U = {}`.

[T,NB] = ntree(...) returns also the number of terminal nodes (leaves) of T.

For more information on object fields, type help ntree/get.

Class NTREE (Parent class: WTBO)

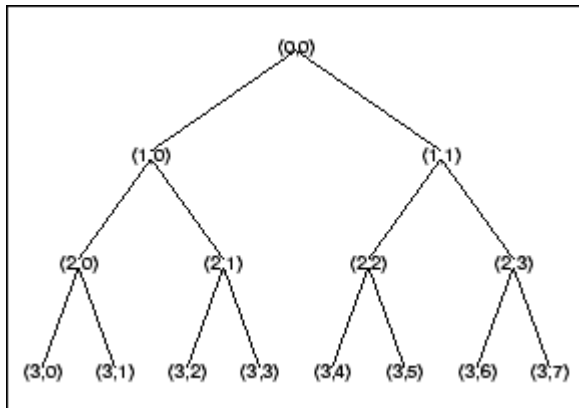
Fields

wtbo	Parent object
order	Tree order
depth	Tree depth
spsch	Split scheme for nodes
tn	Column vector with terminal node indices

Examples

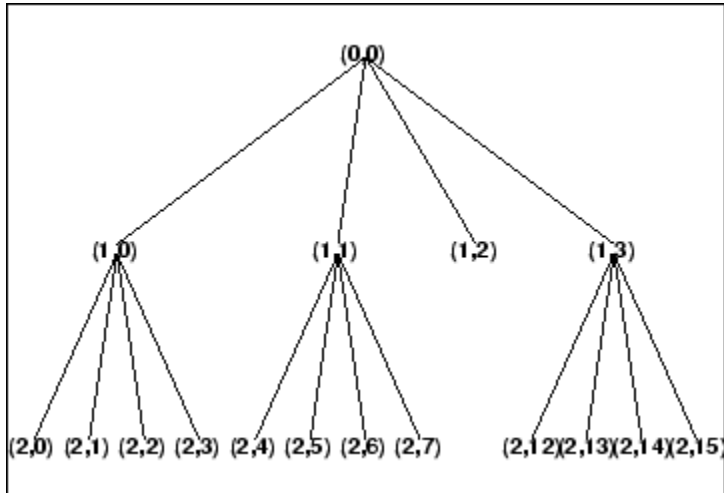
```
% Create binary tree (tree of order 2) of depth 3.
t2 = ntree(2,3);
```

```
% Plot tree t2.
plot(t2)
```

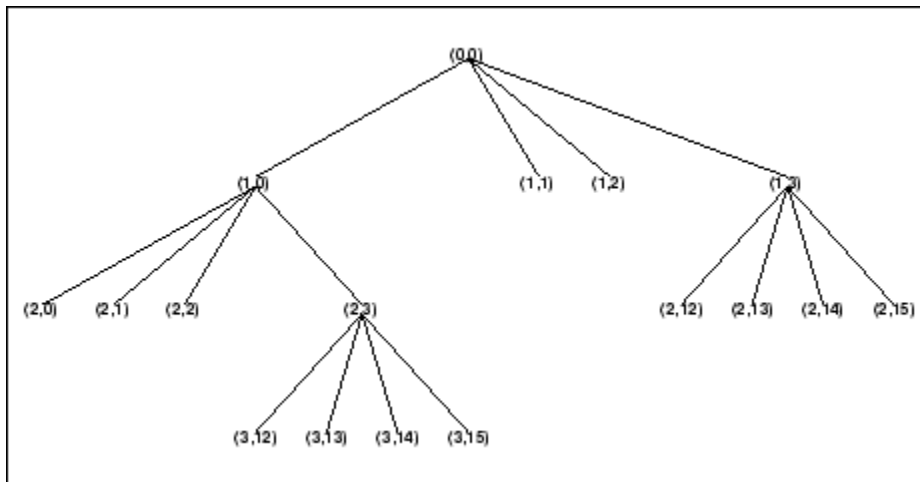


```
% Create a quadtree (tree of order 4) of depth 2.
t4 = ntree(4,2,[1 1 0 1]);
```

```
% Plot tree t4.  
plot(t4)
```



```
% Split and merge some nodes using the gui  
% generated by plot (see the plot function).  
% The figure becomes:
```



See Also

wtbo

Introduced before R2006a

numCoefficients

Number of wavelet scattering coefficients

Syntax

```
ncf = numCoefficients(sf)
```

Description

`ncf = numCoefficients(sf)` returns the number of scattering coefficients for each scattering path in the scattering framework `sf`. The number of scattering coefficients depends on the values of the “SignalLength” on page 1-0 , “InvarianceScale” on page 1-0 , and “OversamplingFactor” on page 1-0 properties of `sf`.

Examples

Oversample 1-D Wavelet Scattering Transform

This example shows how to oversample a 1-D wavelet scattering transform.

Load an ECG signal sampled at 180 Hz, and create a wavelet scattering framework to process the signal. To perform a critically downsampled wavelet scattering transform, do not change the value of the `OversamplingFactor` property in `sf`. Return the number of scattering coefficients for the framework.

```
load wecg
Fs = 180;
sf = waveletScattering('SignalLength',numel(wecg),'SamplingFrequency',Fs);
ncf = numCoefficients(sf)
```

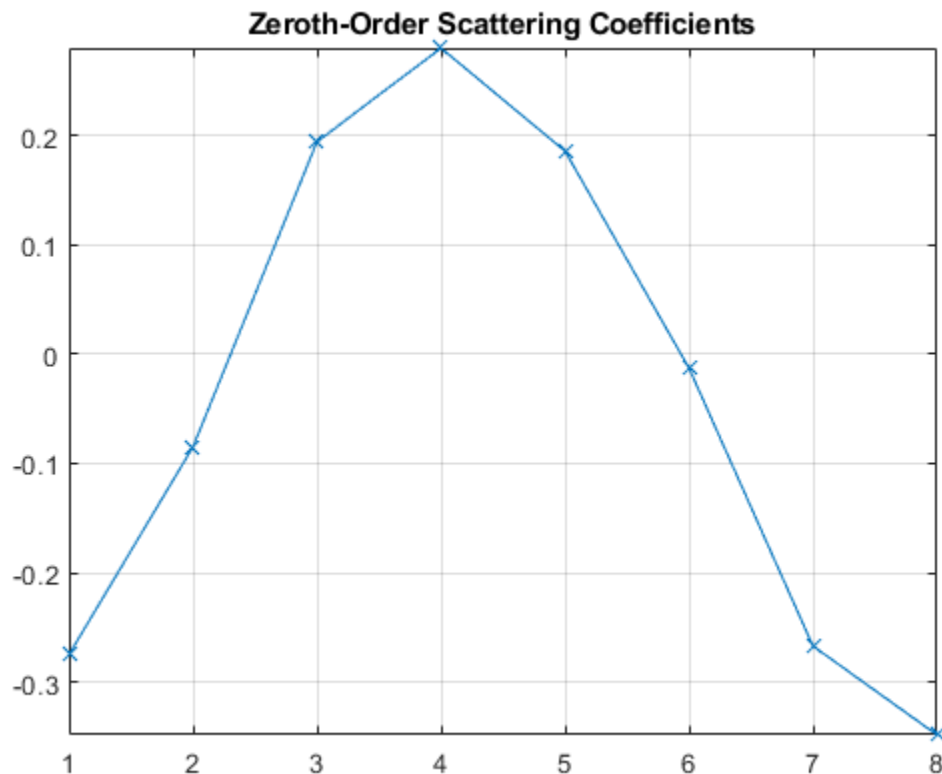
```
ncf = 8
```

Return the 1-D wavelet scattering transform of `wecg`, and plot the zeroth-order scattering coefficients. Confirm the number of zeroth-order scattering coefficients is equal to `ncf`.

```
s = scatteringTransform(sf,wecg);  
display(['Number of zeroth-order scattering coefficients: ',...  
        num2str(numel(s{1}.signals{1}))])
```

Number of zeroth-order scattering coefficients: 8

```
plot(s{1}.signals{1},'x-')  
grid on  
axis tight  
title('Zeroth-Order Scattering Coefficients')
```



To oversample the scattering coefficients by a factor of 2, set the `OversamplingFactor` property of `sf` equal to 1 (because $\log_2 1 = 1$). Return the number of scattering

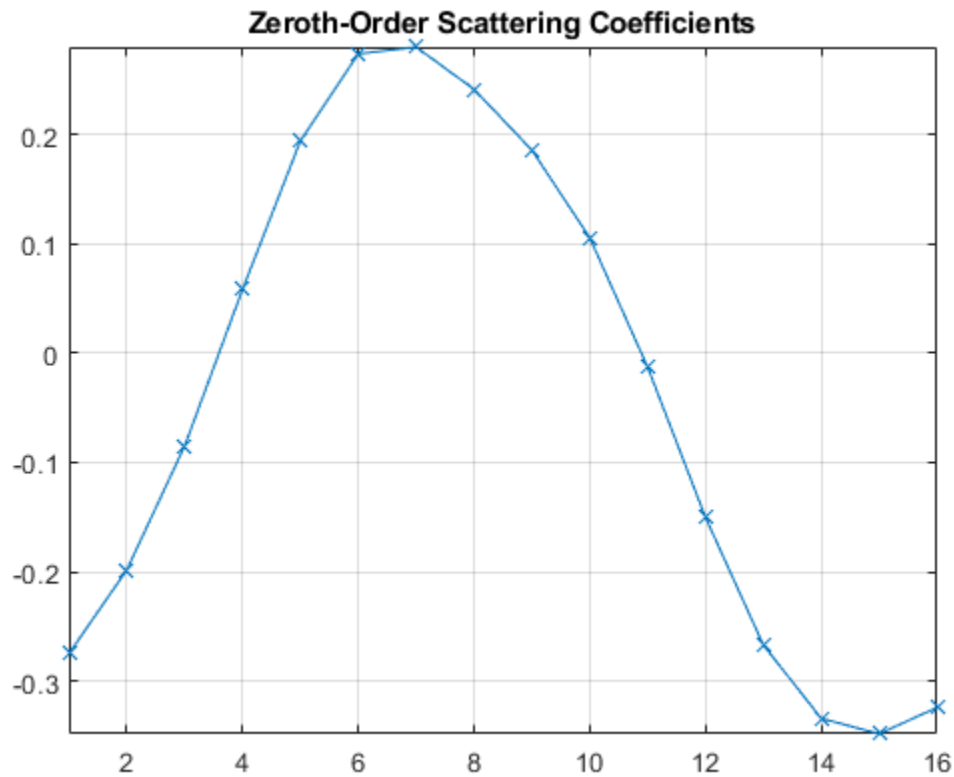
coefficients for the edited framework. Confirm the number of scattering coefficients has doubled.

```
sf.OversamplingFactor = 1;  
ncf = numCoefficients(sf)
```

```
ncf = 16
```

Return the wavelet scattering transform of `wecg` using the edited framework, and plot the zeroth-order scattering coefficients. Since the number of coefficients in the critically sampled transform is equal to 8, confirm that the number of zeroth-order coefficients in the oversampled transform is equal to 16.

```
s = scatteringTransform(sf,wecg);  
figure  
plot(s{1}.signals{1}, 'x-')  
grid on  
axis tight  
title('Zeroth-Order Scattering Coefficients')
```



Input Arguments

sf — Scattering decomposition framework

waveletScattering object

Scattering decomposition framework, specified as a waveletScattering object.

See Also

waveletScattering

Introduced in R2019a

numfilterbanks

Number of scattering filter banks

Syntax

```
nfb = numfilterbanks(sf)
```

Description

`nfb = numfilterbanks(sf)` returns the number of filter banks in the scattering decomposition framework, `sf`. The number of filter banks in a scattering decomposition framework is equal to $ord - 1$ where ord is the number of scattering orders.

Examples

Number of Filter Banks in Scattering Decomposition Framework

Calculate the number of filter banks for the default scattering decomposition framework.

```
sf = waveletScattering
```

```
sf =  
    waveletScattering with properties:
```

```
        SignalLength: 1024  
        InvarianceScale: 512  
        QualityFactors: [8 1]  
            Boundary: 'periodic'  
        SamplingFrequency: 1  
            Precision: 'double'  
        OversamplingFactor: 0
```

```
Nfb = numfilterbanks(sf)
```

```
Nfb = 2
```

Input Arguments

sf — Scattering decomposition framework

`waveletScattering` object | `waveletScattering2` object

Scattering decomposition framework, specified as a `waveletScattering` object or a `waveletScattering2` object.

See Also

`waveletScattering` | `waveletScattering2`

Introduced in R2018b

numorders

Number of scattering orders

Syntax

```
no = numorders(sf)
```

Description

`no = numorders(sf)` returns the number of orders for the scattering decomposition framework, `sf`. The number of orders is equal to $Nfb + 1$, where Nfb is the number of filter banks in `sf`.

Examples

Number of Orders in Scattering Decomposition Framework

Calculate the number of orders for the default scattering decomposition framework.

```
sf = waveletScattering
```

```
sf =  
    waveletScattering with properties:
```

```
        SignalLength: 1024  
        InvarianceScale: 512  
        QualityFactors: [8 1]  
            Boundary: 'periodic'  
        SamplingFrequency: 1  
            Precision: 'double'  
        OversamplingFactor: 0
```

```
no = numorders(sf)
```

```
no = 3
```

Input Arguments

sf – Scattering decomposition framework

`waveletScattering` object | `waveletScattering2` object

Scattering decomposition framework, specified as a `waveletScattering` object or a `waveletScattering2` object.

See Also

`waveletScattering` | `waveletScattering2`

Introduced in R2018b

orthfilt

Orthogonal wavelet filter set

Syntax

[Lo_D,Hi_D,Lo_R,Hi_R] = orthfilt(W)

Description

[Lo_D,Hi_D,Lo_R,Hi_R] = orthfilt(W) computes the four filters associated with the scaling filter W corresponding to a wavelet:

Lo_D	Decomposition low-pass filter
Hi_D	Decomposition high-pass filter
Lo_R	Reconstruction low-pass filter
Hi_R	Reconstruction high-pass filter

For an orthogonal wavelet, in the multiresolution framework, we start with the scaling function ϕ and the wavelet function ψ . One of the fundamental relations is the twin-scale relation:

$$\frac{1}{2}\phi\left(\frac{x}{2}\right) = \sum_{n \in Z} w_n \phi(x - n)$$

All the filters used in `dwt` and `idwt` are intimately related to the sequence $(w_n)_{n \in Z}$. Clearly if ϕ is compactly supported, the sequence (w_n) is finite and can be viewed as a FIR filter. The scaling filter W is

- A low-pass FIR filter
- Of length $2N$
- Of sum 1
- Of norm $\frac{1}{\sqrt{2}}$

For example, for the db3 scaling filter,

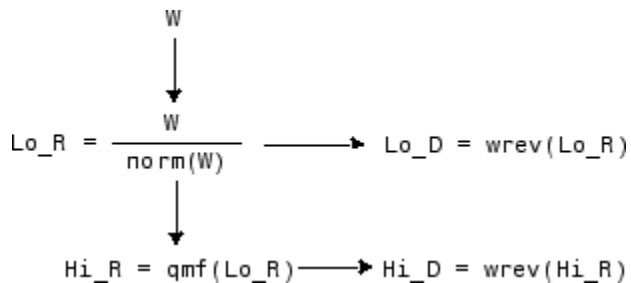
```
load db3
db3
db3 =
    0.2352  0.5706  0.3252 -0.0955 -0.0604  0.0249

sum(db3)
ans =
    1.000
norm(db3)
ans =
    0.7071
```

From filter W , we define four FIR filters, of length $2N$ and norm 1, organized as follows:

Filters	Low-Pass	High-Pass
Decomposition	Lo_D	Hi_D
Reconstruction	Lo_R	Hi_R

The four filters are computed using the following scheme:



where qmf is such that Hi_R and Lo_R are quadrature mirror filters (i.e., $\text{Hi}_R(k) = (-1)^k \text{Lo}_R(2N + 1 - k)$, for $k = 1, 2, \dots, 2N$), and where wrev flips the filter coefficients. So Hi_D and Lo_D are also quadrature mirror filters. The computation of these filters is performed using `orthfilt`.

Examples

```
% Load scaling filter.
load db8; w = db8;
```

```

subplot(421); stem(w);
title('Original scaling filter');

% Compute the four filters.
[Lo_D,Hi_D,Lo_R,Hi_R] = orthfilt(w);
subplot(423); stem(Lo_D);
title('Decomposition low-pass filter');
subplot(424); stem(Hi_D);
title('Decomposition high-pass filter');
subplot(425); stem(Lo_R);
title('Reconstruction low-pass filter');
subplot(426); stem(Hi_R);
title('Reconstruction high-pass filter');

% Check for orthonormality.
df = [Lo_D;Hi_D];
rf = [Lo_R;Hi_R];
id = df*df'

id =
    1.0000         0
         0    1.0000

id = rf*rf'

id =
    1.0000         0
         0    1.0000

% Check for orthogonality by dyadic translation, for example:
df = [Lo_D 0 0;Hi_D 0 0];
dft = [0 0 Lo_D; 0 0 Hi_D];
zer = df*dft'

zer =

    1.0e-12 *
   -0.1883  0.0000
   -0.0000 -0.1883

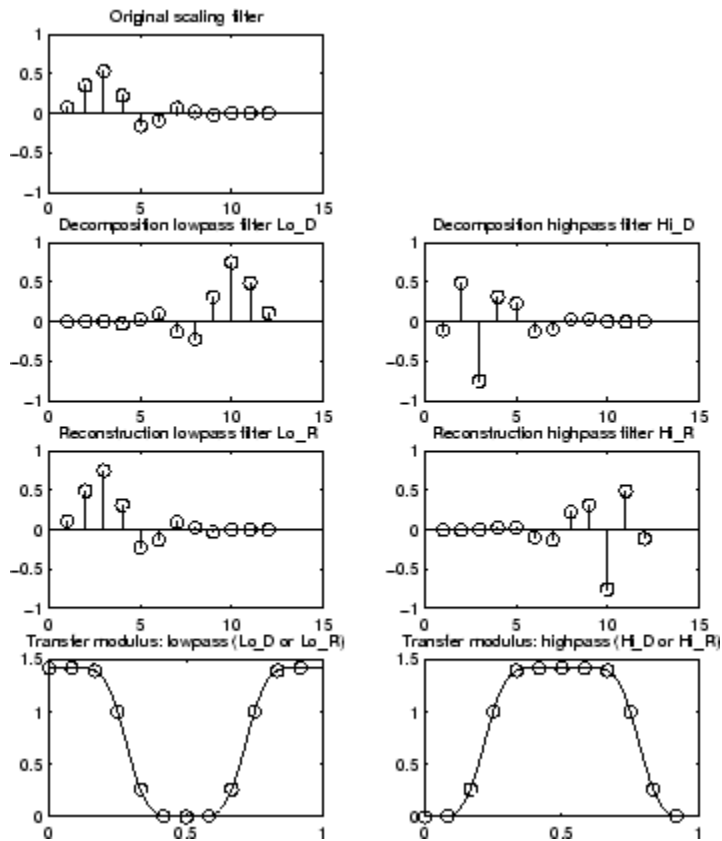
% High- and low-frequency illustration.
fftld = fft(Lo_D); ffthd = fft(Hi_D);
freq = [1:length(Lo_D)]/length(Lo_D);
subplot(427); plot(freq,abs(fftld));

```

```

title('Transfer modulus: low-pass');
subplot(4,2); plot(freq,abs(ffthd));
title('Transfer modulus: high-pass')
% Editing some graphical properties,
% the following figure is generated.

```



References

Daubechies, I. (1992), *Ten lectures on wavelets*, CBMS-NSF conference series in applied mathematics, SIAM Ed. pp. 117-119, 137, 152.

See Also

biorfilt | qmf | wfilters

Introduced before R2006a

otnodes

Order terminal nodes of binary wavelet packet tree

Syntax

```
[Tn_Pal,Tn_Seq] = otnodes(WPT)  
[Tn_Pal,Tn_Seq,I,J] = otnodes(WPT)  
[DP_Pal,DP_Seq] = otnodes(WPT,'dp')
```

Description

[Tn_Pal,Tn_Seq] = otnodes(WPT) returns the terminal nodes of the binary wavelet packet tree, WPT, in Paley (natural) ordering, Tn_Pal, and sequency (frequency) ordering, Tn_Seq. Tn_Pal and Tn_Seq are N -by-1 column vectors where N is the number of terminal nodes.

[Tn_Pal,Tn_Seq,I,J] = otnodes(WPT) returns the permutations of the terminal node indices such that $Tn_Seq = Tn_Pal(I)$ and $Tn_Pal = Tn_Seq(J)$.

[DP_Pal,DP_Seq] = otnodes(WPT,'dp') returns the Paley and frequency-ordered terminal nodes in node depth-position format. DP_Pal and DP_Seq are N -by-2 matrices. The first column contains the depth index, and the second column contains the position index.

Input Arguments

WPT

Binary wavelet packet tree. You can use `treeord` to determine the order of your wavelet packet tree.

dp

Character vector indicating that the Paley-ordered or sequency-ordered nodes are returned in depth-position format.

Output Arguments

Tn_Pal

Terminal nodes in Paley (natural) ordering

Tn_Seq

Terminal nodes in sequency ordering

DP_Pal

Paley-ordered terminal nodes in depth-position format. This output argument only applies when you use the 'dp' input argument.

DP_Seq

Sequency-ordered terminal nodes in depth-position format. This output argument only applies when you use the 'dp' input argument.

Examples

Order Terminal Nodes

Order terminal nodes with Paley and frequency ordering.

```
x = randn(8,1);
wpt = wpdec(x,2,'haar');
[Tn_Pal,Tn_Seq] = otnodes(wpt)
```

```
Tn_Pal = 4×1
```

```
    3
    4
    5
    6
```

```
Tn_Seq = 4×1
```

```
    3
```

```
4  
6  
5
```

Return Permutations for Ordering

Return permutations for Paley and frequency ordering.

```
load noisdopp;  
wpt = wpdec(noisdopp,6,'sym4');  
[Tn_Pal,Tn_Seq,I,J] = otnodes(wpt);  
isequal(Tn_Seq(J),Tn_Pal)
```

```
ans = logical  
1
```

```
isequal(Tn_Seq,Tn_Pal(I))
```

```
ans = logical  
1
```

Order Terminal Nodes by Depth and Position

Order terminal nodes by depth and position.

```
x = randn(8,1);  
wpt = wpdec(x,2,'haar');  
[DP_Pal,DP_Seq] = otnodes(wpt,'dp')
```

```
DP_Pal = 4x2
```

```
2    0  
2    1  
2    2  
2    3
```

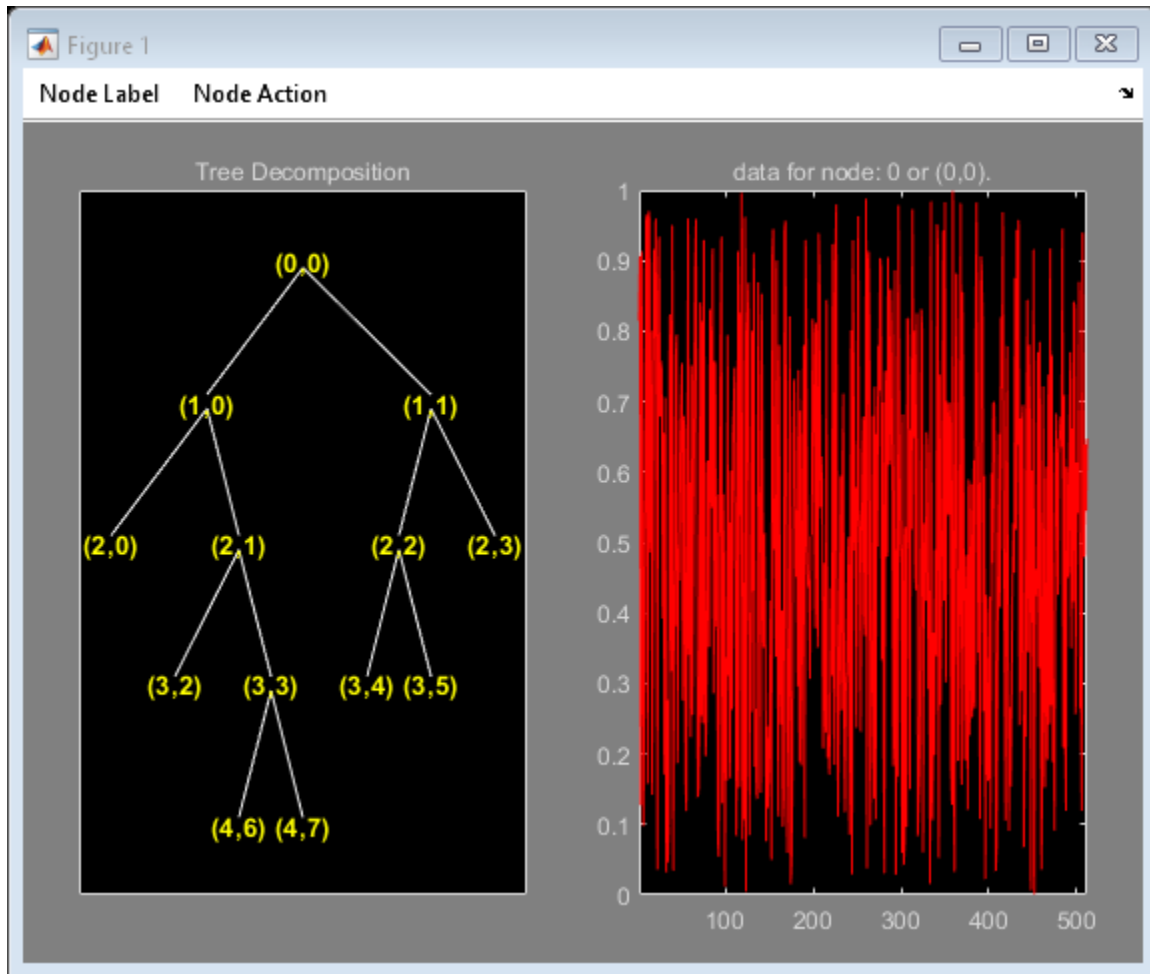
```
DP_Seq = 4x2
```

```
    2    0  
    2    1  
    2    3  
    2    2
```

Order Terminal Nodes from Wavelet Packet Tree

Order terminal nodes from a modified wavelet packet tree.

```
t = wptree(2,2,rand(1,512),'haar');  
t = wpsplt(t,4);  
t = wpsplt(t,5);  
t = wpsplt(t,10);  
plot(t);
```



```
[tn_Pal,tn_Seq,I,J] = otnodes(t)
tn_Pal = 7x1
    3
    9
   21
   22
   11
   12
```

6

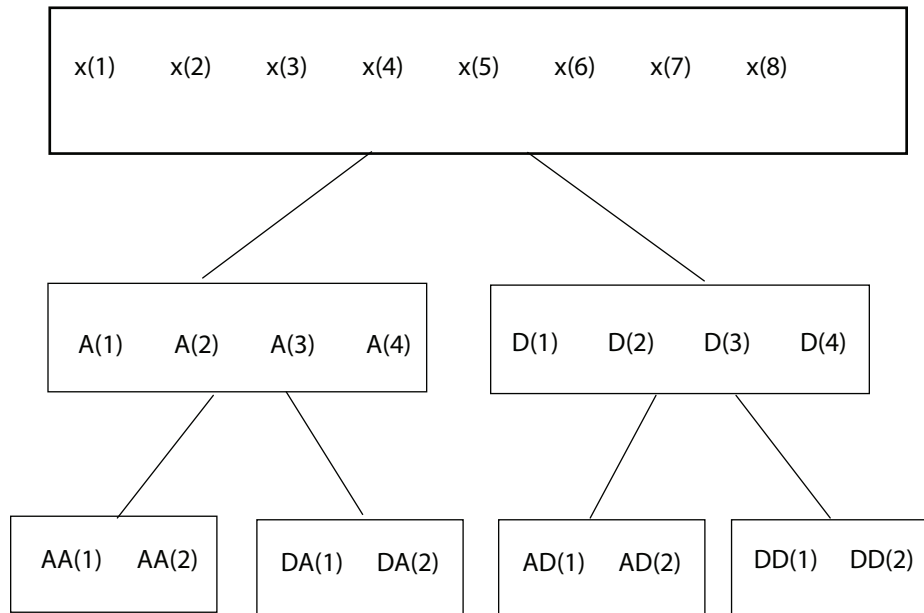
 $tn_Seq = 7 \times 1$ 3
21
22
9
6
12
11 $I = 7 \times 1$ 1
3
4
2
7
6
5 $J = 7 \times 1$ 1
4
2
3
7
6
5

Definitions

Paley (Natural) and Sequency (Frequency) Ordering

The discrete wavelet packet transform iterates on both approximation and detail coefficients at each level. In this transform, A denotes the lowpass (approximation) filter

followed by downsampling. D denotes the highpass (detail) filter followed by downsampling. The following figure represents a wavelet packet transform in Paley ordering acting on a time series of length 8. The transform has a depth of two.



Because of aliasing introduced by downsampling, the frequency content extracted by the operator AD is higher than the frequency content extracted by the DD operator. Therefore, the terminal nodes in frequency (sequency) order are: AA, DA, DD, AD . The terminal nodes in Paley order have the following indices: 3,4,5,6. The frequency order has the indices: 3,4,6,5.

References

Wickerhauser, M.V. *Lectures on Wavelet Packet Algorithms*, Technical Report, Washington University, Department of Mathematics, 1992.

See Also

leaves | treord

Introduced in R2010b

pat2cwav

Build wavelet from pattern

Syntax

```
[PSI,XVAL,NC] = pat2cwav(YPAT,METHOD,POLDEGREE,REGULARITY)
```

Description

`[PSI,XVAL,NC] = pat2cwav(YPAT,METHOD,POLDEGREE,REGULARITY)` computes an admissible wavelet for CWT (given by `XVAL` and `PSI`) adapted to the pattern defined by the vector `YPAT`, and of norm equal to 1.

The underlying x-values pattern is set to

```
xpat = linspace(0,1,length(YPAT))
```

The constant `NC` is such that `NC*PSI` approximates `YPAT` on the interval `[0,1]` by least squares fitting using

- a polynomial of degree `POLDEGREE` when `METHOD` is equal to `'polynomial'`
- a projection on the space of functions orthogonal to constants when `METHOD` is equal to `'orthconst'`

The `REGULARITY` parameter defines the boundary constraints at the points 0 and 1. Allowable values are `'continuous'`, `'differentiable'`, and `'none'`.

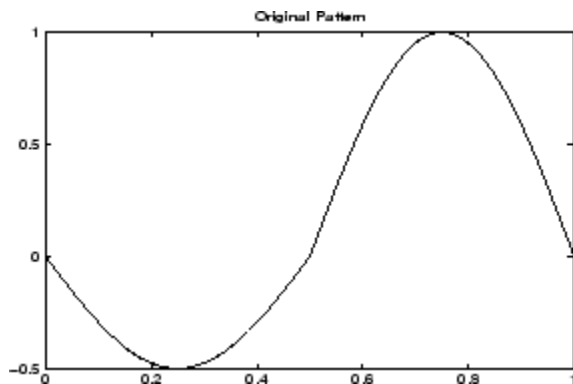
When `METHOD` is equal to `'polynomial'`

- if `REGULARITY` is equal to `'continuous'`, `POLDEGREE` must be greater than or equal to 3.
- if `REGULARITY` is equal to `'differentiable'`, `POLDEGREE` must be greater than or equal to 5.

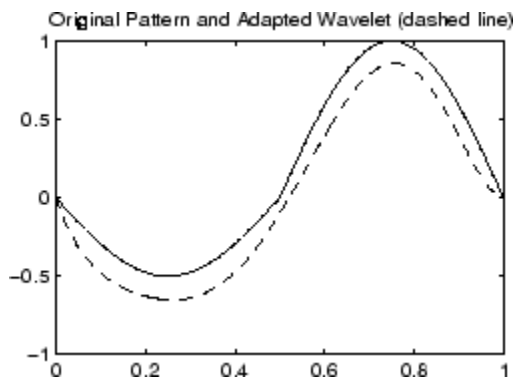
Examples

The principle for designing a new wavelet for CWT is to approximate a given pattern using least squares optimization under constraints leading to an admissible wavelet well suited for the pattern detection using the continuous wavelet transform (see Misiti et al.).

```
load ptpsin1;
plot(X,Y), title('Original Pattern')
```



```
[psi,xval,nc] = pat2cwav(Y, 'polynomial',6, 'continuous') ;
plot(X,Y,'-',xval,nc*psi,'--'),
title('Original Pattern and Adapted Wavelet (dashed line)')
```



You can check that ψ satisfies the definition of a wavelet by noting that it integrates to zero and that its L_2 norm is equal to 1.

```
dx = xval(2)-xval(1);  
Mu = sum(psi*dx)  
L2norm = sum(abs(psi).^2*dx)
```

References

Misiti, M., Y. Misiti, G. Oppenheim, J.-M. Poggi (2003), "Les ondelettes et leurs applications," Hermes.

Introduced before R2006a

paths

Scattering paths

Syntax

```
spaths = paths(sf)  
[spath, npaths] = paths(sf)
```

Description

`spath = paths(sf)` returns the scattering paths for all orders of the scattering framework, `sf`. `spath` is a cell array of MATLAB tables with n elements, where n is the number of orders in the scattering framework.

`[spath, npaths] = paths(sf)` returns the number of paths in each order as n -by-1 column vector, where n is the number of orders in the scattering framework. The sum of the elements of `npaths` is the total number of scattering paths.

Examples

Scattering Paths of Wavelet Image Scattering Framework

This example compares the number of paths in a wavelet image scattering frameworks with three orders.

Create an image scattering framework with an image size of 256-by-256 and invariance scale equal to the minimum of the image size. The default `OptimizePath` value is 1 (`true`).

```
sf = waveletScattering2('ImageSize',[256 256],'InvarianceScale',128)
```

```
sf =
```

waveletScattering2 with properties:

```
    ImageSize: [256 256]
  InvarianceScale: 128
    NumRotations: [6 6]
  QualityFactors: [1 1]
    Precision: "single"
  OversamplingFactor: 0
    OptimizePath: 1
```

Obtain the number of scattering paths in each order. Display the total number of scattering paths.

```
[spaths,npaths] = paths(sf);
sum(npaths)
```

```
ans =
```

```
    391
```

Set the `OptimizePath` value of the framework to `false`. Display the total number of scattering paths. For this framework, the scattering transform does not reduce the number of paths to compute based on a bandwidth consideration.

```
sf.OptimizePath = false;
[spaths,npaths] = paths(sf);
sum(npaths)
```

```
ans =
```

```
    571
```

Wavelets on Scattering Path

This example shows how the `OptimizePath` property can affect the scattering paths that include a specific wavelet.

Create the default wavelet image scattering framework. Obtain all the wavelet filters and center spatial frequencies for the framework. Obtain all the framework scattering paths. Display the total number of paths.

```
sf = waveletScattering2
[~,psifilters,f] = filterbank(sf);
[spaths,npaths] = paths(sf);
disp(['Total Number of Paths: ',num2str(sum(npaths))])
```

```
sf =
```

```
    waveletScattering2 with properties:
```

```
        ImageSize: [128 128]
    InvarianceScale: 64
        NumRotations: [6 6]
    QualityFactors: [1 1]
        Precision: 'single'
    OversamplingFactor: 0
        OptimizePath: 1
```

```
Total Number of Paths: 241
```

Display the number of wavelet filters in each filter bank.

```
disp(['Filter Bank 1: ',num2str(size(psifilters{1},3))]);
disp(['Filter Bank 2: ',num2str(size(psifilters{2},3))]);
```

```
Filter Bank 1: 24
Filter Bank 2: 24
```

Choose a wavelet from the first filter bank and display its spatial center frequency. Use `spaths` to find all the three-element paths that include the chosen wavelet. Display the paths.

```
waveletA = 14;
disp(['Center Frequency: ',num2str(f{1}(waveletA,:))]);
ind = find(spaths{3}.path(:,2)==waveletA);
spaths{3}(ind,:)
```

```
Center Frequency: 0.08119    0.046875
```

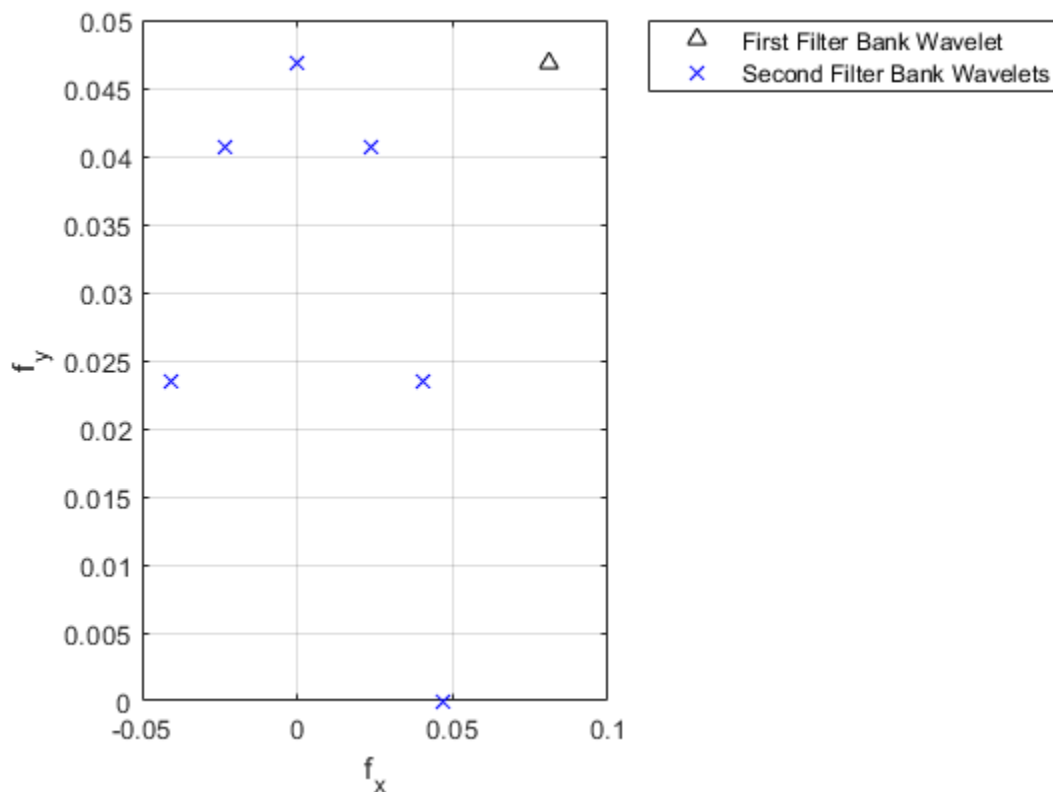
```
ans =
```

6x1 table

	path	
0	14	19
0	14	20
0	14	21
0	14	22
0	14	23
0	14	24

Plot the center frequencies of the wavelet filters on the paths.

```
plot(f{1}(waveletA,1),f{1}(waveletA,2),'k^');  
xlabel('f_x')  
ylabel('f_y')  
hold on  
waveletBs = spaths{3}.path(ind,3);  
plot(f{2}(waveletBs,1),f{2}(waveletBs,2),'bx');  
grid on  
legend('First Filter Bank Wavelet','Second Filter Bank Wavelets',...  
       'Location','northeastoutside')
```



Now set the `OptimizePath` property of the scattering framework `sf` to `false`. Obtain the wavelet filters, center spatial frequencies, and scattering paths of the framework.

```
sf.OptimizePath = false
[~,psifilters2,f2] = filterbank(sf);
[spaths2,npaths2] = paths(sf);
disp(['Total Number of Paths: ',num2str(sum(npaths2))])
```

```
sf =
```

```
waveletScattering2 with properties:
```

```
ImageSize: [128 128]
```

```
InvarianceScale: 64
  NumRotations: [6 6]
  QualityFactors: [1 1]
    Precision: 'single'
OversamplingFactor: 0
  OptimizePath: 0
```

Total Number of Paths: 385

Choose the same wavelet as above. To confirm it is the same wavelet, display its spatial center frequency. Use `spaths` to find all the three-element paths that include the wavelet. Because `OptimizePath` is set to `false`, the wavelet filter has more children.

```
waveletA = 14;
disp(['Center Frequency: ', num2str(f2{1}(waveletA,:))]);
ind = find(spaths2{3}.path(:,2)==waveletA);
spaths2{3}(ind,:)
```

```
Center Frequency: 0.08119    0.046875
```

```
ans =
```

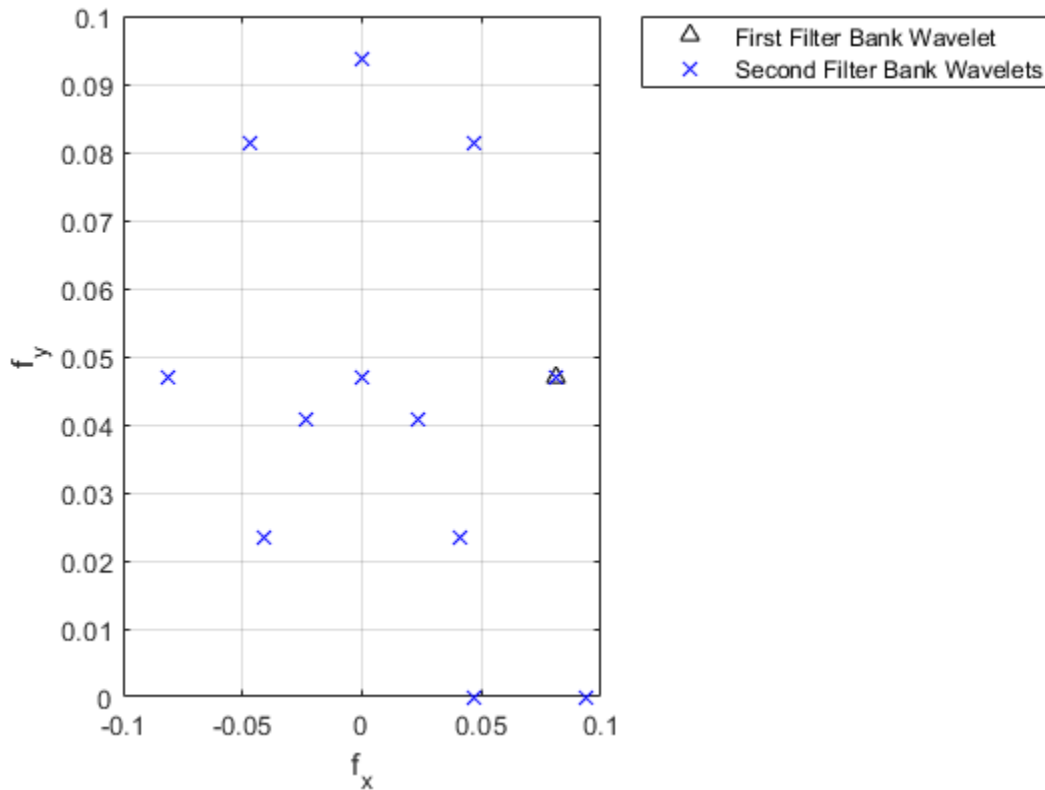
```
12x1 table
```

```
path
```

0	14	13
0	14	14
0	14	15
0	14	16
0	14	17
0	14	18
0	14	19
0	14	20
0	14	21
0	14	22
0	14	23
0	14	24

Plot the center frequencies of the wavelet filters on the paths. Some of child filters have center frequencies higher than the chosen wavelet.


```
figure
plot(f2{1}(waveletA,1),f2{1}(waveletA,2),'k^');
xlabel('f_x')
ylabel('f_y')
hold on
waveletBs = spaths2{3}.path(ind,3);
plot(f2{2}(waveletBs,1),f2{2}(waveletBs,2),'bx');
grid on
legend('First Filter Bank Wavelet','Second Filter Bank Wavelets',...
       'Location','northeastoutside')
```



Input Arguments

sf — Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

Output Arguments

spaths — Scattering paths

cell array

Scattering paths of all orders of the scattering framework, returned as a cell array of MATLAB tables. `spaths` has n elements, where n is the number of orders in the scattering framework.

Each MATLAB table in `spaths` contains a single variable, `path`. The variable `path` is a row vector with one column for each element of the path. The scalar 0 denotes the original image. Positive integers in the L th column denote the corresponding wavelet filter in the $(L-1)$ th filter bank. Wavelet bandpass filters are ordered by decreasing center frequency. There are `NumRotations` wavelets per center frequency pair.

npaths — Number of scattering paths

column vector

Number of scattering paths in each order of the scattering framework. `npaths` is a no -by-1 column vector where no is the number of orders in the scattering framework. The sum of the elements of `npaths` is the total number of scattering paths.

See Also

`coefficientSize` | `waveletScattering2`

Introduced in R2019a

plot

Plot tree GUI

Syntax

```
plot(T)  
plot(T, FIG)
```

Description

`plot` is a graphical tree-management utility.

`plot(T)` plots the tree *T*.

The figure that contains the tree is a GUI tool. It lets you change the **Node Label** to **Depth_Position** or **Index**, and **Node Action** to **Split-Merge** or **Visualize**.

The default values are **Depth_Position** and **Visualize**.

You can click the nodes to execute the current **Node Action**.

`plot(T, FIG)` plots the tree *T* in the figure whose handle is *FIG*. This figure was already used to plot a tree, for example using the command

```
FIG = plot(T)
```

After some split or merge actions, you can get the new tree using its parent figure handle. The following syntax lets you perform this functionality:

```
NEWT = plot(T, 'read', FIG)
```

In fact, the first argument is dummy. The most general syntax is

```
NEWT = plot(DUMMY, 'read', FIG)
```

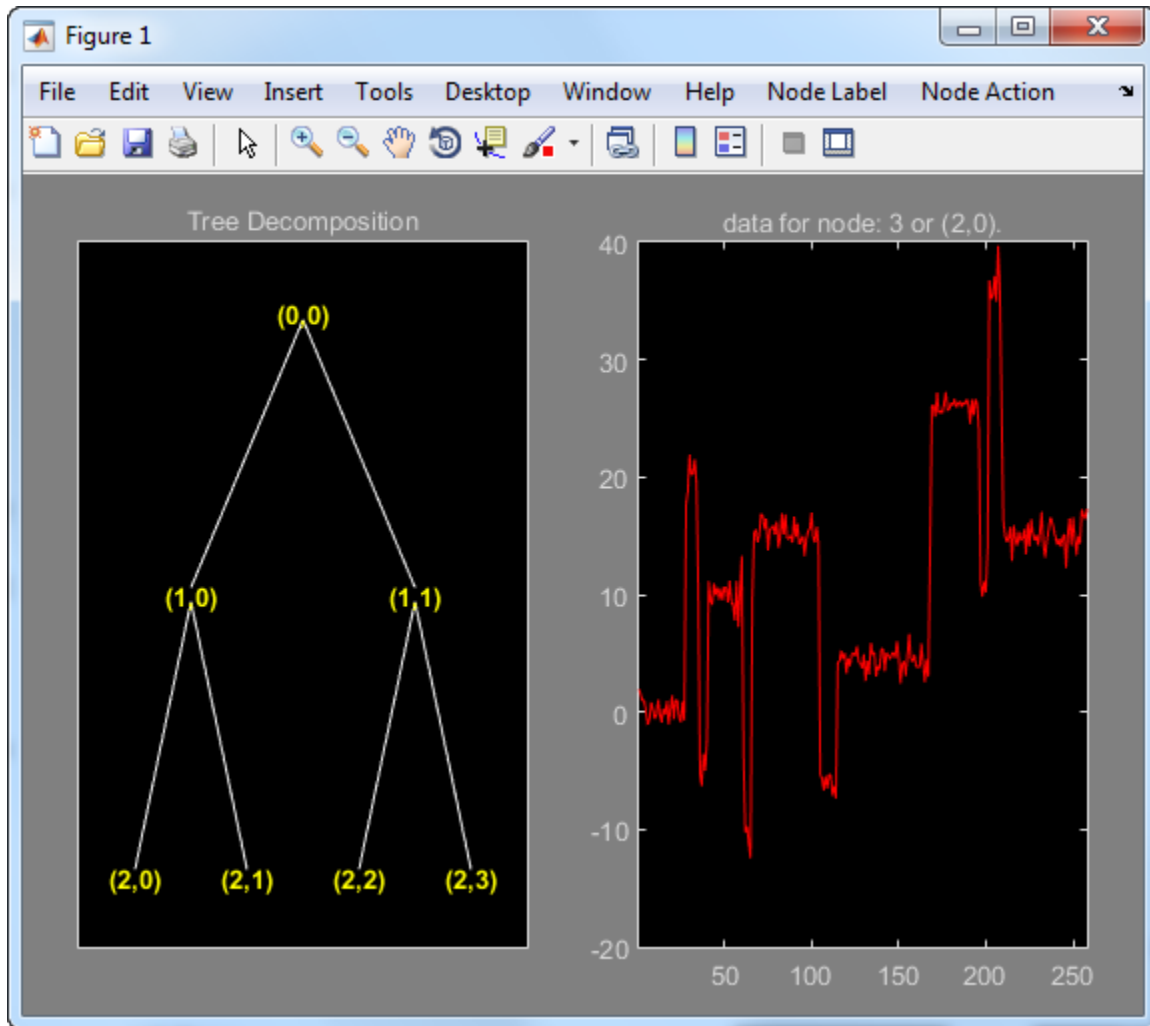
where *DUMMY* is any object parented by an NTREE object. More generally, *DUMMY* can be any object constructor name returning an NTREE parented object. For example:

```
NEWT = plot(ntree,'read',FIG)
NEWT = plot(dtree,'read',FIG)
NEWT = plot(wptree,'read',FIG)
```

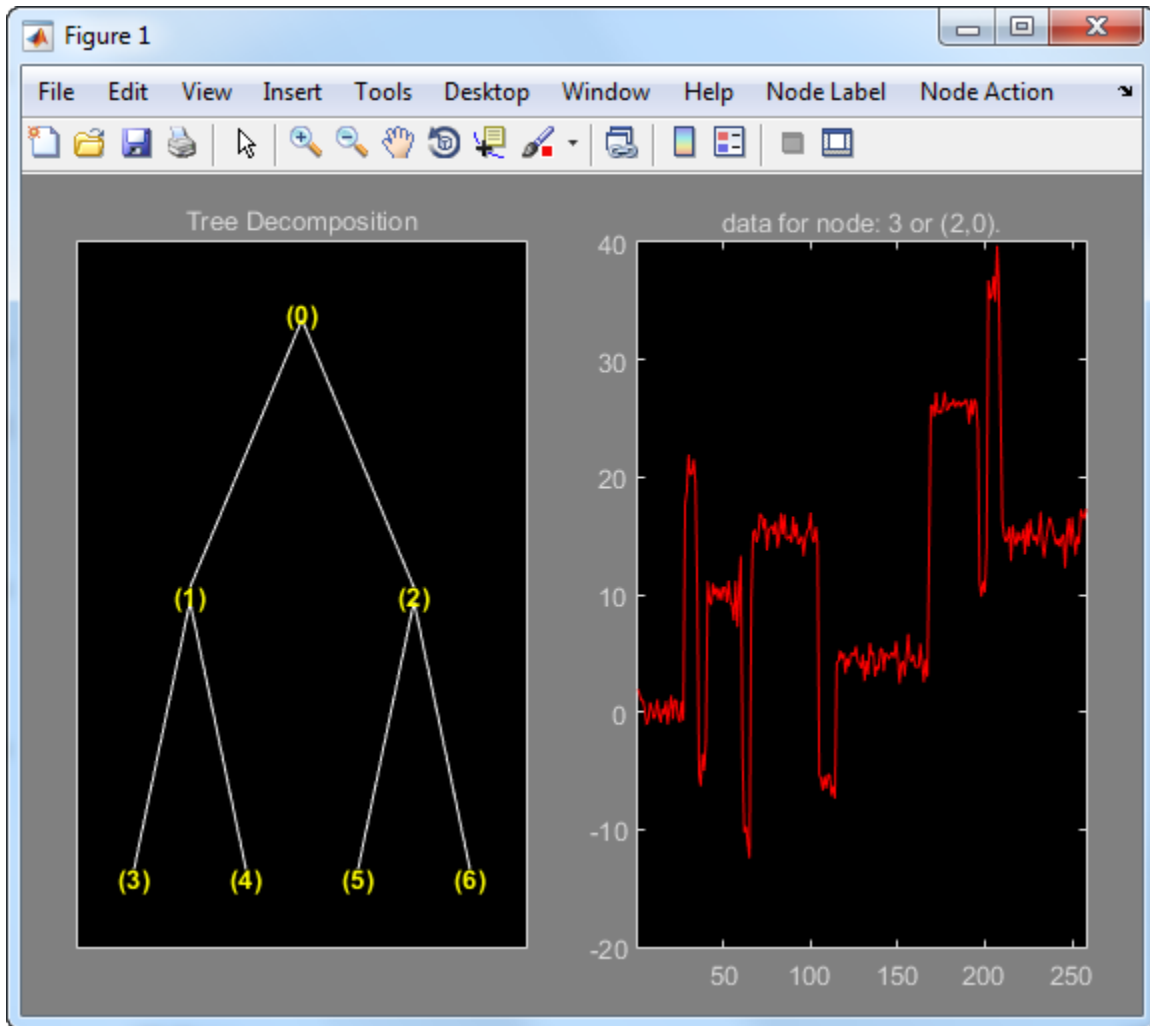
Examples

```
% Create a wavelet packets tree (1-D)
load noisbloc
x = noisbloc;
t = wpdec(x,2,'db2');

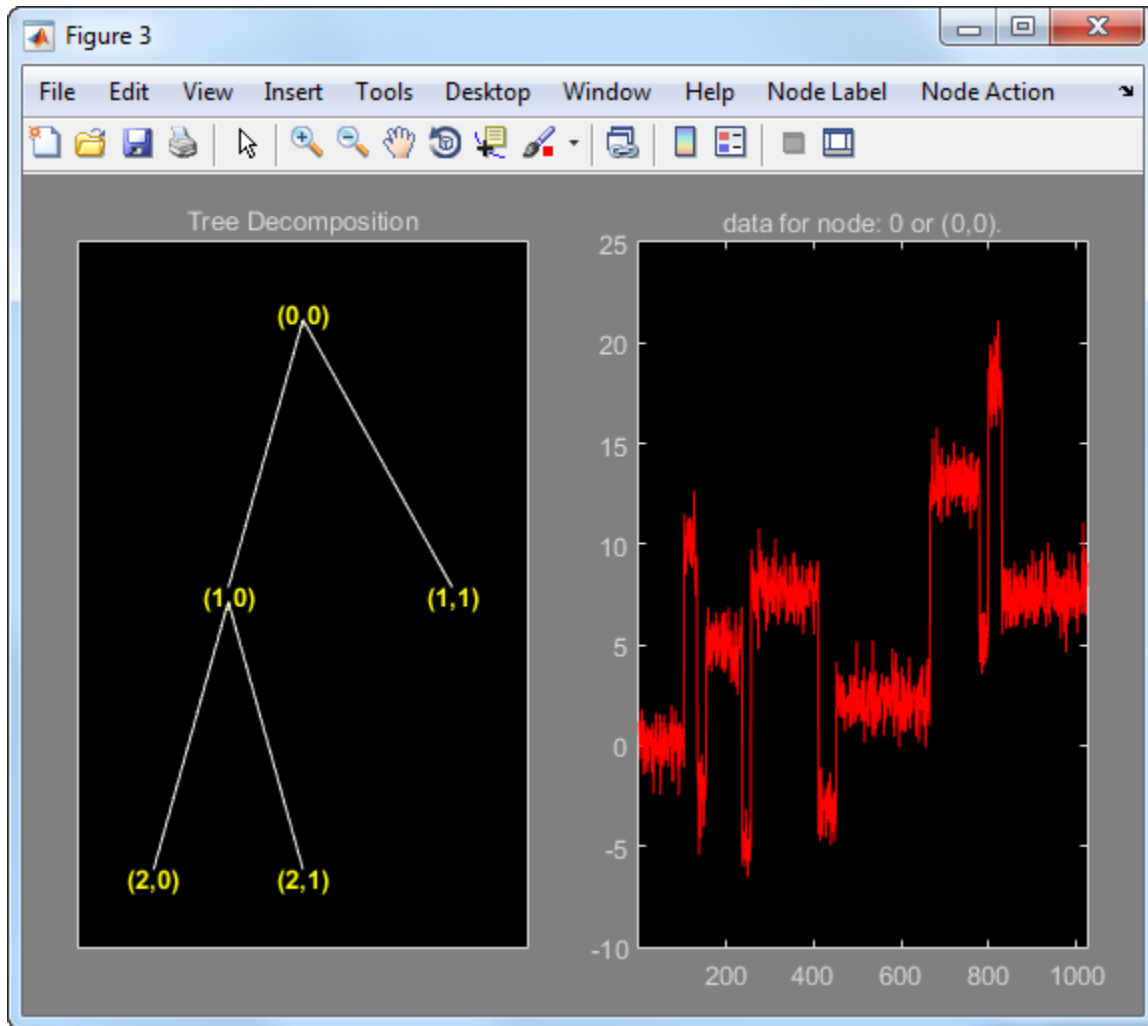
% Plot tree t.
plot(t)
```



```
% Change Node Label from Depth_Position to Index.
% Click the node (3). You get the following figure.
```



Now set the **Node Label** back to **Depth_Position**. Change **Node Action** to **Split-Merge**. Click on the (1, 1) node.



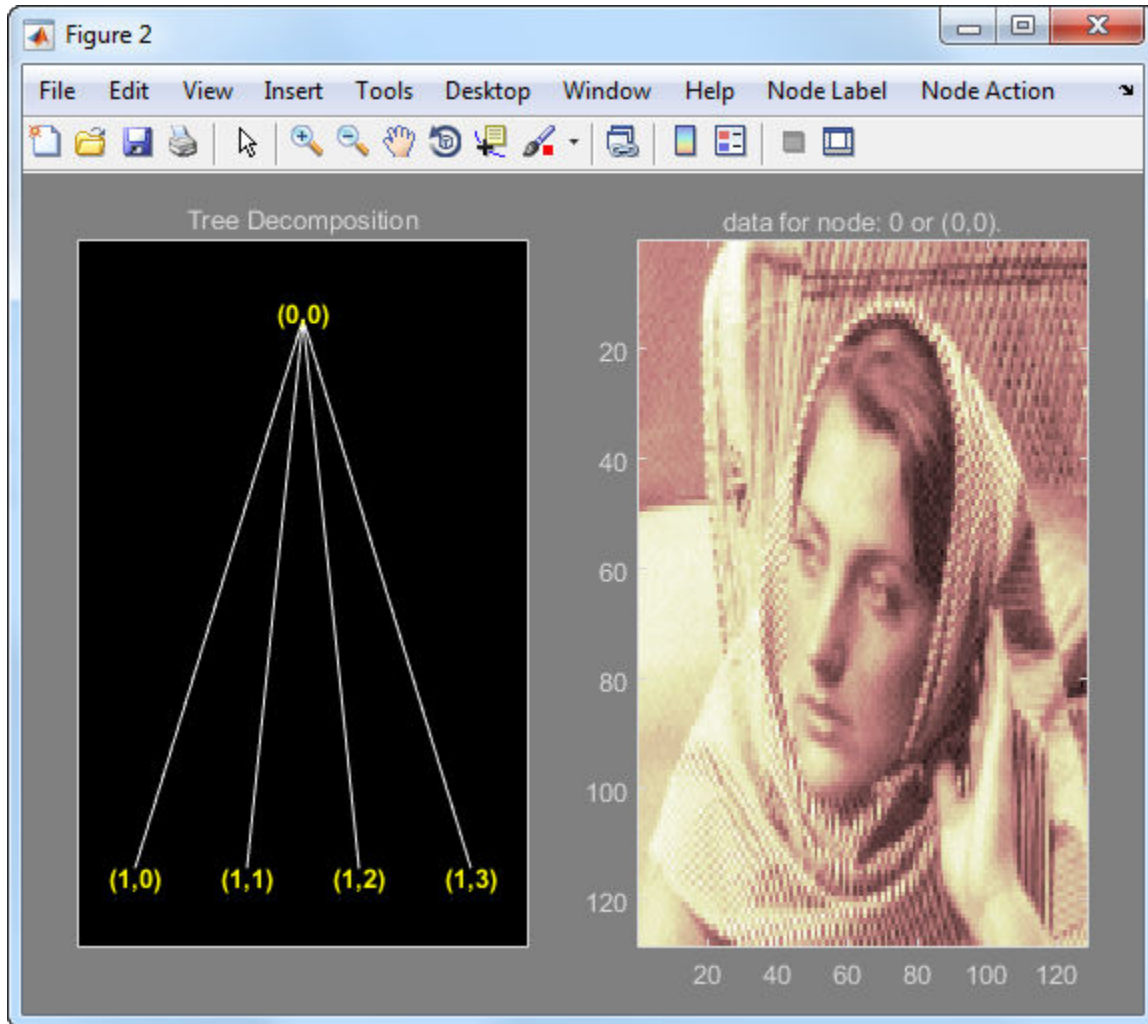
The above figure now shows the discrete wavelet transform down to level 2.

```
% Create a wavelet packets tree (2-D)
load woman2
t = wpdec2(X,1,'sym4');

% Plot tree t.
plot(t)
```



```
% Change Node Label from Depth_Position to Index.  
% Click the node (1). You get the following figure.
```



Introduced before R2006a

plotdt

Plot dual-tree or double-density wavelet transform

Syntax

```
plotdt(wt)
```

Description

`plotdt(wt)` plots the coefficients of the 1-D or 2-D wavelet filter bank decomposition, `wt`.

Examples

Plot Complex Dual-Tree Wavelet Transform of 1-D Signal

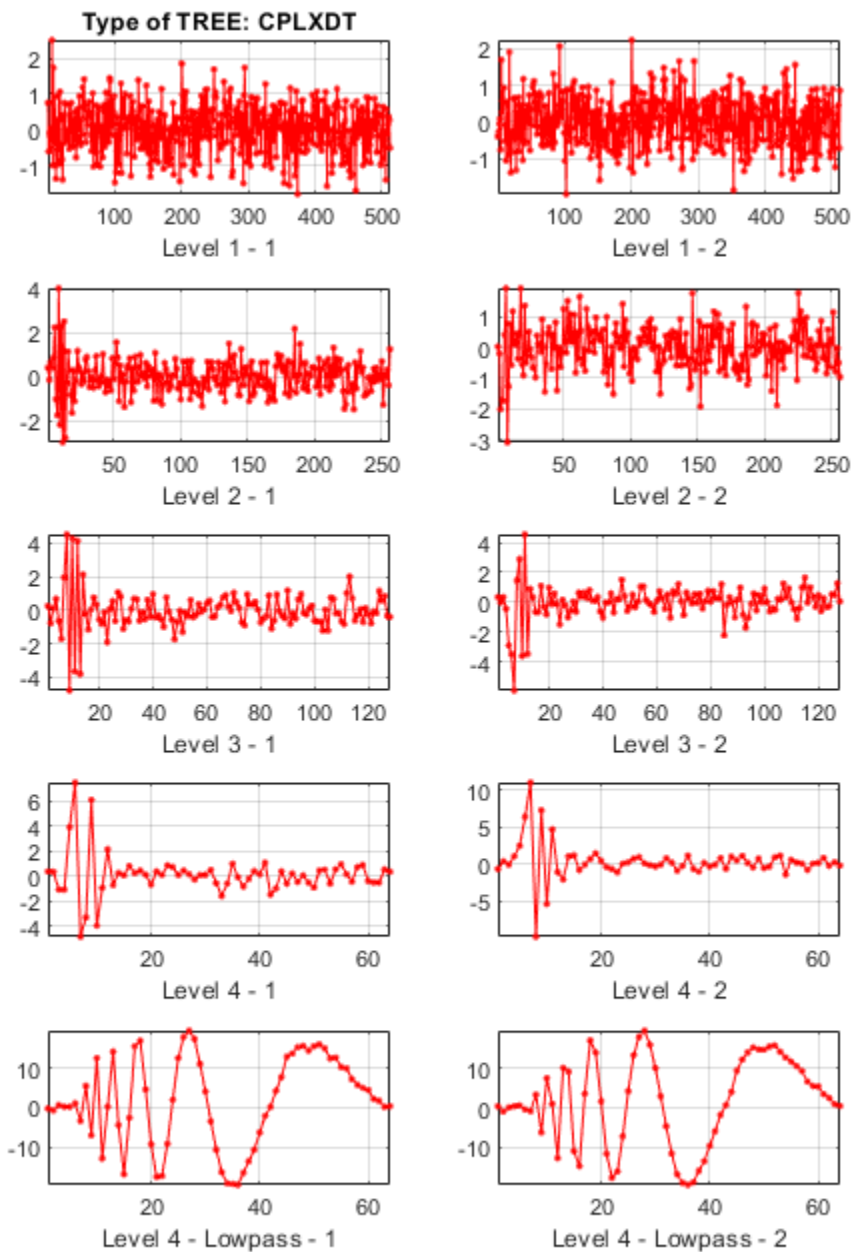
Plot the complex dual-tree wavelet transform of the noisy Doppler signal.

Load the noisy Doppler signal. Obtain the complex dual-tree wavelet transform down to level 4.

```
load noisdopp;  
wt = dddtree('cplxdt',noisdopp,4,'dtf1');
```

Plot the coefficients.

```
plotdt(wt)
```



Plot Complex Oriented Dual-Tree Wavelet Transform of 2-D Image

Plot the complex oriented dual-tree wavelet transform of an image.

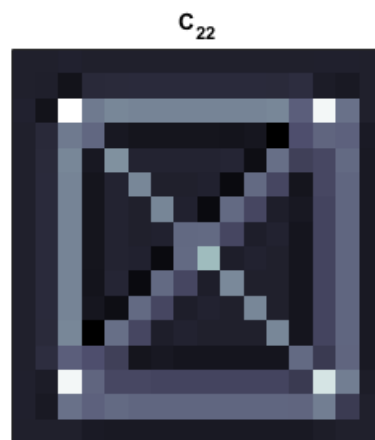
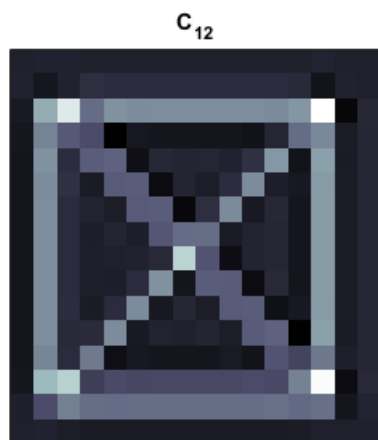
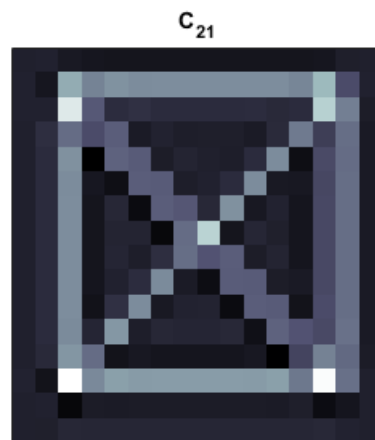
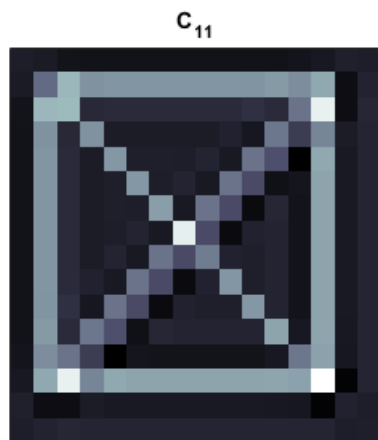
Load the `xbox` image. Obtain the complex oriented dual-tree wavelet transform down to level 3.

```
load xbox;  
wt = dddtree2('cplxdt',xbox,3,'dtf1');
```

Plot the coefficients.

```
plotdt(wt)
```

Coefficients of level 3 - Lowpass

Level 3 - Lowpass

Select the desired level detail coefficients from the drop-down list.

Input Arguments

wt — Wavelet transform

structure

Wavelet transform, returned as a structure from `dddtree` or `dddtree2` with these fields:

type — Type of wavelet decomposition (filter bank)

'dwt' | 'ddt' | 'realdt' | 'cplxdt' | 'realdddt' | 'cplxdddt'

Type of wavelet decomposition (filter bank), specified as one of 'dwt', 'ddt', 'realdt', 'cplxdt', 'realdddt', or 'cplxdddt'. 'realdt' and 'realdddt' are only valid for the 2-D wavelet transform. The type, 'dwt', is a critically sampled (nonredundant) discrete wavelet transform for 1-D data or 2-D images. The other decomposition types are oversampled wavelet transforms. For details about transform types see `dddtree` for 1-D wavelet transforms and `dddtree2` for 2-D wavelet transforms.

level — Level of the wavelet decomposition

positive integer

Level of the wavelet decomposition, specified as a positive integer.

filters — Decomposition (analysis) and reconstruction (synthesis) filters

structure

Decomposition (analysis) and reconstruction (synthesis) filters, specified as a structure with these fields:

Fdf — First-stage analysis filters

matrix | cell array

First level decomposition filters specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet

(highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage analysis filters for the corresponding tree.

Df — Analysis filters for levels > 1

matrix | cell array

Analysis filters for levels > 1, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the analysis filters for the corresponding tree.

Frf — First-level reconstruction filters

matrix | cell array

First-level reconstruction filters, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

Rf — Reconstruction filters for levels > 1

matrix | cell array

Reconstruction filters for levels > 1, specified as an N -by-2 or N -by-3 matrix for single-tree wavelet transforms, or a 1-by-2 cell array of two N -by-2 or N -by-3 matrices for dual-tree wavelet transforms. The matrices are N -by-3 for the double-density wavelet transforms. For an N -by-2 matrix, the first column of the matrix is the scaling (lowpass) filter and the second column is the wavelet (highpass) filter. For an N -by-3 matrix, the first column of the matrix is the scaling (lowpass) filter and the second and third columns are the wavelet (highpass) filters. For the dual-tree transforms, each element of the cell array contains the first-stage synthesis filters for the corresponding tree.

cfs — Wavelet transform coefficients

cell array of matrices

Wavelet transform coefficients, specified as a 1-by-(level+1) cell array of matrices. The size and structure of the matrix elements of the cell array depend on the type of wavelet transform and whether the decomposition is 1-D or 2-D. For a 1-D wavelet transform, the coefficients are organized by transform type as follows:

- 'dwt' — cfs{j}
 - j = 1,2,... level is the level.
 - cfs{level+1} are the lowpass, or scaling, coefficients.
- 'ddt' — cfs{j}(:, :, k)
 - j = 1,2,... level is the level.
 - k = 1,2 is the wavelet filter.
 - cfs{level+1}(:, :,) are the lowpass, or scaling, coefficients.
- 'cplxdt' — cfs{j}(:, :, m)
 - j = 1,2,... level is the level.
 - m = 1,2 are the real and imaginary parts.
 - cfs{level+1}(:, :,) are the lowpass, or scaling, coefficients.
- 'realdddt' — cfs{j}(:, :, d, k)
 - j = 1,2,... level is the level.
 - d = 1,2,3 is the orientation.
 - k = 1,2 is the wavelet transform tree.
 - cfs{level+1}(:, :,) are the lowpass, or scaling, coefficients.
- 'cplxdddt' — cfs{j}(:, :, d, k, m)
 - j = 1,2,... level is the level.
 - k = 1,2 is the wavelet transform tree.
 - m = 1,2 are the real and imaginary parts.
 - cfs{level+1}(:, :,) are the lowpass, or scaling, coefficients.

For a 2-D wavelet transform, the coefficients are organized by transform type as follows:

- 'dwt' — cfs{j}(:, :, d)
 - j = 1,2,... level is the level.

- $d = 1,2,3$ is the orientation.
- $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'ddt' — $cfs\{j\}(:, :, d)$
 - $j = 1,2,\dots$ level is the level.
 - $d = 1,2,3,4,5,6,7,8$ is the orientation.
 - $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'realddt' — $cfs\{j\}(:, :, d, k)$
 - $j = 1,2,\dots$ level is the level.
 - $d = 1,2,3$ is the orientation.
 - $k = 1,2$ is the wavelet transform tree.
 - $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1,2,\dots$ level is the level.
 - $d = 1,2,3$ is the orientation.
 - $k = 1,2$ is the wavelet transform tree.
 - $m = 1,2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'realdddtdt' — $cfs\{j\}(:, :, d, k)$
 - $j = 1,2,\dots$ level is the level.
 - $d = 1,2,3$ is the orientation.
 - $k = 1,2$ is the wavelet transform tree.
 - $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.
- 'cplxdddtdt' — $cfs\{j\}(:, :, d, k, m)$
 - $j = 1,2,\dots$ level is the level.
 - $d = 1,2,3$ is the orientation.
 - $k = 1,2$ is the wavelet transform tree.
 - $m = 1,2$ are the real and imaginary parts.
 - $cfs\{\text{level}+1\}(:, :)$ are the lowpass, or scaling, coefficients.

See Also

dddtree | dddtree2 | dddtreecfs

Topics

“Analytic Wavelets Using the Dual-Tree Wavelet Transform”

“Critically Sampled and Oversampled Wavelet Filter Banks”

Introduced in R2013b

powerbw

CWT filter bank 3-dB bandwidths

Syntax

```
bw = powerbw(fb)
```

Description

`bw = powerbw(fb)` returns 3-dB (half-power) bandwidths for the wavelet filters in the filter bank `fb`. `bw` is a `Ns`-by-4 MATLAB table, where `Ns` is the number of wavelet bandpass frequencies (equal to the number of scales). For every filter in `fb`, the table contains the corresponding bandpass frequency, the 3-dB bandwidth, and the lower frequency and upper frequency limits of the 3-dB bandwidth.

The 3-dB bandwidth limits mark where the filter power is half its peak value. The magnitude frequency response at the limits is equal to $1/\sqrt{2}$ times the peak magnitude. Since the passbands in `fb` are normalized with peak magnitudes approximately equal to 2, the magnitude frequency response at each limit is approximately equal to $2/\sqrt{2}$. The 3-dB bandwidth is also known as the half-power bandwidth because $20 \log_{10} 1/\sqrt{2} \approx -3$.

Examples

Half-Power Wavelet Bandwidths

Create a CWT filter bank.

```
fb = cwtfilterbank;
```

Obtain the 3-dB (half-power) bandwidths of the filter bank. Obtain the frequency responses of the wavelets.

```
bw = powerbw(fb);  
[psidft,f] = freqz(fb);
```

Inspect the first few rows in the table `bw`.

```
bw(1:10, :)
```

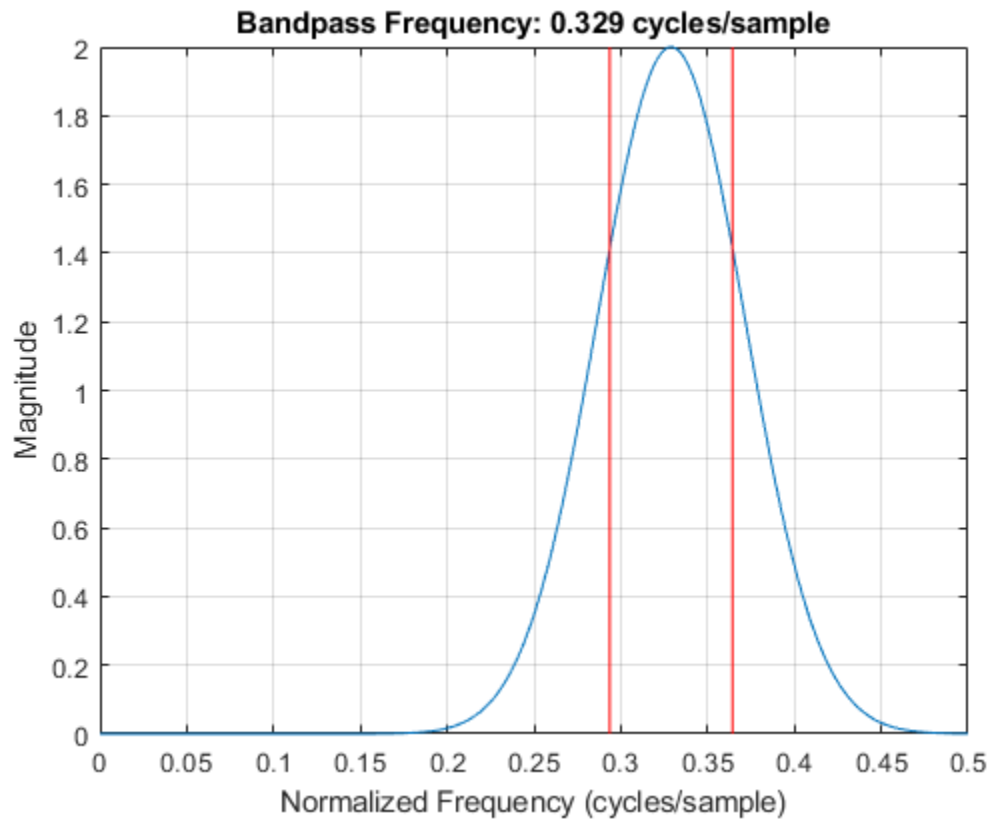
```
ans=10x4 table
    Frequencies    HalfPowerBandwidth    LowFrequencyBorder    HighFrequencyBorder
    _____    _____    _____    _____
    0.43412         0.09323         0.38751         0.48074
    0.40505         0.086988        0.36156         0.44855
    0.37793         0.081162        0.33735         0.41851
    0.35262         0.075726        0.31476         0.39048
    0.329           0.070655        0.29368         0.36434
    0.30697         0.065924        0.27401         0.33994
    0.28642         0.061509        0.25566         0.31717
    0.26723         0.05739         0.23854         0.29593
    0.24934         0.053547        0.22257         0.27611
    0.23264         0.049961        0.20766         0.25762
```

Choose a wavelet bandpass filter from the filter bank. Extract from the table `bw` the 3-dB limits of the bandpass filter.

```
wv = 5;
frq = bw.Frequencies(wv);
lfb = bw.LowFrequencyBorder(wv);
hfb = bw.HighFrequencyBorder(wv);
```

Plot the frequency response and 3-dB limits. Since the frequency response is scaled to have a maximum value equal to 2, inspect the plot to confirm the lower and upper frequency borders cross intersect the frequency response at $\sqrt{2}$.

```
plot(f,psidft(wv,:))
grid on
hold on
plot([lfb lfb],[0 2],'r')
plot([hfb hfb],[0 2],'r')
xlabel('Normalized Frequency (cycles/sample)')
ylabel('Magnitude')
title(['Bandpass Frequency: ' num2str(frq) ' cycles/sample'])
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

bw — 3-dB (half-power) bandwidths

table

3-dB (half-power) bandwidths, returned as a N_s -by-4 table, where N_s is the number of wavelet bandpass frequencies (equal to the number of scales). The table has the following four variables:

Frequencies — Bandpass frequency

positive scalar

Bandpass frequency, returned as a positive scalar (see `centerFrequencies`).

Data Types: double

HalfPowerBandwidth — Half-power bandwidth

positive scalar

Half-power bandwidth, returned as a positive scalar.

Data Types: double

LowFrequencyBorder — Lower frequency edge

positive scalar

Lower frequency edge of the 3-dB bandwidth, returned as a positive scalar.

Data Types: double

HighFrequencyBorder — High frequency edge

positive scalar

High frequency edge of the 3-dB bandwidth, returned as a positive scalar.

Data Types: double

Data Types: table

See Also

`centerFrequencies` | `cwtfilterbank` | `freqz`

Introduced in R2018a

powerbw

DWT filter bank power bandwidth

Syntax

```
bwtable = powerbw(fb)
```

Description

`bwtable = powerbw(fb)` returns a MATLAB table `bwtable` containing the following variables by level:

- Theoretical DWT frequency bands,
- Measured wavelet and scaling filter 3-dB bandwidths,
- Proportions of the total energy in the reported bands.

Examples

DWT Filter Bank Power Bandwidth

Obtain the 3-dB bandwidths of a level-4 discrete wavelet transform with the Fejér-Korovkin `fk18` wavelet. Compare the theoretical and 3-dB bandwidths of the wavelet and scaling filters.

```
fb = dwtfilterbank('Wavelet','fk18','Level',4);  
bw = powerbw(fb);  
bw(:,1:4)
```

```
ans=4x4 table
```

Level	DWTBand		Wavelet3dBBandwidth		Scaling3dBBandwidth	
1	0.25	0.5	0.25	0.5	0	0.25
2	0.125	0.25	0.12496	0.2501	0	0.12499

3	0.0625	0.125	0.062479	0.12504	0	0.062495
4	0.03125	0.0625	0.03124	0.062521	0	0.031247

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

bwtable — Theoretical and measured bandwidths

table

Theoretical and measured bandwidths of the DWT filter bank `fb`, returned as a MATLAB table. `bwtable` is L-by-8 where L is the filter bank `Levels`. Levels are ordered by decreasing resolution. `bwtable` has the following eight variables:

Level — Level of DWT decomposition

positive integer

Level of DWT decomposition, returned as a positive integer less than or equal to the filter bank `Levels`.

DWTBand — Theoretical DWT frequency bands

two-element real-valued vector

Theoretical DWT frequency bands by level, returned as a two-element real-valued vector.

Wavelet3dBBandwidth — Measured wavelet 3-dB bandwidths

two-element real-valued vector

Measured wavelet 3-dB bandwidths by level, returned as a two-element real-valued vector.

Scaling3dBBandwidth — Measured scaling filter 3-dB bandwidths

two-element real-valued vector

Measured scaling filter 3-dB bandwidths by level, returned as a two-element real-valued vector.

WaveletPowerIn3dBBand — Proportion of total wavelet power

positive scalar

Proportion of total wavelet power in the measured 3-dB band by level, returned as a positive scalar.

ScalingPowerIn3dBBand — Proportion of total scaling filter power

positive scalar

Proportion of total scaling filter power in the measured 3-dB band by level, returned as a positive scalar.

WaveletPowerInDWTBand — Proportion of total wavelet power

positive scalar

Proportion of total wavelet power in the theoretical DWT band by level, returned as a positive scalar.

ScalingPowerInDWTBand — Proportion of total scaling filter power

positive scalar

Proportion of total scaling filter power in the theoretical DWT band by level, returned as a positive scalar.

See Also

dwtfilterbank

Introduced in R2018a

qfactor

CWT filter bank quality factor

Syntax

```
qf = qfactor(fb)
```

Description

`qf = qfactor(fb)` returns the quality factor for the wavelet bandpass filters in `fb`. The quality factor is the ratio of the 3-dB bandwidth to the center frequency, where the center frequency is the geometric mean of the bandwidth frequencies. The larger the quality factor, the more frequency localized the wavelet. For reference, a half-band filter has a quality factor of `sqrt(2)`.

Examples

Quality Factor of CWT Filter Bank

Create a CWT filter bank using the default analytic Morse (3,60) wavelet.

```
fb = cwtfilterbank;
```

Compute the quality factor of the filter bank.

```
qf = qfactor(fb)
```

```
qf = 4.6296
```

Create a CWT filter bank using the analytic Morse (3,10) wavelet. Compute the quality factor of the filter bank. The analytic Morlet (3,10) wavelet is not localized in frequency as well as the Morse (3,60) wavelet. Confirm that the quality factor of the second filter bank is smaller than the first filter bank.

```
fb2 = cwtfilterbank('Timebandwidth',10);  
qf2 = qfactor(fb2)  
  
qf2 = 1.8445
```

Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

qf — Quality factor
positive number

Quality factor, returned as a positive real number.

Data Types: `double`

See Also

`cwtfilterbank` | `powerbw`

Introduced in R2018a

qfactor

DWT filter bank quality factor

Syntax

```
qf = qfactor(fb)
```

Description

`qf = qfactor(fb)` returns the quality factor for the wavelet filter bank, `fb`. The quality factor, `qf`, is defined to be the geometric mean frequency of the lower and upper 3-dB bandwidth frequencies divided by the 3-dB bandwidth. For orthogonal wavelets, the measured quality factor approximates the theoretical value of $\sqrt{2}$.

Examples

DWT Filter Bank Quality Factor

Obtain the quality factor for the orthogonal Coiflet `coif4`.

```
wvOrth = 'coif4';  
fb = dwtfilterbank('Wavelet',wvOrth);  
orthogAnalysis = qfactor(fb)
```

```
orthogAnalysis = 1.4142
```

Compare with the quality factor for the biorthogonal wavelet `bior6.8`.

```
wvBior = 'bior6.8';  
fb2 = dwtfilterbank('Wavelet',wvBior);  
biorthogAnalysis = qfactor(fb2)
```

```
biorthogAnalysis = 1.5481
```

By default, `fb` and `fb2` filter banks have the default filter type `Analysis`. Create two new filter banks of filter type `Synthesis` for the same wavelets. Compare the quality factors

with the filter type `Analysis` filter banks. The quality factors using the orthogonal wavelet are equal, while the quality factors using the biorthogonal wavelet are different.

```
fb3 = dwtfilterbank('Wavelet',wvOrth,'FilterType','Synthesis');  
fb4 = dwtfilterbank('Wavelet',wvBior,'FilterType','Synthesis');  
orthogSynthesis = qfactor(fb3)
```

```
orthogSynthesis = 1.4142
```

```
biorthogSynthesis = qfactor(fb4)
```

```
biorthogSynthesis = 1.3002
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

qf — Quality factor

positive scalar

Quality factor of the DWT filter bank `fb`, returned as a positive scalar.

See Also

`dwtfilterbank`

Introduced in R2018a

qmf

Scaling and Wavelet Filter

Syntax

$Y = \text{qmf}(X, P)$

$Y = \text{qmf}(X)$

$Y = \text{qmf}(X, \theta)$

Description

$Y = \text{qmf}(X, P)$ changes the signs of the even index elements of the reversed vector filter coefficients X if P is θ . If P is 1 , the signs of the odd index elements are reversed. Changing P changes the phase of the Fourier transform of the resulting wavelet filter by π radians.

$Y = \text{qmf}(X)$ is equivalent to $Y = \text{qmf}(X, \theta)$.

Let x be a finite energy signal. Two filters F_0 and F_1 are quadrature mirror filters (QMF) if, for any x ,

$$\|y_0\|^2 + \|y_1\|^2 = \|x\|^2$$

where y_0 is a decimated version of the signal x filtered with F_0 so y_0 defined by $x_0 = F_0(x)$ and $y_0(n) = x_0(2n)$, and similarly, y_1 is defined by $x_1 = F_1(x)$ and $y_1(n) = x_1(2n)$. This property ensures a perfect reconstruction of the associated two-channel filter banks scheme (see Strang-Nguyen p. 103).

For example, if F_0 is a Daubechies scaling filter with norm equal to 1 and $F_1 = \text{qmf}(F_0)$, then the transfer functions $F_0(z)$ and $F_1(z)$ of the filters F_0 and F_1 satisfy the condition (see the example for db10):

$$|F_0(z)|^2 + |F_1(z)|^2 = 2.$$

Examples

Create a Quadrature Mirror Filter

This example shows how to create a quadrature mirror filter associated with the `db10` wavelet.

Compute the scaling filter associated with the `db10` wavelet.

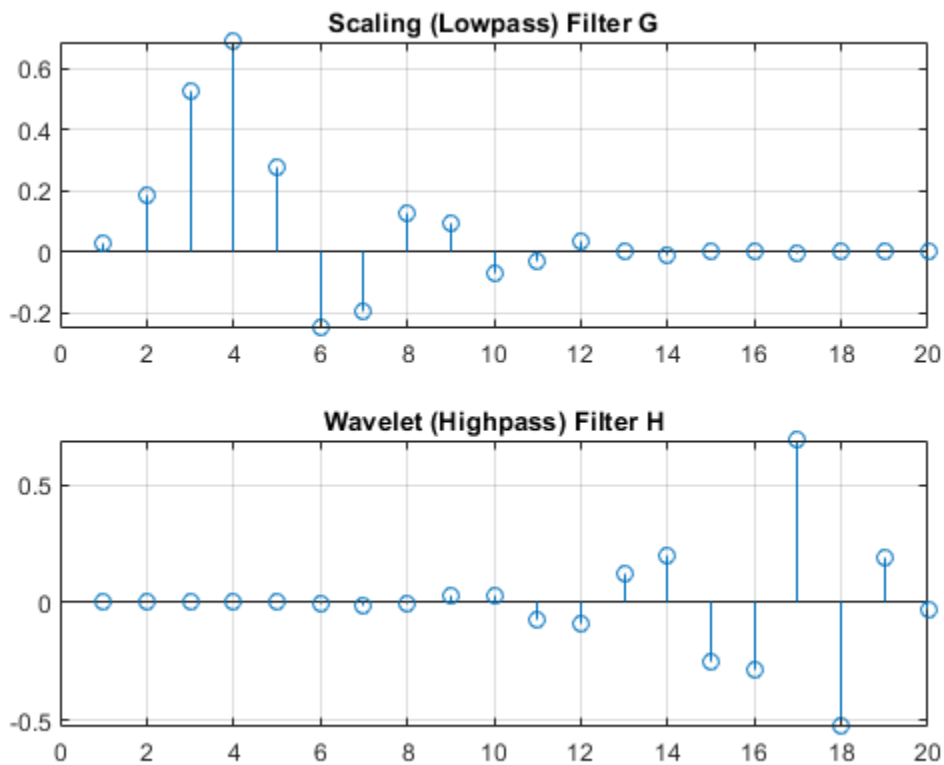
```
sF = dbwavf('db10');
```

`dbwavf` normalizes the filter coefficients so that the norm is equal to $1/\sqrt{2}$. Normalize the coefficients so that the filter has norm equal to 1.

```
G = sqrt(2)*sF;
```

Obtain the wavelet filter coefficients by using `qmf`. Plot the filters.

```
H = qmf(G);  
subplot(2,1,1)  
stem(G)  
title('Scaling (Lowpass) Filter G')  
grid on  
subplot(2,1,2)  
stem(H)  
title('Wavelet (Highpass) Filter H')  
grid on
```

Set the DWT extension mode to Periodization. Generate a random signal of length 64. Perform a single-level wavelet decomposition of the signal using G and H.

```
dwtmode('per')
```

```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
! WARNING: Change DWT Extension Mode !
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
*****
** DWT Extension Mode: Periodization **
*****
```

```
n = 64;
rng 'default'
sig = randn(1,n);
[a,d] = dwt(sig,G,H);
```

The lengths of the approximation and detail coefficients are both 32. Confirm that the filters preserve energy.

```
[sum(sig.^2) sum(a.^2)+sum(d.^2)]
```

```
ans = 1×2
```

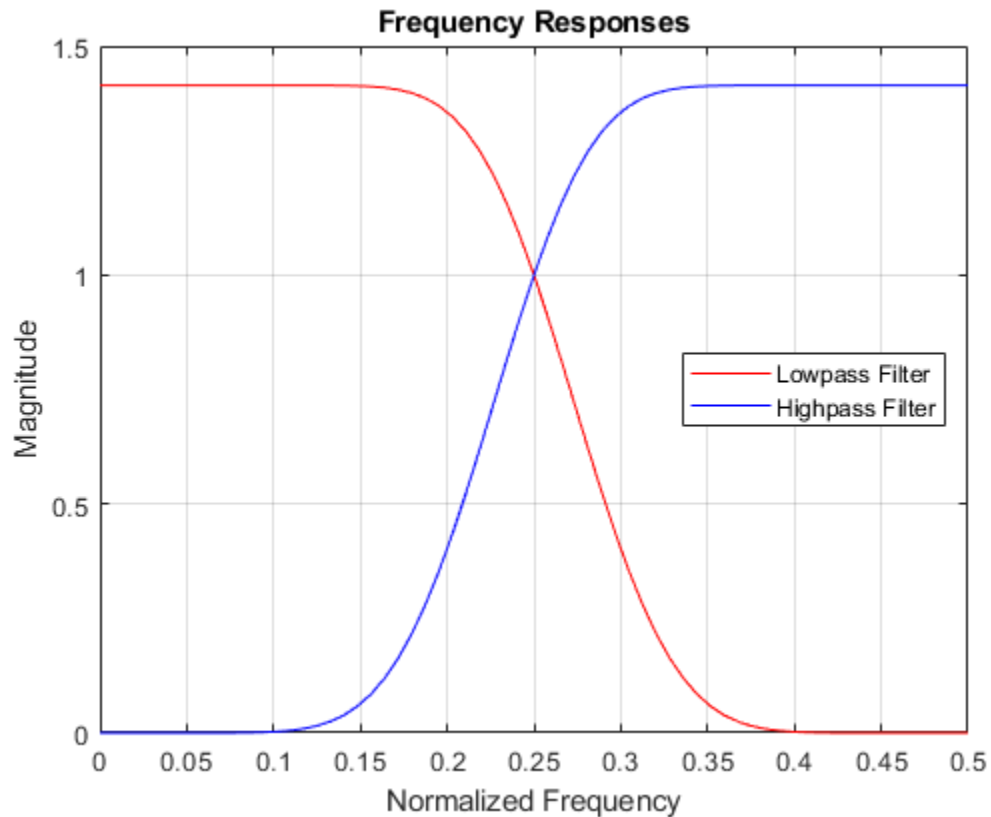
```
92.6872 92.6872
```

Compute the frequency responses of G and H. Zeropad the filters when taking the Fourier transform.

```
n = 128;
F = 0:1/n:1-1/n;
Gdft = fft(G,n);
Hdft = fft(H,n);
```

Plot the magnitude of each frequency response.

```
figure
plot(F(1:n/2+1),abs(Gdft(1:n/2+1)), 'r')
hold on
plot(F(1:n/2+1),abs(Hdft(1:n/2+1)), 'b')
grid on
title('Frequency Responses')
xlabel('Normalized Frequency')
ylabel('Magnitude')
legend('Lowpass Filter', 'Highpass Filter', 'Location', 'east')
```



Confirm the sum of the squared magnitudes of the frequency responses of G and H at each frequency is equal to 2.

```
sumMagnitudes = abs(Gdft).^2+abs(Hdft).^2;
[min(sumMagnitudes) max(sumMagnitudes)]
```

```
ans = 1x2
```

```
2.0000    2.0000
```

Confirm that the filters are orthonormal.

```
df = [G;H];
id = df*df'
```

```
id = 2×2  
  
    1.0000    0.0000  
    0.0000    1.0000
```

Controlling the Phase of a Quadrature Mirror Filter

This example shows the effect of setting the phase parameter of the `qmf` function.

Obtain the decomposition low-pass filter associated with a Daubechies wavelet.

```
lowfilt = wfilters('db4');
```

Use the `qmf` function to obtain the decomposition low-pass filter for a wavelet. Then, compare the signs of the values when the `qmf` phase parameter is set to 0 or 1. The reversed signs indicates a phase shift of π radians, which is the same as multiplying the DFT by $e^{i\pi}$.

```
p0 = qmf(lowfilt,0)
```

```
p0 = 1×8
```

```
    0.2304   -0.7148    0.6309    0.0280   -0.1870   -0.0308    0.0329    0.0106
```

```
p1 = qmf(lowfilt,1)
```

```
p1 = 1×8
```

```
   -0.2304    0.7148   -0.6309   -0.0280    0.1870    0.0308   -0.0329   -0.0106
```

Compute the magnitudes and display the difference between them. Unlike the phase, the magnitude is not affected by the sign reversals.

```
abs(p0) - abs(p1)
```

```
ans = 1×8
```

```
    0    0    0    0    0    0    0    0
```

References

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Introduced before R2006a

rbiowavf

Reverse biorthogonal spline wavelet filters

Syntax

```
[RF,DF] = rbiowavf(wname)
```

Description

[RF,DF] = rbiowavf(wname) returns the reconstruction (synthesis) and decomposition (analysis) scaling filters, RF and DF, respectively, associated with the reverse biorthogonal wavelet specified by wname.

Examples

Reverse Biorthogonal Scaling Filter

Obtain the reverse biorthogonal reconstruction and decomposition scaling filters for the 'rbio3.1' wavelet. The 'rbio3.1' wavelet has three vanishing moments for the decomposition (analysis) wavelet and one vanishing moment for the reconstruction (synthesis) wavelet.

```
[RF,DF] = rbiowavf('rbio3.1');
```

The reconstruction scaling filter, RF, and the decomposition filter, DF, are equal to the filters returned by wfilters scaled by $\sqrt{2}$.

```
[LoD,HiD,LoR,HiR] = wfilters('rbio3.1');  
max(abs(sqrt(2)*DF-LoD))
```

```
ans = 0
```

```
max(abs(sqrt(2)*RF-LoR))
```

```
ans = 0
```

Input Arguments

wname — Name of reverse biorthogonal wavelet

character vector | string scalar

Name of reverse biorthogonal wavelet, specified as 'rbioNd.Nr' where possible values for Nd and Nr are as follows:

Nd = 1	Nr = 1 , 3 or 5
Nd = 2	Nr = 2 , 4 , 6 or 8
Nd = 3	Nr = 1 , 3 , 5 , 7 or 9
Nd = 4	Nr = 4
Nd = 5	Nr = 5
Nd = 6	Nr = 8

Nd and Nr are the numbers of vanishing moments for the decomposition and reconstruction filters, respectively.

Example: 'rbiowavf3.7'

Output Arguments

RF — Reconstruction filter

real-valued vector

Reconstruction filter associated with the reverse biorthogonal wavelet wname, returned as a real-valued vector.

DF — Decomposition filter

real-valued vector

Decomposition filter associated with the reverse biorthogonal wavelet wname, returned as a real-valued vector.

See Also

`biorfilt` | `waveinfo`

Introduced before R2006a

read

Read values of WPTREE

Syntax

VARARGOUT = read(T,VARARGIN)

Description

VARARGOUT = read(T,VARARGIN) is the most general syntax to read one or more property values from the fields of a WPTREE object .

The different ways to call the read function are

PropValue = read(T,'PropName') or
PropValue = read(T,'PropName','PropParam')

or any combination of the previous syntaxes:

[PropValue1,PropValue2,] =
read(T,'PropName1','PropParam1','PropName2','PropParam2',)

where '*PropParam*' is optional.

The valid choices for '*PropName*' and '*PropParam*' are listed in this table.

<i>PropName</i>	<i>PropParam</i>
'ent', 'ento' or 'sizes' (see wptree)	Without ' <i>PropParam</i> ' or with ' <i>PropParam</i> ' = Vector of node indices, PropValue contains the entropy (or optimal entropy, or size) of the tree nodes in ascending node index order.
'cfs'	With ' <i>PropParam</i> ' = One terminal node index. cfs = read(T,'cfs',NODE) is equivalent to cfs = read(T,'data',NODE) and returns the coefficients of the terminal node NODE.

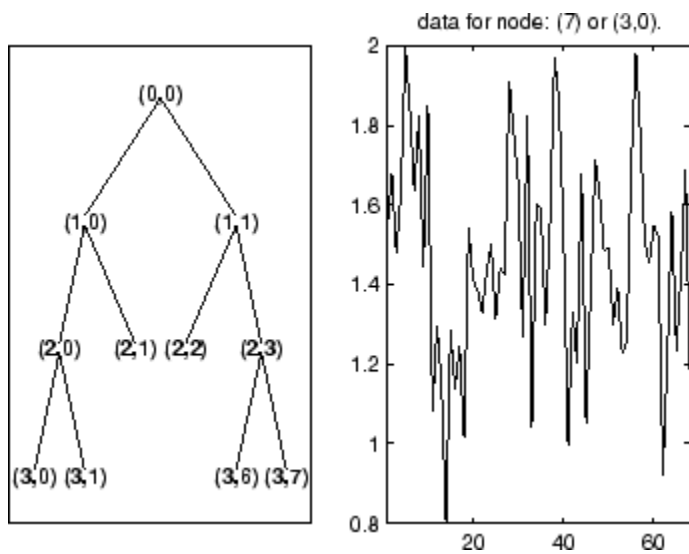
<i>PropName</i>	<i>PropParam</i>
'entName', 'entPar', 'wavName' (see wptree) or 'allcfs'	Without ' <i>PropParam</i> '. <i>cfs</i> = read(T, 'allcfs') is equivalent to <i>cfs</i> = read(T, 'data'). <i>PropValue</i> contains the desired information in ascending node index order of the tree nodes.
'wfilters' (see wfilters)	Without ' <i>PropParam</i> ' or with ' <i>PropParam</i> ' = 'd', 'r', 'l', 'h'.
'data'	Without ' <i>PropParam</i> ' or with ' <i>PropParam</i> ' = One terminal node index or ' <i>PropParam</i> ' = Column vector of terminal node indices. In this last case, <i>PropValue</i> is a cell array. Without ' <i>PropParam</i> ', <i>PropValue</i> contains the coefficients of the tree nodes in ascending node index order.

Examples

```
% Create a wavelet packet tree.
x = rand(1,512);
t = wpdec(x,3,'db3');
t = wpjoin(t,[4;5]);
plot(t);

% Click the node (3,0), (see the plot function).
l% Read values.

sAll = read(t,'sizes');
sNod = read(t,'sizes',[0,4,5]);
eAll = read(t,'ent');
eNod = read(t,'ent',[0,4,5]);
dAll = read(t,'data');
dNod = read(t,'data',[4;5]);
[lo_D,hi_D,lo_R,hi_R] = read(t,'wfilters');
[lo_D,lo_R,hi_D,hi_R] = read(t,'wfilters','l','wfilters','h');
[ent,ento,cfs4,cfs5] = read(t,'ent','ento','cfs',4,'cfs',5);
```



See Also

`disp` | `get` | `set` | `wptree` | `write`

Introduced before R2006a

readtree

Read wavelet packet decomposition tree from figure

Syntax

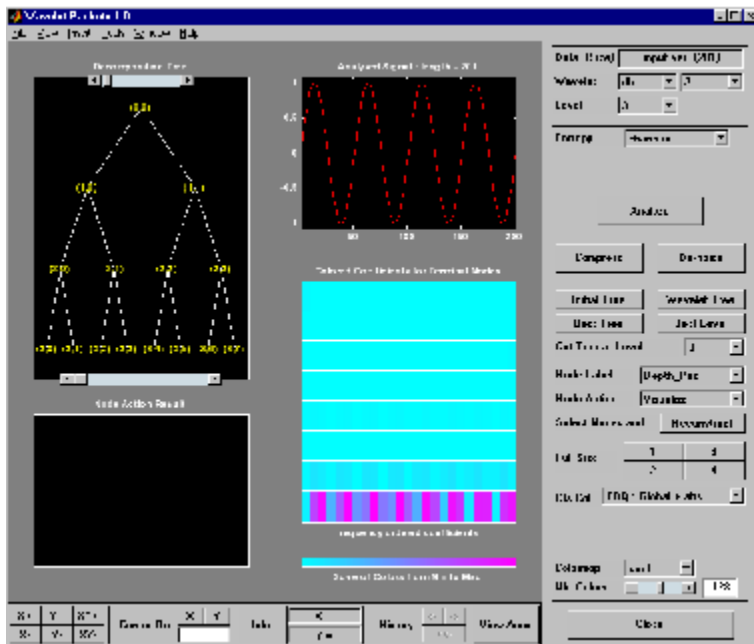
```
T = readtree(F)
```

Description

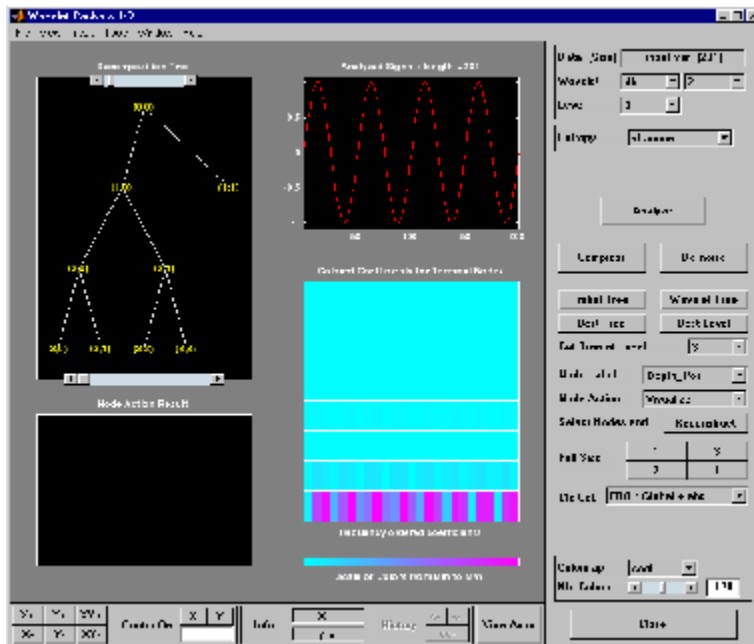
`T = readtree(F)` reads the wavelet packet decomposition tree from the figure whose handle is *F*.

Examples

```
% Create a wavelet packet tree.  
x = sin(8*pi*[0:0.005:1]);  
t = wpdec(x,3,'db2');  
  
% Display the generated tree in a Wavelet Packet 1-D GUI window.  
fig = drawtree(t);
```

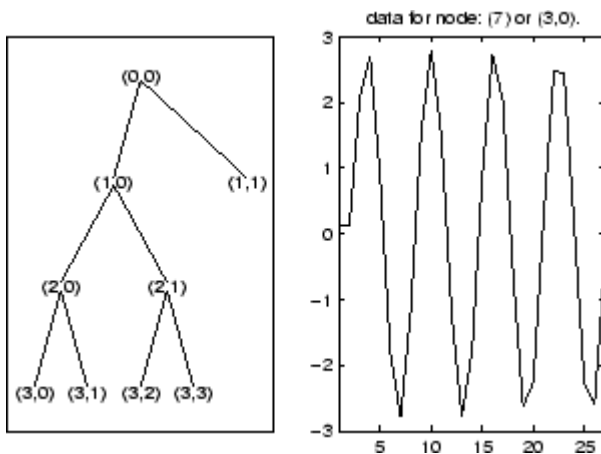


%-----
 % Use the GUI to split or merge Nodes.
 %-----



```
t = readtree(fig);
plot(t)
```

% Click the node (3,0), (see the plot function).



See Also

drawtree

Introduced before R2006a

removeLabelDefinition

Remove label definition from labeled signal set

Syntax

```
removeLabelDefinition(lss, lblname)
```

Description

`removeLabelDefinition(lss, lblname)` removes the label definition `lblname` from the labeled signal set `lss`. If you want to remove a sublabel, specify `lblname` as a two-element string array or two-element cell array of character vectors:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Examples

Remove Label Definition

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss

lss =
    labeledSignalSet with properties:
        Source: {2x1 cell}
        NumMembers: 2
        TimeInformation: "sampleRate"
        SampleRate: 4000
        Labels: [2x3 table]
        Description: "Characterize wave song regions"
```


Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Retrieve a hierarchical list of labels and sublabels.

```
labelDefinitionsHierarchy(lss)
```

```
ans =  
    'WhaleType  
      Sublabels: []  
    MoanRegions  
      Sublabels: []  
    TrillRegions  
      Sublabels: TrillPeaks  
,
```

Remove the sublabel that labels peaks in the trill regions.

```
removeLabelDefinition(lss,{'TrillRegions' 'TrillPeaks'})
```

```
labelDefinitionsHierarchy(lss)
```

```
ans =  
    'WhaleType  
      Sublabels: []  
    MoanRegions  
      Sublabels: []  
    TrillRegions  
      Sublabels: []  
,
```

Remove the label that specifies the whale type.

```
removeLabelDefinition(lss,"WhaleType")
```

```
getLabelNames(lss)
```

```
ans = 2x1 string array  
    "MoanRegions"  
    "TrillRegions"
```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

removeMembers

Remove members from labeled signal set

Syntax

```
removeMembers(lss,midxvect)
```

Description

`removeMembers(lss,midxvect)` removes the members specified in `midxvect` from the labeled signal set `lss`.

Examples

Remove Member

Load a labeled signal set containing recordings of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
      NumMembers: 2  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [2x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Remove the second member of the set.

```
removeMembers(lss,2)
lss
lss =
  labeledSignalSet with properties:
        Source: {[79572x1 double]}
      NumMembers: 1
    TimeInformation: "sampleRate"
      SampleRate: 4000
        Labels: [1x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Input Arguments

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1),randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midxvect — Subset member row numbers

vector of positive integers

Subset member row numbers, specified as a vector of positive integers. Each element of `midxvect` specifies a member row number as it appears in the “Labels” on page 1-0 table of the `labeledSignalSet` object `lss`.

Example: `[2 3 5 7 11 13 17]` chooses a subset of signals indexed by prime numbers.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

removePointValue

Remove row from point label

Syntax

```
removePointValue(lss,midx,lblname)
removePointValue(lss,midx,lblname,'LabelRowIndex',ridx)
removePointValue(lss,midx,lblname,'SublabelRowIndex',sridx)
removePointValue(lss,midx,lblname,'LabelRowIndex',
ridx,'SublabelRowIndex',sridx)
```

Description

`removePointValue(lss,midx,lblname)` removes all rows of the point label `lblname` for the member specified by `midx`.

- If `lblname` is a character vector or a string scalar, the function targets a parent label.
- If `lblname` is a two-element string array or a two-element cell array of character vectors, the function:
 - Interprets the first element as the name of a parent label.
 - Interprets the second element as the sublabel name of a point label.
 - Removes all the points of the sublabel.

`removePointValue(lss,midx,lblname,'LabelRowIndex',ridx)` removes a row, specified by `ridx`, of the point label `lblname` for the member `midx`.

If `lblname` is a two-element string array or a two-element cell array of character vectors, the function:

- Interprets the first element as the name of a parent label.
- Interprets the second element as the sublabel name of a point label.
- Removes all the point of the sublabel contained in row `ridx`.

`removePointValue(lss, midx, lblname, 'SublabelRowIndex', sridx)` removes the sublabel row specified by `sridx`. In this case, `lblname` must be a two-element string array or a two-element cell array of character vectors:

- The first element is the name of a parent attribute label.
- The second element is the sublabel name of a point label.

`removePointValue(lss, midx, lblname, 'LabelRowIndex', ridx, 'SublabelRowIndex', sridx)` removes the sublabel row specified by `sridx` of the ROI or point label row specified by `ridx`. In this case, `lblname` must be a two-element string array or a two-element cell array of character vectors:

- The first element is the name of a parent ROI or point label.
- The second element is the sublabel name of a point label.

Examples

Remove Point Value

Load a labeled signal set containing recordings of whale songs. Get the names of the labels and the number of members.

```
load whales
lss
```

```
lss =
  labeledSignalSet with properties:
        Source: {2x1 cell}
    NumMembers: 2
  TimeInformation: "sampleRate"
    SampleRate: 4000
        Labels: [2x3 table]
  Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

```
nm = lss.NumMembers;
```

Define a point label associated with the signal maximum.

```
themax = signalLabelDefinition('Maximum','LabelType','point', ...  
    'LabelDataType','numeric')
```

```
themax =  
    signalLabelDefinition with properties:
```

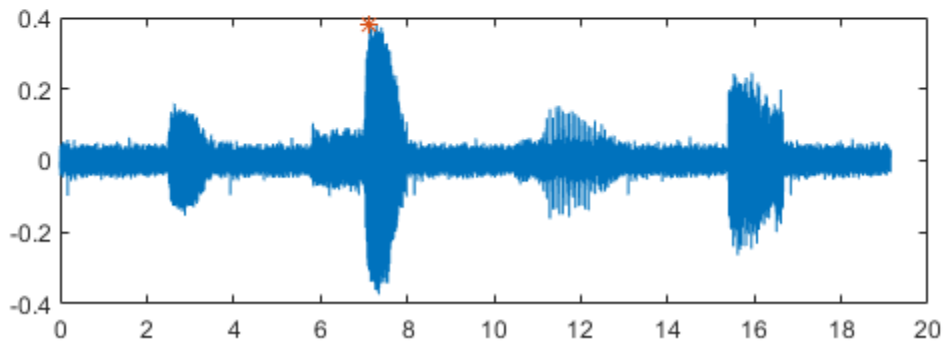
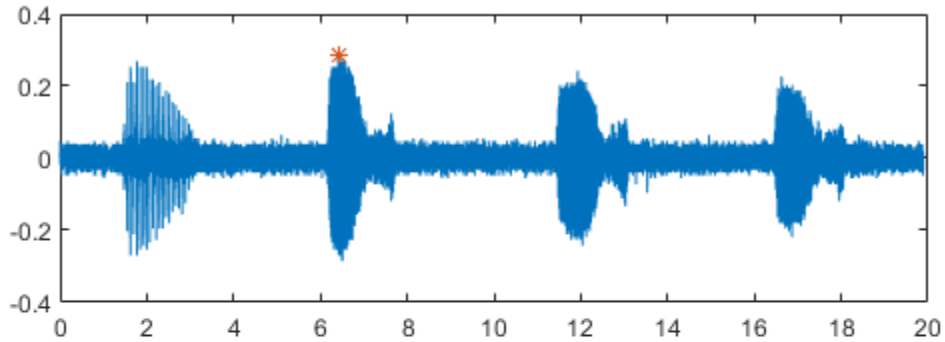
```
                Name: "Maximum"  
                LabelType: "point"  
                LabelDataType: "numeric"  
                ValidationFunction: []  
                PointLocationsDataType: "double"  
                DefaultValue: []  
                Sublabels: [0x0 signalLabelDefinition]  
                Tag: ""  
                Description: ""
```

Use `labeledSignalSet` to create a labeled signal set.

```
addLabelDefinitions(lss,themax)
```

Find the maxima of the signals and add their values to the labeled set.

```
figure  
for idx = 1:nm  
    sg = getSignal(lss,idx);  
    [mx,ix] = max(sg);  
    setLabelValue(lss,idx,'Maximum',ix,mx)  
  
    subplot(nm,1,idx)  
    plot((0:length(sg)-1)/lss.SampleRate,sg,ix/lss.SampleRate,mx,'*')  
end
```

Verify that the set includes the new point label.

```
getLabelValues(lss)
```

```
ans=2x4 table
```

	WhaleType	MoanRegions	TrillRegions	Maximum
Member{1}	blue	[3x2 table]	[1x3 table]	[1x2 table]
Member{2}	blue	[3x2 table]	[1x3 table]	[1x2 table]

Remove the 'Maximum' value for the first member of the set. Verify that the label is empty for the first member.

```
removePointValue(lss,1,'Maximum')
```

```
getLabelValues(lss,1)
```

```
ans=1x4 table
```

	WhaleType	MoanRegions	TrillRegions	Maximum
Member{1}	blue	[3x2 table]	[1x3 table]	[0x2 table]

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1),randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep",'LabelType','roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: {'Asleep' 'REM'} or ["Asleep" "REM"] specifies a region of a signal in which a patient undergoes REM sleep.

ridx — Label row index

positive integer

Label row index, specified as a positive integer. This argument applies only for ROI and point labels.

sridx — Sublabel row index

positive integer

Sublabel row index, specified as a positive integer. This argument applies only when a label and sublabel pair has been specified in `lblname` and the sublabel is of type ROI or point.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

removeRegionValue

Remove row from ROI label

Syntax

```
removeRegionValue(lss, midx, lblname)
removeRegionValue(lss, midx, lblname, 'LabelRowIndex', ridx)
removeRegionValue(lss, midx, lblname, 'SublabelRowIndex', sridx)
removeRegionValue(lss, midx, lblname, 'LabelRowIndex',
ridx, 'SublabelRowIndex', sridx)
```

Description

`removeRegionValue(lss, midx, lblname)` removes all rows of the ROI label `lblname` for the member specified by `midx`.

- If `lblname` is a character vector or a string scalar, the function targets a parent label.
- If `lblname` is a two-element string array or a two-element cell array of character vectors, the function:
 - Interprets the first element as the name of a parent label.
 - Interprets the second element as the sublabel name of an ROI label.
 - Removes all the regions of the sublabel.

`removeRegionValue(lss, midx, lblname, 'LabelRowIndex', ridx)` removes a row, specified by `ridx`, of the ROI label `lblname` for the member `midx`.

If `lblname` is a two-element string array or a two-element cell array of character vectors, the function:

- Interprets the first element as the name of a parent label.
- Interprets the second element as the sublabel name of an ROI label.
- Removes all the regions of the sublabel contained in row `ridx`.

`removeRegionValue(lss, midx, lblname, 'SublabelRowIndex', sridx)` removes the sublabel row specified by `sridx`. In this case, `lblname` must be a two-element string array or a two-element cell array of character vectors:

- The first element is the name of a parent attribute label.
- The second element is the sublabel name of an ROI label.

`removeRegionValue(lss, midx, lblname, 'LabelRowIndex', ridx, 'SublabelRowIndex', sridx)` removes the sublabel row specified by `sridx` of the ROI or point label row specified by `ridx`. In this case, `lblname` must be a two-element string array or a two-element cell array of character vectors:

- The first element is the name of a parent ROI or point label.
- The second element is the sublabel name of an ROI label.

Examples

Remove Region Value

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss
```

```
lss =
  LabeledSignalSet with properties:
    Source: {2x1 cell}
    NumMembers: 2
    TimeInformation: "sampleRate"
    SampleRate: 4000
    Labels: [2x3 table]
    Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Get the names and values of the labels in the set. For the following, concentrate on the second member of the set.

```
lbldefs = getLabelValues(lss)
```

```
lbldefs=2x3 table
           WhaleType   MoanRegions   TrillRegions
Member{1}   blue      [3x2 table]   [1x3 table]
Member{2}   blue      [3x2 table]   [1x3 table]
```

```
idx = 2;
```

Retrieve the signal and the time information. Plot the signal.

```
[lbs,info] = getLabeledSignal(lss,idx)
```

```
lbs=1x4 table
           Signal           WhaleType   MoanRegions   TrillRegions
Member{2} [76579x1 double]   blue          [3x2 table]   [1x3 table]
```

```
info = struct with fields:
    TimeInformation: "sampleRate"
    SampleRate: 4000
```

```
fs = info.SampleRate;
sg = getSignal(lss,idx);
t = (0:length(sg)-1)/fs;
```

```
plot(t,sg)
```

Highlight the moans and trills of the signal.

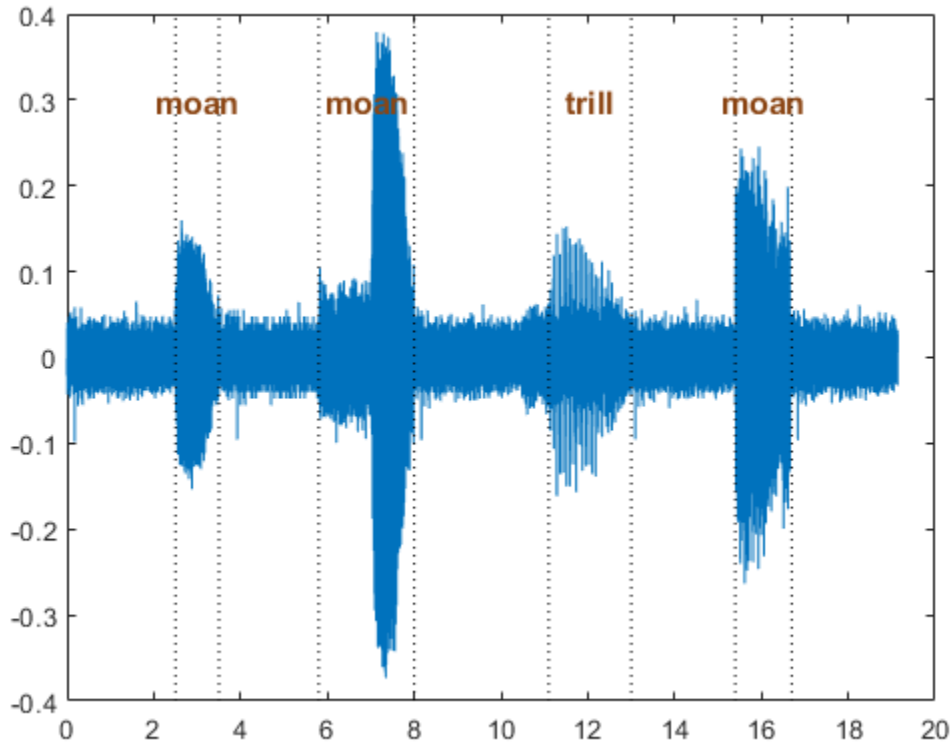
```
mvals = getLabelValues(lss,idx,'MoanRegions');
tvals = getLabelValues(lss,idx,'TrillRegions');
```

```
hold on
[X,Y] = meshgrid([mvals.ROIlimits;tvals.ROIlimits],ylim);
plot(X,Y,':k')
topts = {'HorizontalAlignment','center','FontWeight','bold', ...
        'FontSize',12,'Color',[139 69 19]/255};
text((X(1,1:4)+X(1,5:end))/2,Y(2,5:end)-0.1, ...
```

```

    ["moan" "moan" "moan" "trill"],topts{:})
hold off

```



Remove the second moan from the labels. Plot the signal again. Highlight the moans and trills.

```

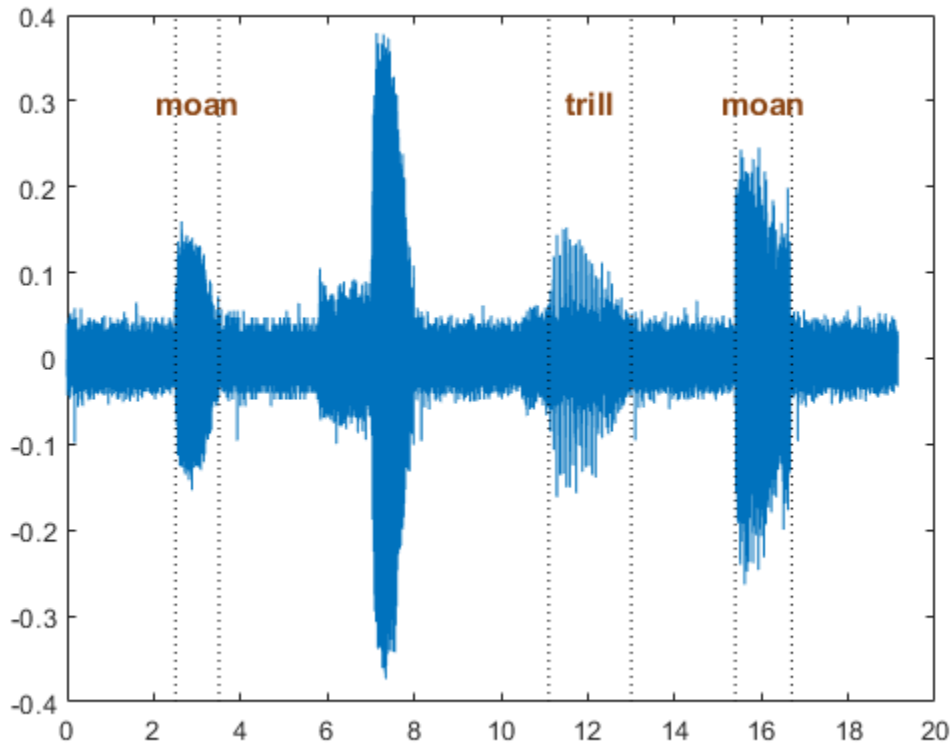
removeRegionValue(lss,idx,'MoanRegions','LabelRowIndex',2)

plot(t,sg)
mvals = getLabelValues(lss,idx,'MoanRegions');

hold on
[X,Y] = meshgrid([mvals.ROIlimits;tvals.ROIlimits],ylim);
plot(X,Y,':k')

```

```
topts = {'HorizontalAlignment','center','FontWeight','bold', ...  
        'FontSize',12,'Color',[139 69 19]/255};  
text((X(1,1:3)+X(1,4:end))/2,Y(2,4:end)-0.1, ...  
     ["moan" "moan" "trill"],topts{:})  
hold off
```



Input Arguments

lss — Labeled signal set
labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

ridx — Label row index

positive integer

Label row index, specified as a positive integer. This argument applies only for ROI and point labels.

sridx — Sublabel row index

positive integer

Sublabel row index, specified as a positive integer. This argument applies only when a label and sublabel pair has been specified in `lblname` and the sublabel is of type ROI or point.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

resetLabelValues

Reset labels to default values

Syntax

```
resetLabelValues(lss)
resetLabelValues(lss,midx)

resetLabelValues(lss,midx,lblname)
resetLabelValues( ____, 'LabelRowIndex', ridx)
```

Description

`resetLabelValues(lss)` resets all label values for all members of the labeled signal set `lss`.

`resetLabelValues(lss,midx)` resets all label values for the signals in the member specified by `midx`.

`resetLabelValues(lss,midx,lblname)` resets the values of label `lblname` for the signals in the member specified by `midx`. To reset a sublabel, make `lblname` a two-element string array or a two-element cell array of character vectors, with the first element containing the parent label name and the second element containing the sublabel name.

By default, the function resets all sublabels of a parent label. To target a sublabel of an ROI or point parent label, specify the parent label row index using `ridx`.

`resetLabelValues(____, 'LabelRowIndex', ridx)` specifies the row index of the ROI or point parent label for which you want to reset a sublabel value.

Examples

Reset Label Values

Load a labeled signal set containing recordings of whale songs. Get the names of the labels.

```
load whales
lss
```

```
lss =
  LabeledSignalSet with properties:
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

```
getLabelNames(lss)
```

```
ans = 3x1 string array
    "WhaleType"
    "MoanRegions"
    "TrillRegions"
```

Get the label values corresponding to the trill regions for the second signal in the set.

```
idx = 2;
getLabelValues(lss,idx,'TrillRegions')
```

```
ans=1x2 table
    ROIlimits    Value
    _____    _____
    11.1         13      [1]
```

Reset the values. Verify that 'TrillRegions' becomes an empty array.

```

resetLabelValues(lss,idx,'TrillRegions')
getLabelValues(lss,idx,'TrillRegions')
ans =
    0x2 empty table
getLabelValues(lss,idx)
ans=1x3 table
           WhaleType    MoanRegions    TrillRegions
           _____    _____    _____
Member{2}    blue        [3x2 table]    [0x3 table]

```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female')) specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. midx specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

ridx – Label row index

positive integer

Label row index, specified as a positive integer. This argument applies only for ROI and point labels.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

scal2frq

Scale to frequency

Syntax

```
F = scal2frq(A, 'wname', DELTA)
scal2frq(A, 'wname')
scal2frq(A, 'wname', 1)
```

Description

`F = scal2frq(A, 'wname', DELTA)` returns the pseudo-frequencies corresponding to the scales given by `A` and the wavelet function `'wname'` (see `wavefun` for more information) and the sampling period `DELTA`.

`scal2frq(A, 'wname')` is equivalent to `scal2frq(A, 'wname', 1)`.

There is only an approximate answer for the relationship between scale and frequency.

In wavelet analysis, the way to relate scale to frequency is to determine the center frequency of the wavelet, F_c , and use the following relationship:

$$F_a = \frac{F_c}{a}$$

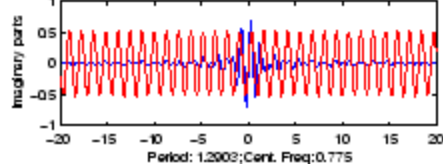
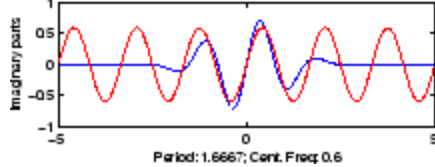
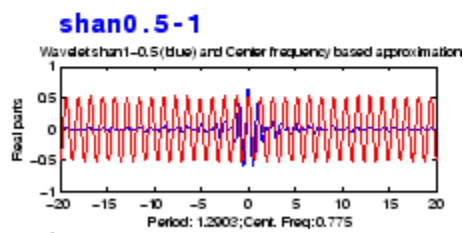
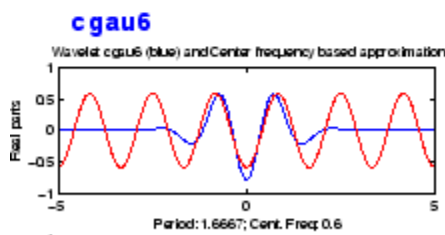
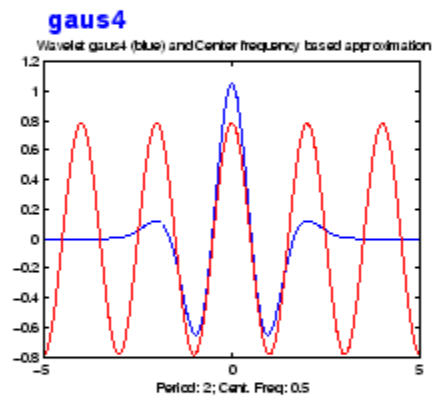
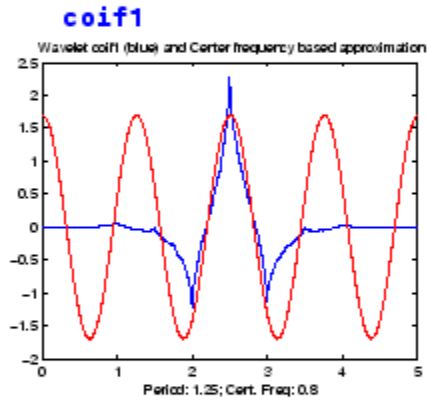
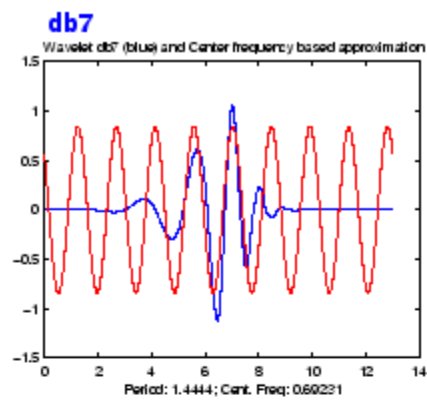
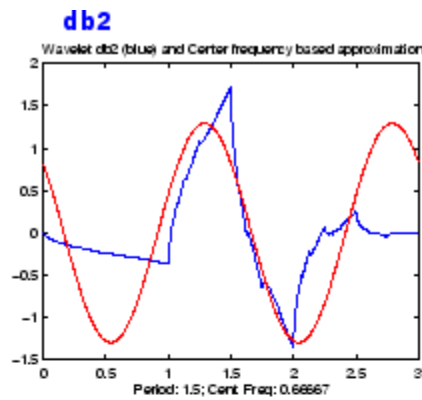
where

- a is a scale.
- F_c is the center frequency of the wavelet in Hz.
- F_a is the pseudo-frequency corresponding to the scale a , in Hz.

The idea is to associate with a given wavelet a purely periodic signal of frequency F_c . The frequency maximizing the Fourier transform of the wavelet modulus is F_c . `centfrq` computes the center frequency for a specified wavelet. From the above relationship, it can be seen that scale is inversely proportional to pseudo-frequency. For example, if the

scale increases, the wavelet becomes more spread out, resulting in a lower pseudo-frequency.

Some examples of the correspondence between the center frequency and the wavelet are shown in the following figure.



Center Frequencies for Real and Complex Wavelets

As you can see, the center frequency-based approximation captures the main wavelet oscillations. The center frequency is a convenient and simple characterization of the dominant frequency of the wavelet.

Examples

Scales To Frequencies

Construct a vector of scales with 10 voices per octave over five octaves. Assume the data are sampled at 10 kHz.

```
voicesperoctave = 10;
numoctaves = 5;
a0 = 2^(1/voicesperoctave);
Fs = 1e4;
scales = ...
    a0.^(voicesperoctave:numoctaves*voicesperoctave);
```

Convert the scales to approximate frequencies in hertz for the Morlet wavelet.

```
Frq = scal2frq(scales, 'morl', 1/Fs);
```

Determine the corresponding periods. Construct a table with the scales, the corresponding frequencies, and periods. Display the smallest 20 scales along with their corresponding frequencies and periods.

```
Frq = Frq(:);
scales = scales(:);
T = [scales.*(1/Fs) Frq 1./Frq];
T = array2table(T, 'VariableNames', {'Scale', 'Frequency', 'Period'});
T(1:20, :)
```

```
ans=20x3 table
      Scale      Frequency      Period
-----
      0.0002      4062.5      0.00024615
0.00021435      3790.4      0.00026382
0.00022974      3536.6      0.00028276
0.00024623      3299.8      0.00030305
  0.0002639      3078.8      0.0003248
0.00028284      2872.6      0.00034811
```

```

0.00030314    2680.3    0.0003731
0.0003249    2500.8    0.00039988
0.00034822    2333.3    0.00042858
0.00037321     2177    0.00045934
0.0004     2031.3    0.00049231
0.00042871    1895.2    0.00052764
0.00045948    1768.3    0.00056551
0.00049246    1649.9    0.0006061
0.0005278     1539.4    0.0006496
0.00056569    1436.3    0.00069623
:

```

Plot CWT with Frequencies in a Contour Plot

The example shows how to create a contour plot of the CWT using approximate frequencies in Hz.

Create a signal consisting of two sine waves with disjoint support in additive noise. Assume the signal is sampled at 1 kHz.

```

Fs = 1000;
t = 0:1/Fs:1-1/Fs;
x = 1.5*cos(2*pi*100*t).*(t<0.25)+1.5*cos(2*pi*50*t).*(t>0.5 & t<=0.75);
x = x+0.05*randn(size(t));

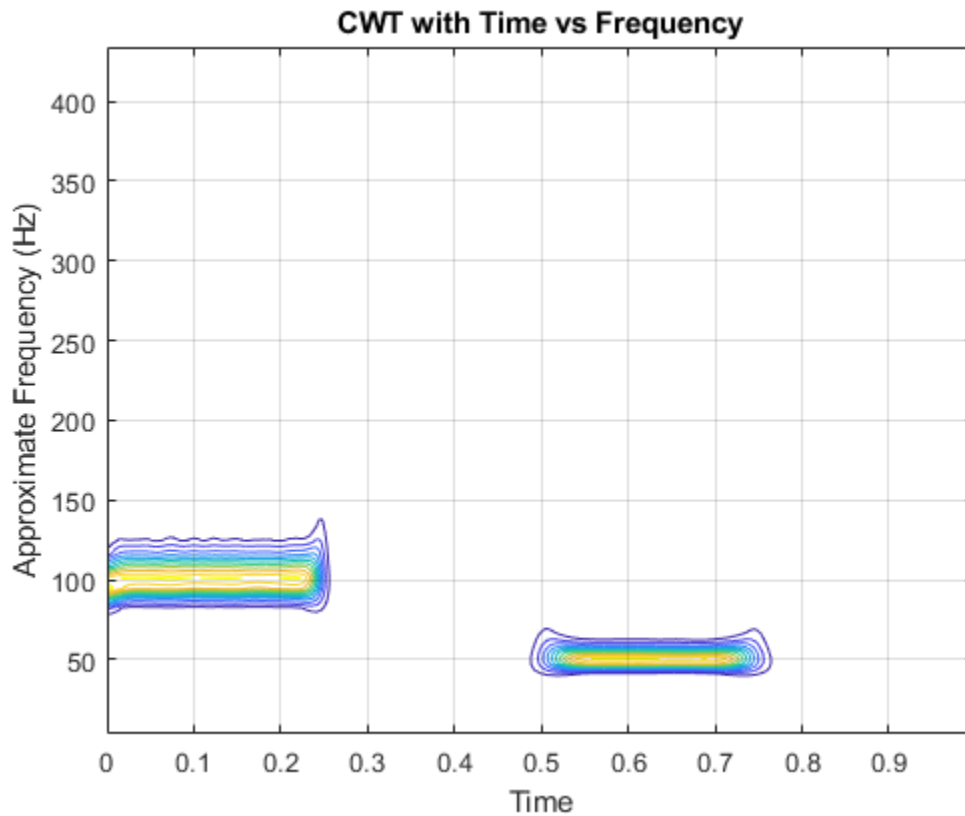
```

Obtain the CWT of the input signal and plot the result.

```

[cfs,f] = cwt(x,Fs);
contour(t,f,abs(cfs).^2);
axis tight;
grid on;
xlabel('Time');
ylabel('Approximate Frequency (Hz)');
title('CWT with Time vs Frequency');

```



References

Abry, P. (1997), *Ondelettes et turbulence. Multirésolutions, algorithmes de décomposition, invariance d'échelles*, Diderot Editeur, Paris.

See Also

centfrq

Introduced before R2006a

scales

CWT filter bank scales

Syntax

```
rs = scales(fb)  
[rs,cs] = scales(fb)
```

Description

`rs = scales(fb)` returns the raw (unitless) scales used in creating the wavelet bandpass filters. Scales are ordered from finest scale to coarsest scale.

`[rs,cs] = scales(fb)` returns the wavelet scales converted to units of the sampling frequency or sampling period.

Examples

CWT Filter Bank Scales

Create a CWT filter bank with sampling period equal to 0.001 seconds.

```
fb = cwtfilterbank('SamplingPeriod',seconds(0.001));
```

Obtain the raw and converted scales used in creating the wavelet bandpass filters.

```
[rs,cs] = scales(fb);
```

Obtain the filter bank bandpass center periods.

```
P = centerPeriods(fb);
```

Compare the finest converted scale with the smallest bandpass center period normalized by the sampling period.

```
min(cs)
```

```
ans = 2.3035
```

```
min(P)/seconds(0.001)
```

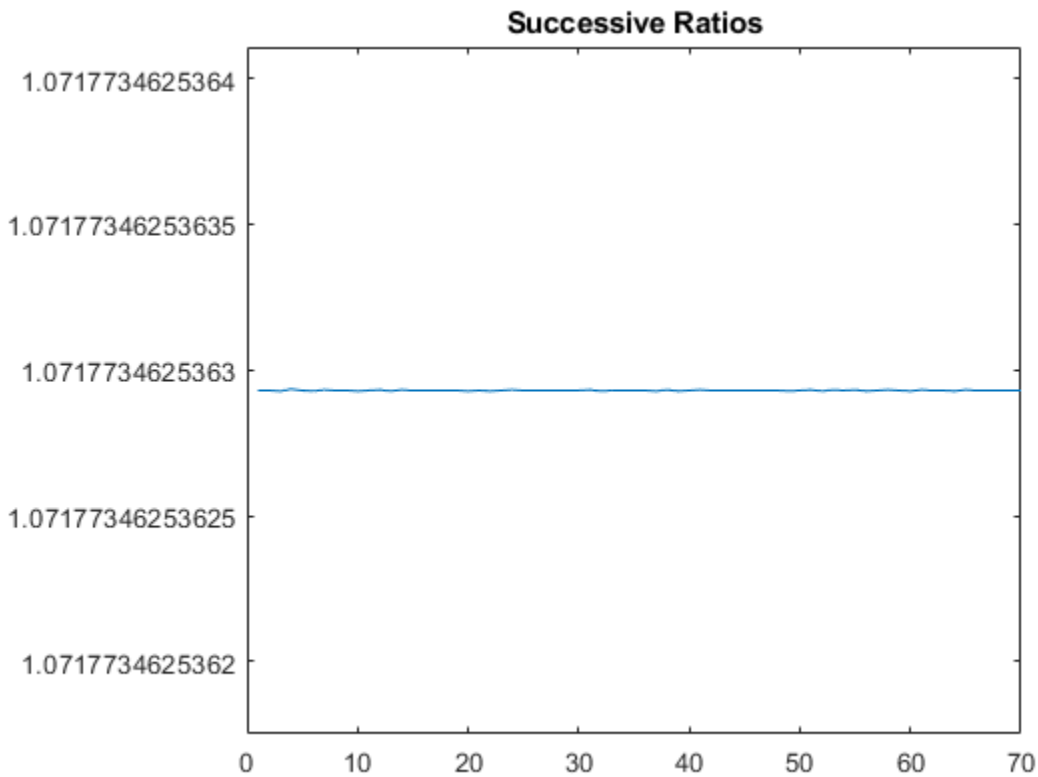
```
ans = 2.3035
```

The scales should increase by a factor of approximately $2^{(1/|NV|)}$ where **NV** is the number of voices per octave. The default value of **NV** is 10. Plot the ratios of successive scales, and compare with $2^{(1/10)}$.

```
2^(1/10)
```

```
ans = 1.0718
```

```
len = length(rs);  
plot(rs(2:len)./rs(1:len-1))  
title('Successive Ratios')
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

rs — Raw scales

real-valued vector

Raw scales used in creating the wavelet bandpass filters, returned as a real-valued vector of length N_s where N_s is the number of wavelet bandpass frequencies (equal to the number of scales).

Data Types: double

cs — Converted scales

real-valued vector

Converted scales used in creating the wavelet bandpass filters, returned as a real-valued vector of length N_s where N_s is the number of wavelet bandpass frequencies (equal to the number of scales). cs is in units of the sampling frequency or sampling period.

Data Types: double

See Also

[centerFrequencies](#) | [centerPeriods](#) | [cwtfilterbank](#)

Introduced in R2018a

scalingfunctions

DWT filter bank time-domain scaling functions

Syntax

```
phi = scalingfunctions(fb)  
[phi,t] = scalingfunctions(fb)
```

Description

`phi = scalingfunctions(fb)` returns the time-centered scaling functions for each level of the discrete wavelet transform (DWT) filter bank `fb`. `phi` is an L-by-N matrix where L is the filter bank `Level` and N is the filter bank `SignalLength`.

`[phi,t] = scalingfunctions(fb)` returns the sampling instants, `t`. `t` are returned in the interval $[-(N*DT)/2,(N*DT)/2]$, where N is the filter bank `SignalLength` and DT is the filter bank sampling period (reciprocal of the filter bank sampling frequency).

Examples

DWT Filter Bank Scaling Functions

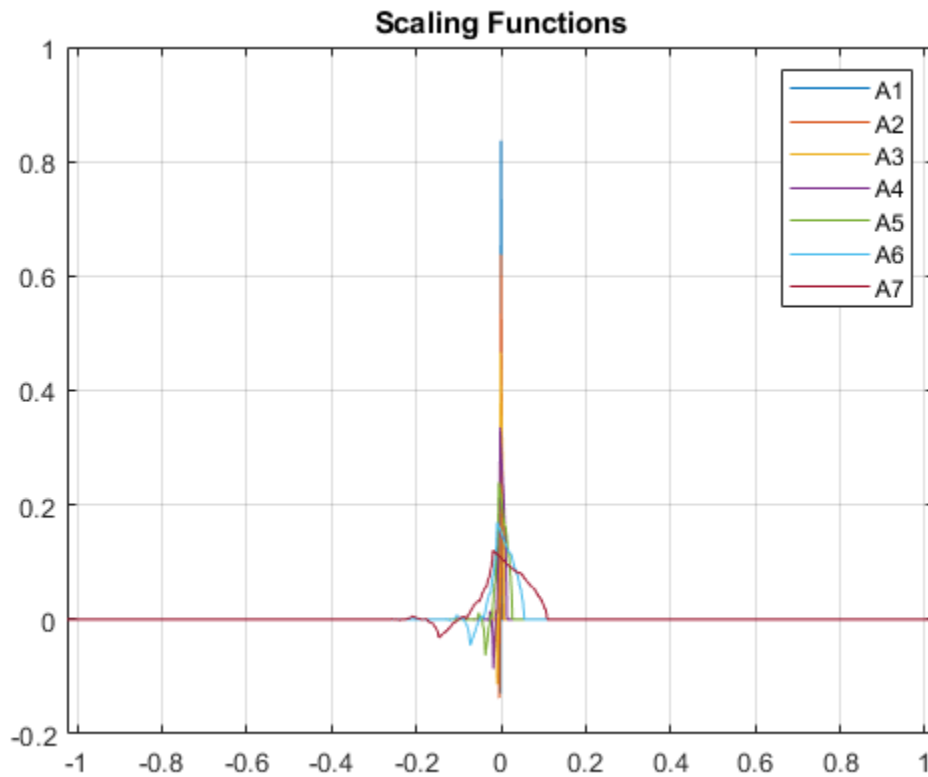
Create a seven level DWT filter bank for a length 2048 signal, using the Daubechies `db2` wavelet, and a sampling frequency of 1kHz.

```
wv = "db2";  
len = 2048;  
Fs = 1e3;  
lev = 7;  
fb = dwtfilterbank('SignalLength',len,'Wavelet',wv,'Level',lev,'SamplingFrequency',Fs)
```

Obtain and plot the scaling functions for each level of the filter bank.

```
[phi,t] = scalingfunctions(fb);  
plot(t,phi')
```

```
grid on
xlim([-len/2*1e-3 len/2*1e-3])
title('Scaling Functions')
legend('A1','A2','A3','A4','A5','A6','A7')
```



Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

phi — Time-centered scaling functions

real-valued matrix

Time-centered scaling functions of the filter bank `fb`, returned as a real-valued L-by-N matrix, where L is the filter bank `Level` and N is the `SignalLength`. The scaling functions are ordered in `phi` from the finest scale resolution to the coarsest scale resolution.

t — Sampling instants

real-valued vector

Sampling instants, returned as a real-valued vector `t` of length N, where N is the filter bank `SignalLength`. Sampling instants lie in the interval $-(N*DT)/2 \leq t < (N*DT)/2$, where DT is the filter bank sampling period (reciprocal of the filter bank sampling frequency).

See Also

`dwtfilterbank` | `wavelets`

Introduced in R2018a

scattergram

Visualize scattering or scalogram coefficients

Syntax

```
img = scattergram(sf,S)
img = scattergram(sf,U)
img = scattergram( ____,Name,Value)
scattergram( ____ )
```

Description

`img = scattergram(sf,S)` returns the scattergram as a matrix for the first-order scattering coefficients, `S`. The matrix `S` is the output of `scatteringTransform` computed using the scattering framework, `sf`.

`img = scattergram(sf,U)` returns the scattergram as a matrix for the first-order scalogram coefficients, `U`. The matrix `U` is the output of `scatteringTransform` computed using the scattering framework, `sf`.

`img = scattergram(____,Name,Value)` returns the scattergram with additional options specified by one or more `Name,Value` pair arguments. You can use this syntax with any of the input syntaxes shown previously.

`scattergram(____)` with no output arguments plots the scattergram in the current figure. You can use any of the input syntaxes shown previously.

Examples

Visualize Scattergram

Load an ECG signal sampled at 180 Hz. Create a scattering decomposition framework that can be used with the signal.

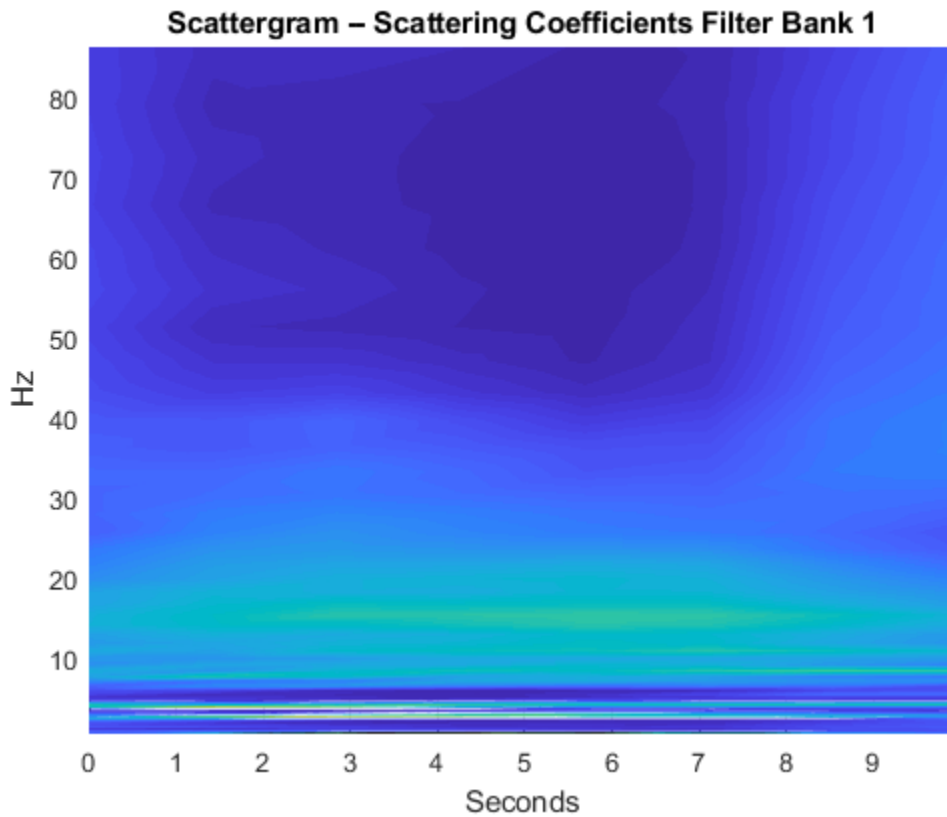
```
load wecg
Fs = 180;
sf = waveletScattering('SignalLength', numel(wecg), ...
    'SamplingFrequency', Fs);
```

Calculate the scattering transform of the signal.

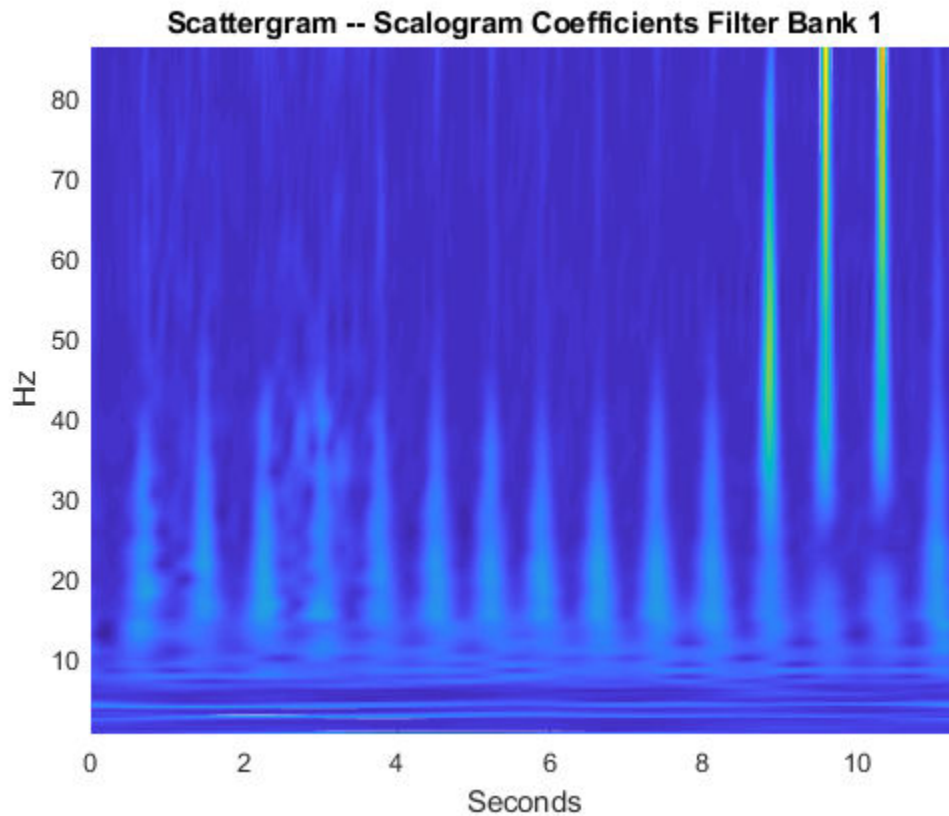
```
[S,U] = scatteringTransform(sf,wecg);
```

Visualize the scattergram for the first-order scattering and scalogram coefficients.

```
scattergram(sf,S,'FilterBank',1)
```



```
figure
scattergram(sf,U,'FilterBank',1)
```



Input Arguments

sf – Scattering decomposition framework

`waveletScattering` object

Scattering decomposition framework, specified as a `waveletScattering` object.

S – Scattering coefficients

cell array

Scattering coefficients, specified as a cell array. *S* is the output of `scatteringTransform` computed using the scattering framework, *sf*. For more information, see `scatteringTransform`.

U — Scalogram coefficients

cell array

Scalogram coefficients, specified as a cell array. *U* is the output of `scatteringTransform` computed using the scattering framework, *sf*. For more information, see `scatteringTransform`.

Name-Value Pair Arguments

Specify optional comma-separated pairs of *Name*, *Value* arguments. *Name* is the argument name and *Value* is the corresponding value. *Name* must appear inside quotes. You can specify several name and value pair arguments in any order as *Name1*, *Value1*, . . . , *NameN*, *ValueN*.

Example: `'FilterBank', 1` specifies the first filter bank.

FilterBank — Filter bank index

positive integer between 1 and the number of filter banks in *sf* inclusive

Filter bank index, specified as a positive number between 1 and the number of filter banks in *sf* inclusive. `scattergram` returns the scattergram for the specified filter bank in *sf*. The number of filter banks in *sf* is equal to the number of specified `QualityFactors` in *sf*.

If *FilterBank* is greater than 1, `scattergram` averages the scalogram or scattering coefficients over all paths terminating at each wavelet bandpass filter. To obtain paths with a common parent, use the `'Parent'` name-value pair.

P — Path parent index

nonnegative integer

Path parent index, specified as a nonnegative integer. The scalar *P* is a nonnegative integer representing the *P*-th wavelet filter at the filter bank `FilterBank - 1`. `scattergram` returns the scattergram for the path at the specified filter bank with parent *P*. If *FilterBank* is equal to 1, the zeroth filter bank corresponds to the input signal in the case of the scalogram coefficients and the lowpass filtering of the input signal with the scaling function in the case of the scattering coefficients. Lower values of *P* correspond to wavelets with higher bandpass frequencies.

If you specify `P`, you must specify the `FilterBank` name-value pair.

If you specify a value for `P` which results in a single child, the output `img` is a vector. The scattergram of a single child is a line plot. If you specify a value for `P` that results in no children, `scattergram` returns the scattergram for the filter bank specified by `FilterBank`.

Output Arguments

`img` — Scattergram

real-valued matrix | real-valued vector

Scattergram, returned as a real-valued matrix or vector. If you use the `Parent` name-value pair and specify a value which results in a single child, `img` is a vector. If the parent has more than one child, `img` is a matrix.

See Also

`waveletScattering`

Introduced in R2018b

scatteringTransform

Wavelet 1-D scattering transform

Syntax

```
s = scatteringTransform(sf,x)  
[s,u] = scatteringTransform(sf,x)
```

Description

`s = scatteringTransform(sf,x)` returns the wavelet 1-D scattering transform of `x` for the scattering decomposition framework, `sf`. `x` is a real-valued vector or matrix. If `x` is a matrix, `scatteringTransform` applies the scattering transform to each column of `x` separately. `s` is a cell array with $Nfb + 1$ elements, where Nfb is the number of filter banks in `sf`. The number of filter banks is also equal to the number of elements in the `QualityFactors` value of `sf`. Equivalently, the number of elements in `s` is equal to the number of orders in the scattering decomposition. Each element of `s` is a MATLAB table.

The precision of the scattering coefficients depends on the precision specified in the framework `sf`.

`[s,u] = scatteringTransform(sf,x)` returns `s`, the wavelet 1-D scattering transform, and `u`, the scalogram coefficients for each of the scattering orders. `u` is a cell array with $Nfb + 1$ elements, where Nfb is the number of filter banks in the scattering framework. The number of filter banks is also equal to the number of elements in the `QualityFactors` value of `sf`. Equivalently, the number of elements in `u` is equal to the number of orders in the scattering decomposition. Each element of `u` is a MATLAB table.

The precision of the scalogram coefficients depends on the precision specified in the framework `sf`.

Examples

Scattering Transform of ECG Signal

This example shows how to return the wavelet 1-D scattering transform of a real-valued signal.

Load an ECG signal sampled at 180 Hz.

```
load wecg
Fs = 180;
```

Create a scattering decomposition framework to apply to the signal. Compute the scattering transform of the signal.

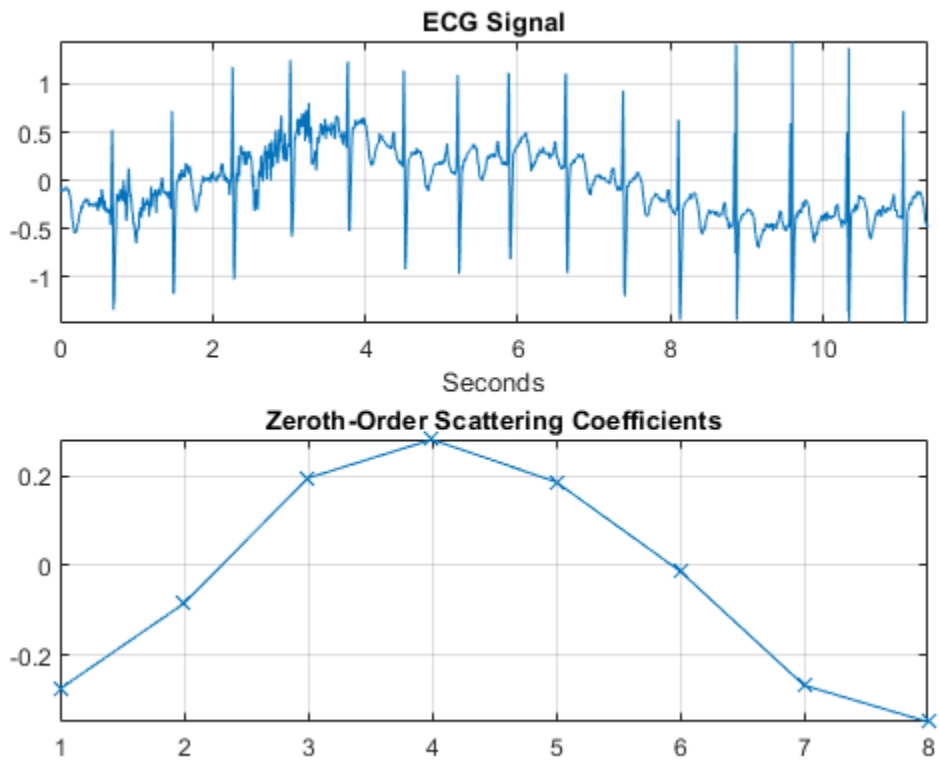
```
sf = waveletScattering('SignalLength', numel(wecg), ...
    'SamplingFrequency', Fs)
```

```
sf =
    waveletScattering with properties:
        SignalLength: 2048
        InvarianceScale: 5.6889
        QualityFactors: [8 1]
        Boundary: "periodic"
        SamplingFrequency: 180
        Precision: "double"
        OversamplingFactor: 0
```

```
[S,U] = scatteringTransform(sf,wecg);
```

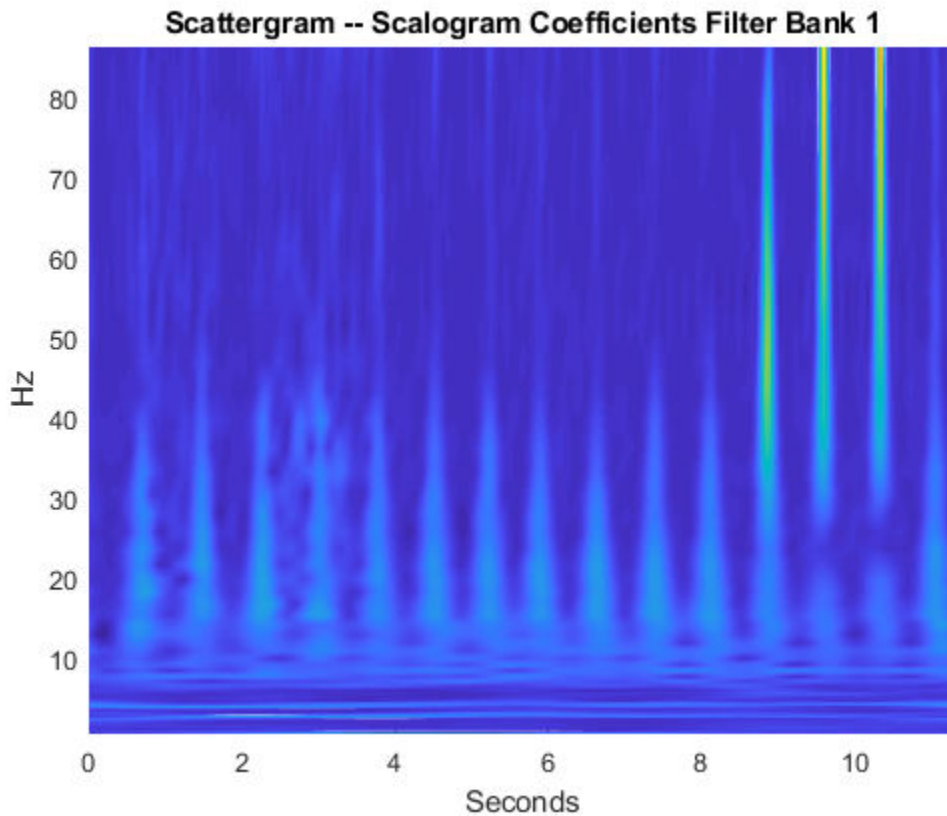
Plot the signal and the zeroth-order scattering coefficients. Note that the invariance scale is one half the duration of the signal.

```
t = [0:length(wecg)-1]/Fs;
subplot(2,1,1)
plot(t,wecg)
grid on
axis tight
xlabel('Seconds')
title('ECG Signal')
subplot(2,1,2)
plot(S{1}.signals{1}, 'x-')
grid on
axis tight
title('Zeroth-Order Scattering Coefficients')
```



Visualize the scattergram for the first-order scalogram coefficients.

```
figure  
scattergram(sf,U,'FilterBank',1)
```



Input Arguments

sf — **Scattering decomposition framework**

`waveletScattering` object

Scattering decomposition framework, specified as a `waveletScattering` object.

x — **Input data**

real-valued vector | real-valued matrix

Input data, specified as a real-valued vector or matrix. If x is a vector, the number of samples in x must equal the `SignalLength` value of `sf`. If x is a matrix, the number of rows in x must equal the `SignalLength` value of `sf`.

Data Types: `double` | `single`

Output Arguments

s — Scattering coefficients

cell array

Scattering coefficients, returned as a cell array. `s` is a cell array with $Nfb+1$ elements where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` value of `sf`. Equivalently, the number of elements of `s` is equal to the number of orders in `sf`.

The precision of the scattering coefficients depends on the precision specified in the framework `sf`.

Each element of `s` is a MATLAB table with the following variables:

signals — Scattering coefficients

cell array

Scattering coefficients, returned as a cell array. If x is a vector, each element of `signals` is a column vector. If x has N columns, each element of `signals` is a M -by- N matrix, where M is the number of scattering coefficients.

Data Types: `double` | `single`

path — Scattering path

row vector

Scattering path used to obtain the scattering coefficients, returned as a row vector. Each column of `path` corresponds to one element of the path. The scalar 0 denotes the original signal. Positive integers in the L^{th} column denote the corresponding wavelet filter in the $(L-1)^{\text{th}}$ filter bank. Wavelet bandpass filters are ordered by decreasing center frequency.

Data Types: `double`

bandwidth — Bandwidth of scattering coefficients

scalar

Bandwidth of the scattering coefficients, returned as a scalar. If you specify a sampling frequency in the scattering framework, the bandwidth is in hertz. Otherwise, the bandwidth is in cycles/sample.

Data Types: double

resolution — Base-2 log resolution

scalar

Base-2 log resolution of the scattering coefficients, returned as a scalar.

Data Types: double

u — Scalogram coefficients

cell array

Scalogram coefficients, returned as a cell array. `u` is a cell array with $Nfb+1$ elements, where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` value of `sf`. Equivalently, the number of elements of `u` is equal to the number of orders in `sf`.

The precision of the scalogram coefficients depends on the precision specified in the framework `sf`.

Each element of `u` is a MATLAB table with the following variables:

coefficients — Scalogram coefficients

cell array

Scalogram coefficients, returned as a cell array. If `x` is a vector, each element of `coefficients` is a column vector. If `x` has N columns, each element of `coefficients` is a M -by- N matrix, where M is the number of scalogram coefficients.

Data Types: double | single

path — Scattering path

row vector

Scattering path used to obtain the scalogram coefficients, returned as a row vector. Each column of `path` corresponds to one element of the path. The scalar 0 denotes the original signal. Positive integers in the L^{th} column denote the corresponding wavelet filter in the $(L-1)^{\text{th}}$ filter bank. Wavelet bandpass filters are ordered by decreasing center frequency.

Data Types: double

bandwidth — Bandwidth of scalogram coefficients

scalar

Bandwidth of the scalogram coefficients, returned as a scalar. If you specify a sampling frequency in the scattering decomposition framework, the bandwidth is in hertz. Otherwise, the bandwidth is in cycles/sample.

Data Types: double

resolution — Base-2 log resolution

scalar

Base-2 log resolution of the scalogram coefficients, returned as a scalar.

Data Types: double

See Also

`featureMatrix` | `waveletScattering`

Introduced in R2018b

scatteringTransform

Wavelet 2-D scattering transform

Syntax

```
s = scatteringTransform(sf,im)
[s,u] = scatteringTransform(sf,im)
```

Description

`s = scatteringTransform(sf,im)` returns the wavelet 2-D scattering transform of `im` for `sf`, the scattering decomposition framework. `im` is a real-valued 2-D matrix or 3-D matrix. If `im` is 3-D, the size of the third dimension must equal 3. The row and column sizes of `im` must match the `ImageSize` value of `sf`. The output `s` is a cell array with $Nfb + 1$ elements, where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` property of `sf`. Equivalently, the number of elements in `s` is equal to the number of orders in the scattering decomposition. Each element of `s` is a MATLAB table.

`[s,u] = scatteringTransform(sf,im)` also returns the wavelet scalogram coefficients for `im`. The output `u` is a cell array with $Nfb + 1$ elements, where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` property of `sf`. Equivalently, the number of elements in `u` is equal to the number of orders in the scattering decomposition. Each element of `u` is a MATLAB table.

Examples

Compare Scattering and Scalogram Coefficients

This example shows that scattering coefficients are lowpassed versions of scalogram coefficients.

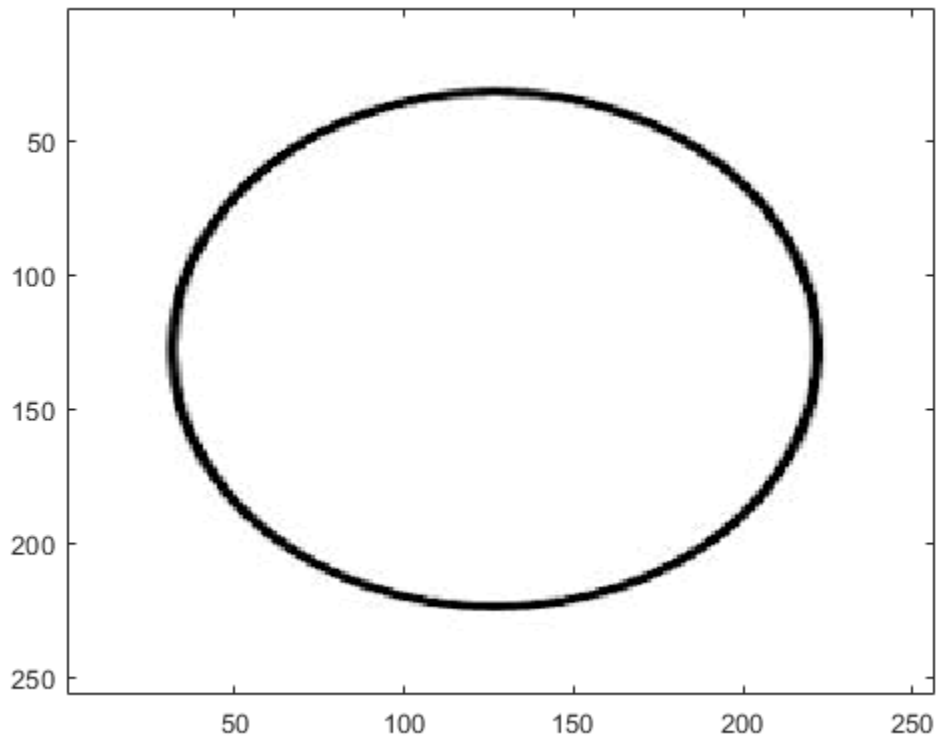
Load an RGB image. Display the red channel.

```
im = imread('circle.jpg');  
size(im)
```

```
ans = 1×3
```

```
    256    256     3
```

```
figure  
imagesc(im(:,:,1))  
colormap gray;
```



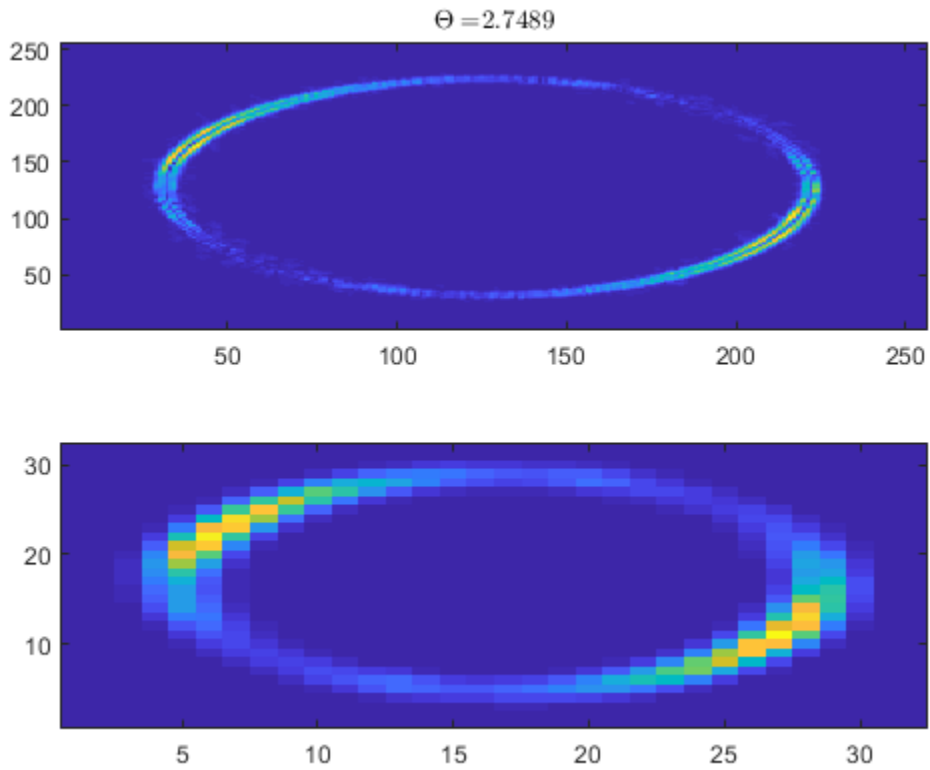
For RGB images, the size of the third dimension must be 3. You only have to specify the row and column sizes of the image when you create the scattering framework. Create a scattering framework to apply to the image and take the scattering transform.

```
sf = waveletScattering2('ImageSize',[256 256],'InvarianceScale',32,...
    'NumRotations',[8 8]);
[S,U] = scatteringTransform(sf,im);
```

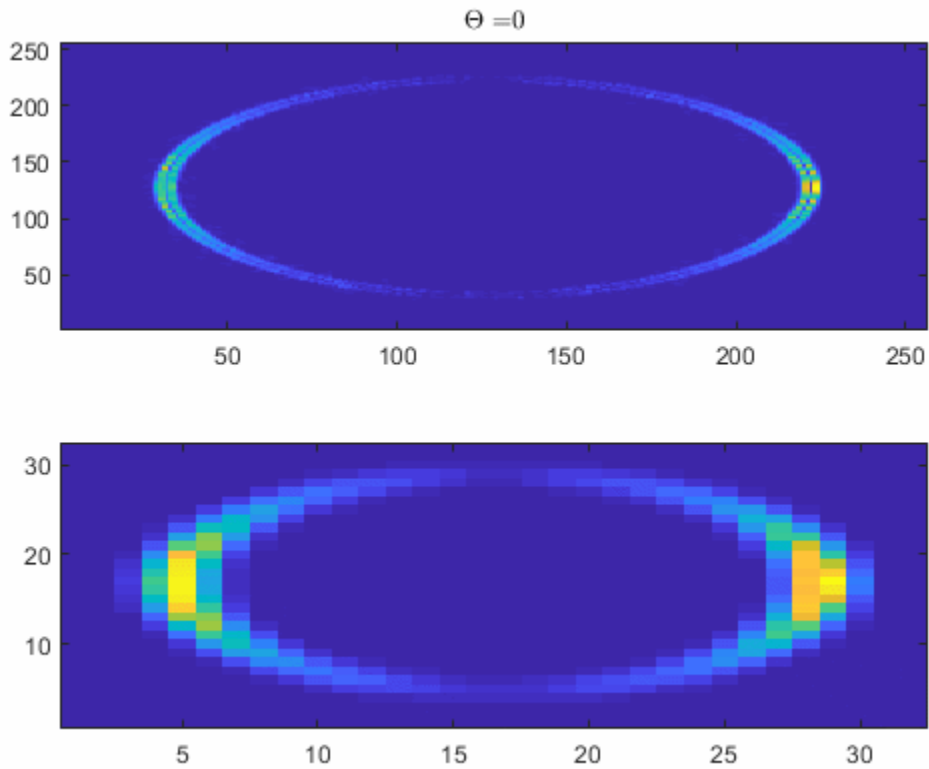
The image and coefficient fields in S and U are M -by- N -by-3. The M -by- N dimensions are constant only in the scattering images because the scaling function has fixed bandwidth, while the wavelets have different bandwidths.

Use a for-loop and plot the red channel for the scalogram and scattering coefficients for the 8 rotation angles in the scattering transform. Note how the scattering coefficients are lowpass versions of the scalogram coefficients.

```
[~,~,~,filterparams] = sf.filterbank();
theta = filterparams{1}.rotations;
figure
for k = 1:numel(theta)
    subplot(2,1,1)
    imagesc(U{2}.coefficients{k}(:,:,1));
    axis xy
    title(['$$\Theta = $$' num2str(theta(k))],'Interpreter','Latex');
    subplot(2,1,2)
    imagesc(S{2}.images{k}(:,:,1));
    axis xy
    pause(1)
end
```



The above for-loop results in an animation identical to the one below.



Input Arguments

sf – Scattering decomposition framework

waveletScattering2 object

Scattering decomposition framework, specified as a waveletScattering2 object.

im – Input image

real-valued matrix

Input image, specified as a real-valued 2-D matrix or 3-D matrix. If `im` is 3-D, `im` is assumed to be a color image in the RGB color space, and the size of the third dimension must equal 3. The row and column sizes of `im` must match the `ImageSize` property of `sf`.

Output Arguments

s — Scattering coefficients

cell array

Scattering coefficients, returned as a cell array. `s` is a cell array with $Nfb+1$ elements where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` property of `sf`. Equivalently, the number of elements in `s` is equal to the number of orders in the scattering decomposition. Each element of `s` is a MATLAB table with these variables:

images — Scattering coefficients

cell array

Scattering coefficients, returned as a cell array. Each element of `images` is an M -by- N or M -by- N -by-3 matrix.

path — Scattering path

row vector

Scattering path used to obtain the scattering coefficients, returned as a row vector. Each column of `path` corresponds to one element of the path. The scalar 0 denotes the original image. Positive integers in the L th column denote the corresponding wavelet filter in the $(L-1)$ th filter bank. Wavelet bandpass filters are ordered by decreasing center frequency.

There are `NumRotations` wavelets per center frequency pair.

bandwidth — Bandwidth of scattering coefficients

scalar

Bandwidth of scattering coefficients, returned as a scalar. The bandwidth is symmetric in the x and y directions.

resolution — Base-2 log resolution

scalar

Base-2 log resolution of the scattering coefficients, returned as a scalar.

u — Scalogram coefficients

cell array

Scalogram coefficients, returned as a cell array. `u` is a cell array with $Nfb+1$ elements, where Nfb is the number of filter banks in the scattering decomposition framework. Nfb is equal to the number of elements in the `QualityFactors` property of `sf`. Equivalently, the number of elements in `u` is equal to the number of orders in the scattering decomposition. Each element of `u` is a MATLAB table with these variables:

coefficients — Scalogram coefficients

cell array

Scalogram coefficients, returned as a cell array. Each element of `coefficients` is an M -by- N or M -by- N -by-3 matrix.

path — Scattering path

row vector

Scattering path used to obtain the scalogram coefficients, returned as a row vector. Each column of `path` corresponds to one element of the path. The scalar 0 denotes the original image. Positive integers in the L th column denote the corresponding wavelet filter in the $(L-1)$ th filter bank. Wavelet bandpass filters are ordered by decreasing center frequency.

There are `NumRotations` wavelets per center frequency pair.

bandwidth — Bandwidth of scalogram coefficients

scalar

Bandwidth of scalogram coefficients, returned as a scalar.

resolution — Base-2 log resolution

scalar

Base-2 log resolution of the scattering coefficients, returned as a scalar.

See Also`waveletScattering2`**Introduced in R2019a**

set

WPTREE field contents

Syntax

```
T = set(T,'FieldName1',FieldValue1,'FieldName2',FieldValue2, ...)
```

Description

`T = set(T,'FieldName1',FieldValue1,'FieldName2',FieldValue2, ...)` sets the content of the specified fields for the WPTREE object T.

For the fields that are objects or structures, you can set the subfield contents, giving the name of these subfields as '*FieldName*' values.

The valid choices for '*FieldName*' are

'dtree'	DTREE parent object
'wavInfo'	Structure (wavelet information)

The fields of the wavelet information structure, 'wavInfo', are also valid for '*FieldName*':

'wavName'	Wavelet name
'Lo_D'	Low Decomposition filter
'Hi_D'	High Decomposition filter
'Lo_R'	Low Reconstruction filter
'Hi_R'	High Reconstruction filter

'entInfo'	Structure (entropy information)
-----------	---------------------------------

The fields of the entropy information structure, 'entInfo', are also valid for '*FieldName*':

'entName'	Entropy name
'entPar'	Entropy parameter

Or fields of DTREE parent object:

'ntree'	NTREE parent object
'allNI'	All nodes information
'terNI'	Terminal nodes information

Or fields of NTREE parent object:

'wtbo'	WTBO parent object
'order'	Order of the tree
'depth'	Depth of the tree
'spsch'	Split scheme for nodes
'tn'	Array of terminal nodes of the tree

Or fields of WTBO parent object:

'wtboInfo'	Object information
'ud'	Userdata field

Caution The set function should only be used to set the field 'ud'.

See Also

disp | get | read | write

Introduced before R2006a

setLabelValue

Set label value in labeled signal set

Syntax

```
setLabelValue(lss,midx,lblname,val)
setLabelValue(lss,midx,lblname,limits,val)
setLabelValue(lss,midx,lblname,locs,val)
setLabelValue( ____, 'LabelRowIndex',ridx)
setLabelValue( ____, 'SublabelRowIndex',sridx)
```

Description

setLabelValue(lss,midx,lblname,val) sets the attribute label lblname to value val, for the member of labeled signal set lss specified in midx. Omit val if lblname has a default value and you want to set the label to the default value.

setLabelValue(lss,midx,lblname,limits,val) adds regions delimited by limits to the ROI label named lblname. The number of rows of limits specifies the number of added regions.

setLabelValue(lss,midx,lblname,locs,val) adds points to the point label named lblname. locs specifies the number of added points and their locations.

setLabelValue(____, 'LabelRowIndex',ridx) specifies the row index, ridx, of an ROI or point label. The specified value replaces the current value of that row. If you omit this argument, the function appends ROI or point values to any existing label values.

setLabelValue(____, 'SublabelRowIndex',sridx) specifies the row index, sridx, of an ROI or point sublabel. The specified value replaces the current value of that sublabel row.

Examples

Set Label Value

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss

lss =
    LabeledSignalSet with properties:

        Source: {2x1 cell}
        NumMembers: 2
        TimeInformation: "sampleRate"
        SampleRate: 4000
        Labels: [2x3 table]
        Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Add a new label to the signal set, corresponding to the maximum value of each member.

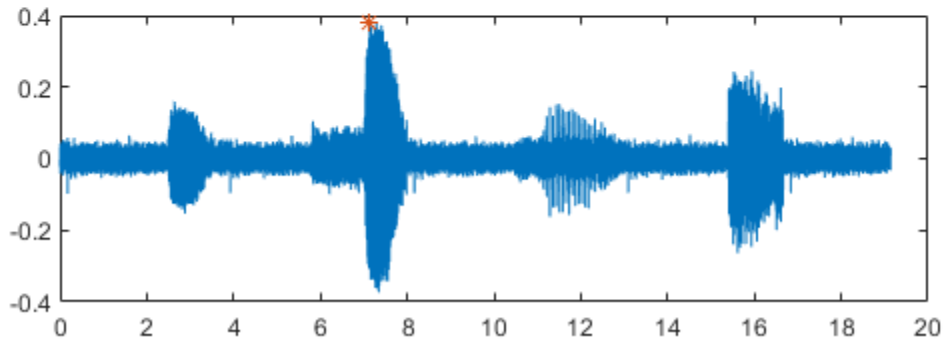
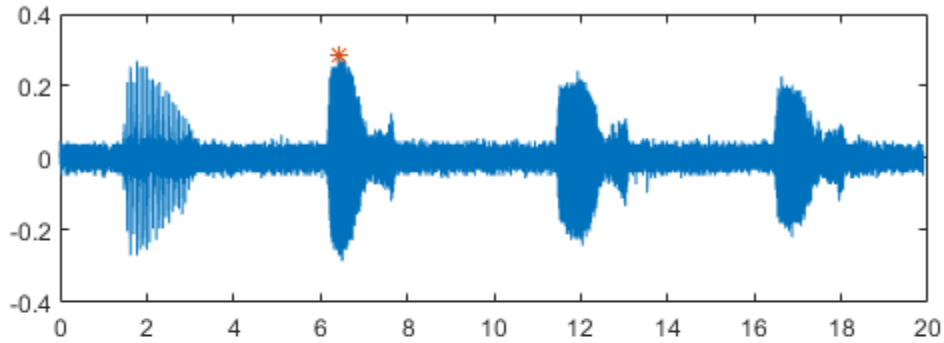
```
theMax = signalLabelDefinition('Maximum', ...
    'LabelDataType','numeric', ...
    'Description','Maximum value of the signal');
addLabelDefinitions(lss,theMax)
```

For each labeled signal, set the value of the new label to the signal maximum. Plot the signals and their maxima.

```
fs = lss.SampleRate;
for k = 1:lss.NumMembers
    sg = getSignal(lss,k);
    [mx,ix] = max(sg);

    setLabelValue(lss,k,'Maximum',mx)

    subplot(2,1,k)
    plot((0:length(sg)-1)/fs,sg,ix/fs,mx,'*')
end
```



Display the names and values of the labels in the set.

```
lbldefs = getLabelValues(lss)
```

```
lbldefs=2x4 table
```

	WhaleType	MoanRegions	TrillRegions	Maximum
Member{1}	blue	[3x2 table]	[1x3 table]	[0.2850]
Member{2}	blue	[3x2 table]	[1x3 table]	[0.3791]

Decide that the signal maximum is better represented as a point label than as an attribute. Remove the numeric definition and redefine the maximum.

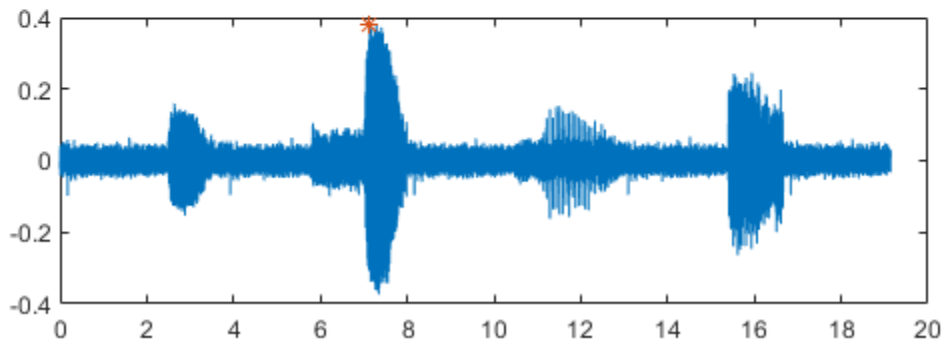
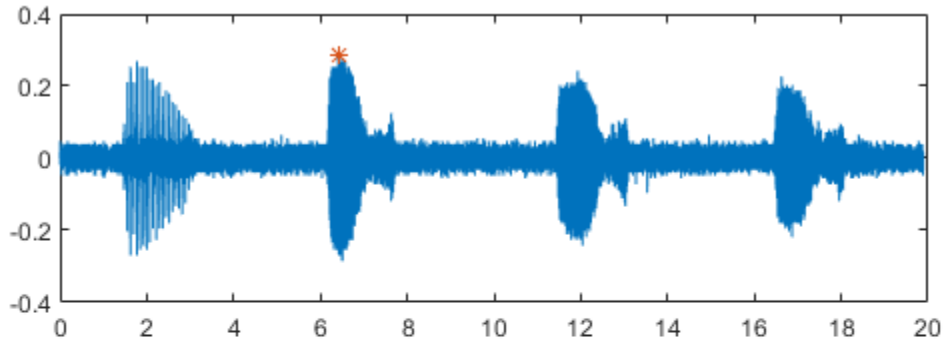
```
removeLabelDefinition(lss, 'Maximum')
theMax = signalLabelDefinition('Maximum', ...
    'LabelType', 'point', 'LabelDataType', 'numeric', ...
    'Description', 'Maximum value of the signal');
addLabelDefinitions(lss, theMax)
```

For each labeled signal, set the value of the new label to the signal maximum.

```
for k = 1:lss.NumMembers
    sg = getSignal(lss,k);
    [mx,ix] = max(sg);
    setLabelValue(lss,k, 'Maximum', ix/fs, mx)
end
```

Plot the signals and their maxima.

```
for k = 1:lss.NumMembers
    subplot(2,1,k)
    sg = getSignal(lss,k);
    peaks = getLabelValues(lss,k, 'Maximum');
    plot((0:length(sg)-1)/fs, sg, ...
        peaks.Location, cell2mat(peaks.Value), '*')
end
```



Input Arguments

lss — Labeled signal set
labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

lblname — Label or sublabel name

character vector | string scalar | cell array of character vectors | string array

Label name, specified as a character vector or string scalar.

Label or sublabel name. To specify a label, use a character vector or a string scalar. To specify a sublabel, use a two-element cell array of character vectors or a two-element string array:

- The first element is the name of the parent label.
- The second element is the name of the sublabel.

When targeting a sublabel of an ROI or point label, you must also specify the 'LabelRowIndex' of the parent label whose label you want to set. The row of the parent must already exist before you can set a sublabel value to it.

Example: `signalLabelDefinition("Asleep", 'LabelType', 'roi')` specifies a label of name "Asleep" for a region of a signal in which a patient is asleep during a clinical trial.

Example: `{'Asleep' 'REM'}` or `["Asleep" "REM"]` specifies a region of a signal in which a patient undergoes REM sleep.

val — Label values

numeric value or array | logical value or array | categorical value or array | character vector or cell array of character vectors | string or string array | table or table array | timetable or timetable array

Label values, specified as a numeric, logical, or categorical value, as a string, as a table, or as a timetable. `val` can also be an array of any of the previous types. `val` must be of the data type specified for `lblname`.

- If you specify `locs`, then `val` must have the same number of elements as `locs`.
- If you specify `limits`, then `val` must have a number of elements equal to the number of rows in `limits`.

- If `limits` has more than one row, and `lblname` is of type `'numeric'` or `'logical'`, then `val` must be a vector or a cell array.
- If `limits` has more than one row, and `lblname` is of type `'string'` or `'categorical'`, then `val` must be a string array or a cell array of character vectors.
- If `limits` has more than one row, and `lblname` is of type `'table'` or `'timetable'`, then `val` must be a cell array of tables or timetables.

limits — Region limits

two-column matrix

Region limits, specified as a two-column matrix.

- If `lss` does not have time information, then `limits` defines the minimum and maximum indices over which the regions are defined.
- If `lss` has time information, then `limits` defines the minimum and maximum instants over which the regions are defined.

`limits` must be of the data type specified by the “`ROIlimitsDataType`” on page 1-0 property of the label definition for `lblname`.

Example: `seconds([0:3;1:4]')`

Example: `[0:3;1:4]'`

locs — Point locations

vector

Point locations, specified as a vector.

- If `lss` does not have time information, then `locs` defines the indices corresponding to the point locations.
- If `lss` has time information, then `locs` defines the instants corresponding to the point locations.

`locs` must be of the data type specified by the “`PointLocationsDataType`” on page 1-0 property of the label definition for `lblname`.

ridx — Label row index

positive integer

Label row index, specified as a positive integer. This argument applies only for ROI and point labels.

sridx — Sublabel row index

positive integer

Sublabel row index, specified as a positive integer. This argument applies only when a label and sublabel pair has been specified in `lblname` and the sublabel is of type ROI or point.

See Also

`labeledSignalSet` | `signalLabelDefinition`

Introduced in R2018b

setMemberNames

Set member names in labeled signal set

Syntax

```
setMemberNames(lss, mnames)
setMemberNames(lss, mnames, midx)
```

Description

`setMemberNames(lss, mnames)` sets the names of the members of the labeled signal set `lss` to `mnames`. The length of `mnames` must be equal to the number of members.

`setMemberNames(lss, mnames, midx)` sets the name of the member specified by `midx`.

Examples

Set Member Names

Load a labeled signal set containing recordings of whale songs.

```
load whales
lss
```

```
lss =
  labeledSignalSet with properties:
```

```
      Source: {2x1 cell}
      NumMembers: 2
      TimeInformation: "sampleRate"
      SampleRate: 4000
      Labels: [2x3 table]
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.

Use `setLabelValue` to add data to the set.

Set the names of the set members to the whales' nicknames.

```
setMemberNames(lss, {'Brutus' 'Lucy'})
```

Return a string array with the names of the members.

```
getMemberNames(lss)
```

```
ans = 2x1 string array
    "Brutus"
    "Lucy"
```

Input Arguments

lss — Labeled signal set

labeledSignalSet object

Labeled signal set, specified as a labeledSignalSet object.

Example: `labeledSignalSet({randn(100,1) randn(10,1)}, signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

mnames — Member names

character vector | string scalar | cell array of character vectors | string array

Member names, specified as a character vector, a string scalar, a cell array of character vectors, or a string array.

Example: `labeledSignalSet({randn(100,1) randn(10,1)}, 'MemberNames', {'llama' 'alpaca'})` specifies a set of random signals with two members, 'llama' and 'alpaca'.

midx — Member row number

positive integer

Member row number, specified as a positive integer. `midx` specifies the member row number as it appears in the “Labels” on page 1-0 table of a labeled signal set.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2019a

shanwavf

Complex Shannon wavelet

Syntax

[PSI,X] = shanwavf(LB,UB,N,FB,FC)

Description

[PSI,X] = shanwavf(LB,UB,N,FB,FC) returns values of the complex Shannon wavelet. The complex Shannon wavelet is defined by a bandwidth parameter FB, a wavelet center frequency FC, and the expression

$$\text{PSI}(X) = (\text{FB}^{\wedge}0.5) * (\text{sinc}(\text{FB}*X) .* \text{exp}(2*i*pi*FC*X))$$

on an N point regular grid in the interval [LB,UB].

FB and FC must be such that $\text{FC} > 0$ and $\text{FB} > 0$.

Output arguments are the wavelet function PSI computed on the grid X.

Examples

Complex Shannon Wavelet

Obtain and plot a complex Shannon wavelet. Set the bandwidth and center frequency parameters.

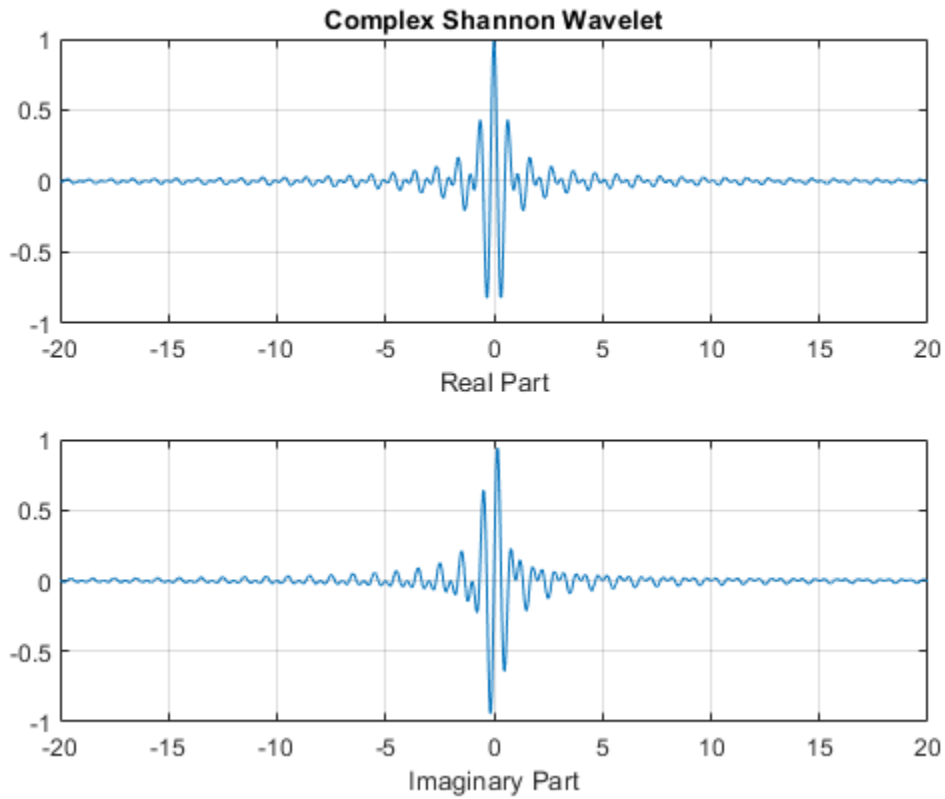
```
fb = 1;  
fc = 1.5;
```

Set the effective support and number of sample points.

```
lb = -20;  
ub = 20;  
n = 1000;
```

Obtain the complex-valued Shannon wavelet and plot the real and imaginary parts.

```
[psi,x] = shanwavf(lb,ub,n,fb,fc);  
subplot(2,1,1)  
plot(x,real(psi))  
title('Complex Shannon Wavelet')  
xlabel('Real Part')  
grid on  
subplot(2,1,2)  
plot(x,imag(psi))  
xlabel('Imaginary Part')  
grid on
```



References

Teolis, A. (1998), *Computational signal processing with wavelets*, Birkäuser, p. 62.

See Also

waveinfo

Introduced before R2006a

signalLabelDefinition

Create signal label definition

Description

Use `signalLabelDefinition` to create signal label definitions for data sets. The labels can correspond to attributes, regions, or points of interest. Use a vector of `signalLabelDefinition` objects to create a `labeledSignalSet`.

Creation

Syntax

```
sld = signalLabelDefinition(name)
sld = signalLabelDefinition(name,Name,Value)
```

Description

`sld = signalLabelDefinition(name)` creates a signal label definition object, `sld`, with the “Name” on page 1-0 property set to `name` and other properties set to default values.

`sld = signalLabelDefinition(name,Name,Value)` sets “Properties” on page 1-1099 using name-value pairs. You can specify multiple name-value pairs. Enclose each property name in quotes.

Input Arguments

name — Label name

character vector | string scalar

Label name, specified as a character vector or string scalar.

Data Types: `char` | `string`

Properties

Name — Name of label

character vector | string scalar

Name of label, specified as a character vector or string scalar.

Data Types: char | string

LabelType — Type of label

'attribute' (default) | 'roi' | 'point'

Type of label, specified as one of the following:

- 'attribute' — Define signal characteristics.
- 'roi' — Define signal characteristics over regions of interest.
- 'point' — Define signal characteristics over points of interest.

Data Types: char | string

LabelDataType — Data type of label

'logical' (default) | 'categorical' | 'numeric' | 'string' | 'table' | 'timetable'

Data type of label, specified as 'logical', 'categorical', 'numeric', 'string', 'table', or 'timetable'. Use the “Categories” on page 1-0 property to specify the array of categories when this property is set to 'categorical'.

Data Types: char | string

Categories — Label category names

string array | cell array of character vectors

Label category names, specified as a string array or a cell array of character vectors. The array must have unique elements. This property applies only when the “LabelDataType” on page 1-0 property is set to 'categorical'.

Example: 'LabelDataType', 'categorical', 'Categories', ["apple", "orange"]

Data Types: char | string

ROIlimitsDataType — Data type of ROI limits

'double' (default) | 'duration'

Data type of ROI limits, specified as either 'double' or 'duration'. This property applies only when “LabelType” on page 1-0 is set to 'roi'.

Data Types: char | string

PointLocationsDataType — Data type of point locations

'double' (default) | 'duration'

Data type of point locations, specified as either 'double' or 'duration'. This property applies only when “LabelType” on page 1-0 is set to 'point'.

Data Types: char | string

ValidationFunction — Validation function

function handle

Validation function, specified as a function handle and used when setting label values in a labeledSignalSet object. This property applies only when “LabelDataType” on page 1-0 is set to 'categorical', 'logical', 'numeric', 'table', or 'timetable'. If not specified, the function checks only that its input values are of the correct data type. If “LabelDataType” on page 1-0 is set to 'categorical', the function checks that the input is one of the values specified using “Categories” on page 1-0. The function takes an input value and returns true if the value is valid and false if the value is invalid.

Example:

```
'LabelDataType','numeric','DefaultValue',1,'ValidationFunction',@(x)  
)x<2
```

Data Types: function_handle

DefaultValue — Default value of label

[] (default) | LabelDataType value

Default value of label, specified as a value of the type specified using “LabelDataType” on page 1-0. If “LabelDataType” on page 1-0 is set to 'categorical', then “DefaultValue” on page 1-0 must be one of the values specified using “Categories” on page 1-0.

Example: 'LabelDataType','categorical','Categories',
["apple","orange"],'DefaultValue',"apple"

Data Types: char | double | logical | string | table

Description — Label description

character vector | string scalar

Label description, specified as a character vector or string scalar.

Example: 'Description', 'Patient is asleep'

Data Types: char | string

Tag — Label tag identifier

character vector | string scalar

Label tag identifier, specified as a character vector or string scalar. Use this property to identify the same label in a larger labeling scheme or public labeling set.

Example: 'Tag', 'Peak1'

Data Types: char | string

Sublabels — Array of sublabels

signal label definition object

Array of sublabels, specified as a signal label definition object. To specify more than one sublabel, set this property to a vector of signal label definition objects. Use this property to create a relationship between a parent label and its children.

Note Sublabels cannot have sublabels.

Example: 'Sublabels',
[signalLabelDefinition("negative"),signalLabelDefinition("positive")
]

Object Functions

labelDefinitionsHierarchy	Get hierarchical list of label and sublabel names
labelDefinitionsSummary	Get summary table of signal label definitions

Examples

Label Definitions for Whale Songs

Consider a set of whale sound recordings. The recorded whale sounds consist of trills and moans. You want to look at each signal and label it to identify the whale type, the trill regions, and the moan regions. For each trill region, you also want to label the signal peaks higher than a certain threshold.

Signal Label Definitions

Define an attribute label to store whale types. The possible categories are blue whale, humpback whale, and white whale.

```
dWhaleType = signalLabelDefinition('WhaleType', ...
    'LabelType', 'attribute', ...
    'LabelDataType', 'categorical', ...
    'Categories', ["blue" "humpback" "white"], ...
    'Description', 'Whale type');
```

Define a region-of-interest (ROI) label to capture moan regions. Define another ROI label to capture trill regions.

```
dMoans = signalLabelDefinition('MoanRegions', ...
    'LabelType', 'roi', ...
    'LabelDataType', 'logical', ...
    'Description', 'Regions where moans occur');
```

```
dTrills = signalLabelDefinition('TrillRegions', ...
    'LabelType', 'roi', ...
    'LabelDataType', 'logical', ...
    'Description', 'Regions where trills occur');
```

Finally, define a point label to capture the trill peaks. Set this label as a sublabel of the dTrills definition.

```
dTrillPeaks = signalLabelDefinition('TrillPeaks', ...
    'LabelType', 'point', ...
    'LabelDataType', 'numeric', ...
    'Description', 'Trill peaks');
```

```
dTrills.Sublabels = dTrillPeaks;
```

Labeled Signal Set

Create a `labeledSignalSet` with the whale signals and the label definitions. Add label values to identify the whale type, the moan and trill regions, and the peaks of the trills.

```
load labelwhalesignals
lbldefs = [dWhaleType dMoans dTrills];

lss = labeledSignalSet({whale1 whale2},lbldefs,...
    'SampleRate',Fs,'Description','Characterize wave song regions');
```

Visualize the label hierarchy and label properties using `labelDefinitionsHierarchy` and `labelDefinitionsSummary`.

```
labelDefinitionsHierarchy(lss)
```

```
ans =
    'WhaleType
      Sublabels: []
    MoanRegions
      Sublabels: []
    TrillRegions
      Sublabels: TrillPeaks
    ,
```

```
labelDefinitionsSummary(lss)
```

```
ans=3x9 table
    LabelName      LabelType      LabelDataType      Categories      ValidationFunction
    _____      _____      _____      _____      _____
    "WhaleType"     "attribute"     "categorical"     [3x1 string]     ["N/A"]
    "MoanRegions"   "roi"           "logical"         ["N/A" ]         []
    "TrillRegions"  "roi"           "logical"         ["N/A" ]         []
```

The signals in the loaded data correspond to songs of two blue whales. Set the 'WhaleType' values for both signals.

```
setLabelValue(lss,1,'WhaleType','blue');
setLabelValue(lss,2,'WhaleType','blue');
```

Visualize the 'Labels' property. The table has the newly added 'WhaleType' values for both signals.

```
lss.Labels
```

```
ans=2x3 table
    WhaleType      MoanRegions      TrillRegions
```

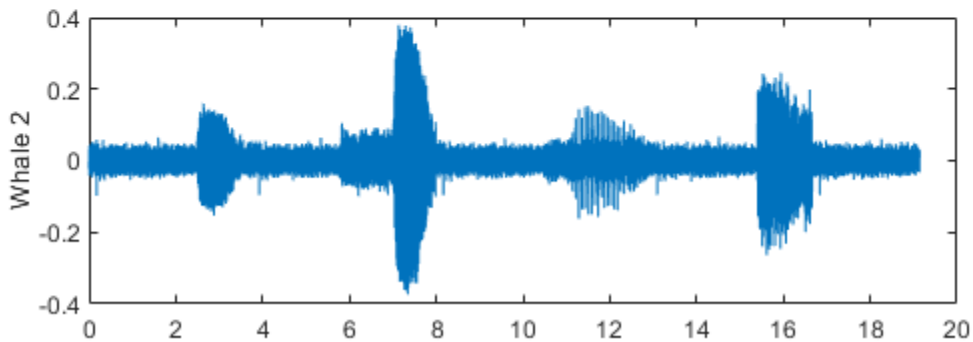
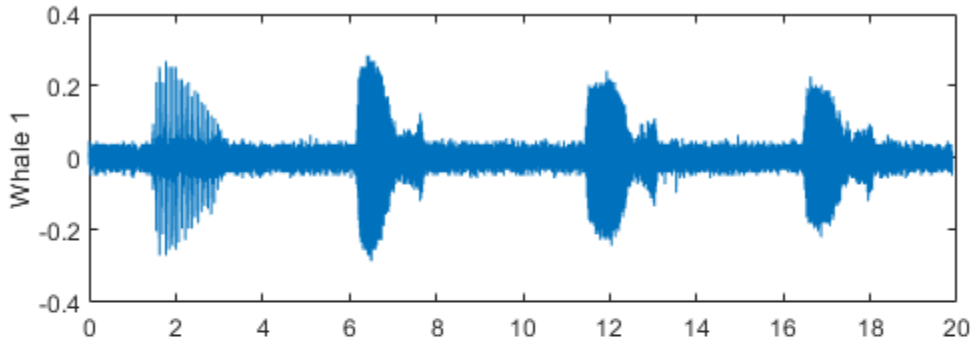
Member{1}	blue	[0x2 table]	[0x3 table]
Member{2}	blue	[0x2 table]	[0x3 table]

Visualize Regions and Points

Visualize the whale songs to identify the trill and moan regions.

```
subplot(2,1,1)
plot((0:length(whale1)-1)/Fs,whale1)
ylabel('Whale 1')
hold on

subplot(2,1,2)
plot((0:length(whale2)-1)/Fs,whale2)
ylabel('Whale 2')
hold on
```



Add the moan and trill regions to the labeled set. For ROI labels, specify the ROI limits in seconds and the label values. Label the different regions in the plots using an auxiliary function.

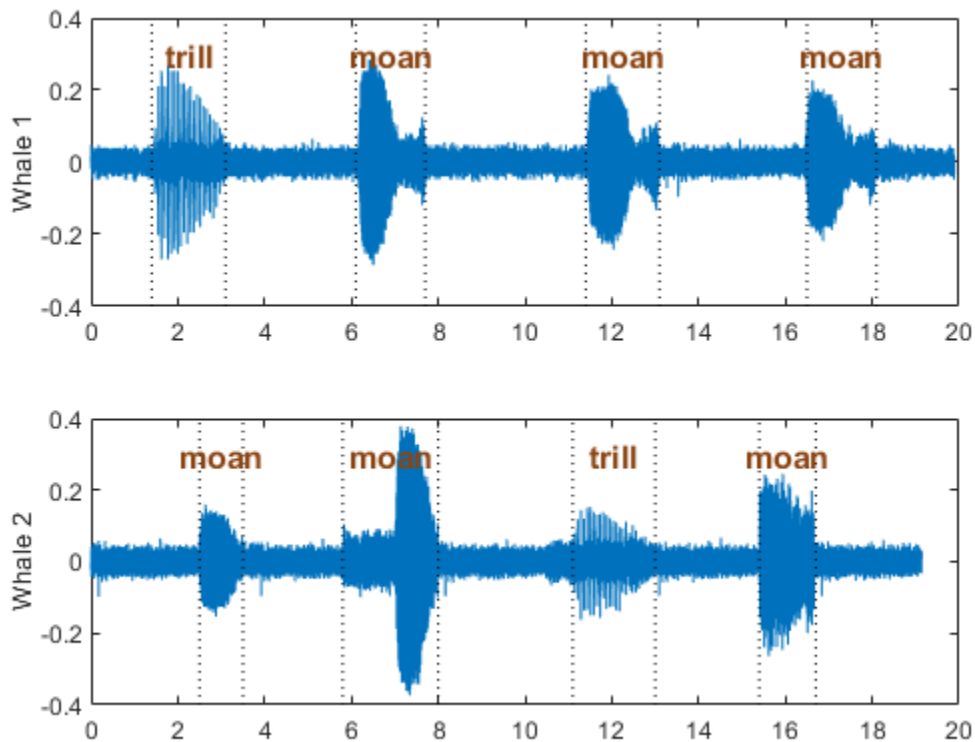
```
moanRegionsWhale1 = [6.1 7.7; 11.4 13.1; 16.5 18.1];
setLabelValue(lss,1,'MoanRegions',moanRegionsWhale1,[true true true]);
```

```
trillRegionWhale1 = [1.4 3.1];
setLabelValue(lss,1,'TrillRegions',trillRegionWhale1,true);
```

```
subplot(2,1,1)
labelIntervals([moanRegionsWhale1;trillRegionWhale1])
```

```
moanRegionsWhale2 = [2.5 3.5; 5.8 8; 15.4 16.7];
```

```
setLabelValue(lss,2,'MoanRegions',moanRegionsWhale2,[true true true]);  
trillRegionWhale2 = [11.1 13];  
setLabelValue(lss,2,'TrillRegions',trillRegionWhale2,true);  
subplot(2,1,2)  
labelIntervals([moanRegionsWhale2;trillRegionWhale2])
```



Label three peaks for each trill region. For point labels, you specify the point locations and the label values. In this example, the point locations are in seconds.

```
peakLocsWhale1 = [1.553 1.626 1.7];  
peakValsWhale1 = [0.211 0.254 0.211];
```



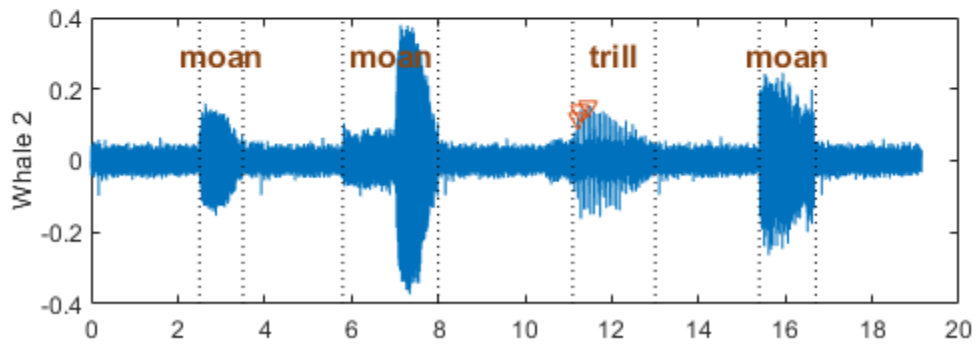
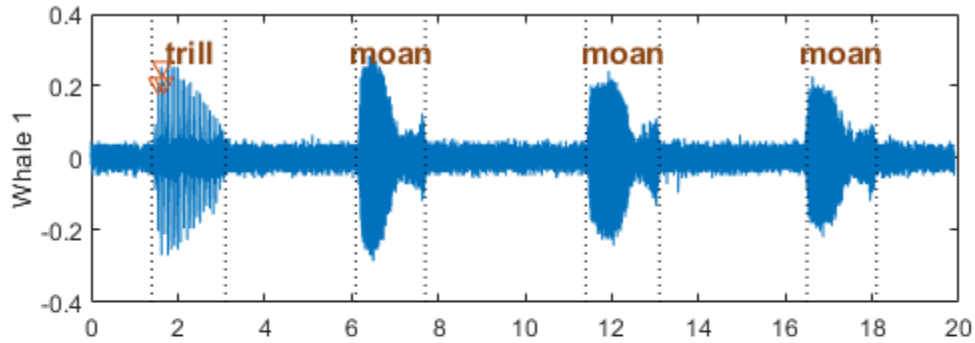
```
setLabelValue(lss,1,["TrillRegions" "TrillPeaks"],...
    peakLocsWhale1,peakValsWhale1,'LabelRowIndex',1);

subplot(2,1,1)
plot(peakLocsWhale1,peakValsWhale1,'v')
hold off

peakLocsWhale2 = [11.214 11.288 11.437];
peakValsWhale2 = [0.119 0.14 0.15];

setLabelValue(lss,2,["TrillRegions" "TrillPeaks"],...
    peakLocsWhale2,peakValsWhale2,'LabelRowIndex',1);

subplot(2,1,2)
plot(peakLocsWhale2,peakValsWhale2,'v')
hold off
```



Explore Label Values

Explore the label values using `getLabelValues`.

```
getLabelValues(lss)
```

```
ans=2x3 table
```

	WhaleType	MoanRegions	TrillRegions
Member{1}	blue	[3x2 table]	[1x3 table]
Member{2}	blue	[3x2 table]	[1x3 table]

Retrieve the moan regions for the first member of the labeled set.

```
getLabelValues(lss,1, 'MoanRegions')
```

```
ans=3x2 table
  ROIlimits      Value
  _____      _____
    6.1      7.7      [1]
   11.4     13.1      [1]
   16.5     18.1      [1]
```

Use a second output argument to list the sublabels of a label.

```
[value,valueWithSublabel] = getLabelValues(lss,1, 'TrillRegions')
```

```
value=1x2 table
  ROIlimits      Value
  _____      _____
    1.4      3.1      [1]
```

```
valueWithSublabel=1x3 table
  ROIlimits      Value      Sublabels
  _____      _____      _____
    1.4      3.1      [1]      [3x2 table]
```

To retrieve the values in a sublabel, express the label name as a two-element array.

```
getLabelValues(lss,1,["TrillRegions" "TrillPeaks"])
```

```
ans=3x2 table
  Location      Value
  _____      _____
    1.553      [0.2110]
    1.626      [0.2540]
    1.7        [0.2110]
```

Find the value of the third trill peak corresponding to the second member of the set.

```
getLabelValues(lss,2,["TrillRegions" "TrillPeaks"], ...  
    'LabelRowIndex',1,'SublabelRowIndex',3)
```

```
ans=1x2 table  
  Location      Value  
  _____  _____  
      11.437      [0.1500]
```

```
function labelIntervals(moansTrills)  
% Auxiliary function to label moan and trill regions in plots  
[X,Y] = meshgrid(moansTrills,ylim);  
plot(X,Y,'k:');  
topts = {'HorizontalAlignment','center','FontWeight','bold', ...  
    'FontSize',12,'Color',[139 69 19]/255};  
text((X(1,1:4)+X(1,5:end))/2,Y(2,5:end)-0.1, ...  
    ["moan" "moan" "moan" "trill"],topts{:})  
end
```

See Also

labeledSignalSet

Introduced in R2018b

Signal Multiresolution Analyzer


Decompose signals into time-aligned components

Description

The **Signal Multiresolution Analyzer** app is an interactive tool for visualizing multilevel wavelet and empirical mode decompositions of real-valued 1-D signals and comparing results. With the app, you can:

- Access all the signals in the MATLAB workspace.
- Adjust default parameters and generate multiple decompositions using `modwt` and `modwtmra` (default) or `emd` methods.
- Choose decomposition levels to include in the signal reconstruction.
- Visualize and compare results.
- Obtain frequency ranges of the decomposition levels. (See `powerbw` for more information.)
- Determine the relative energy of the signal across levels.
- Export reconstructed signals and decompositions to your workspace.
- Recreate the decomposition in your workspace by generating a MATLAB script.

Open the Signal Multiresolution Analyzer App

- MATLAB Toolstrip: On the **Apps** tab, under **Signal Processing and Communications**, click **Signal Multiresolution Analyzer** .
- MATLAB command prompt: Enter `signalMultiresolutionAnalyzer`.

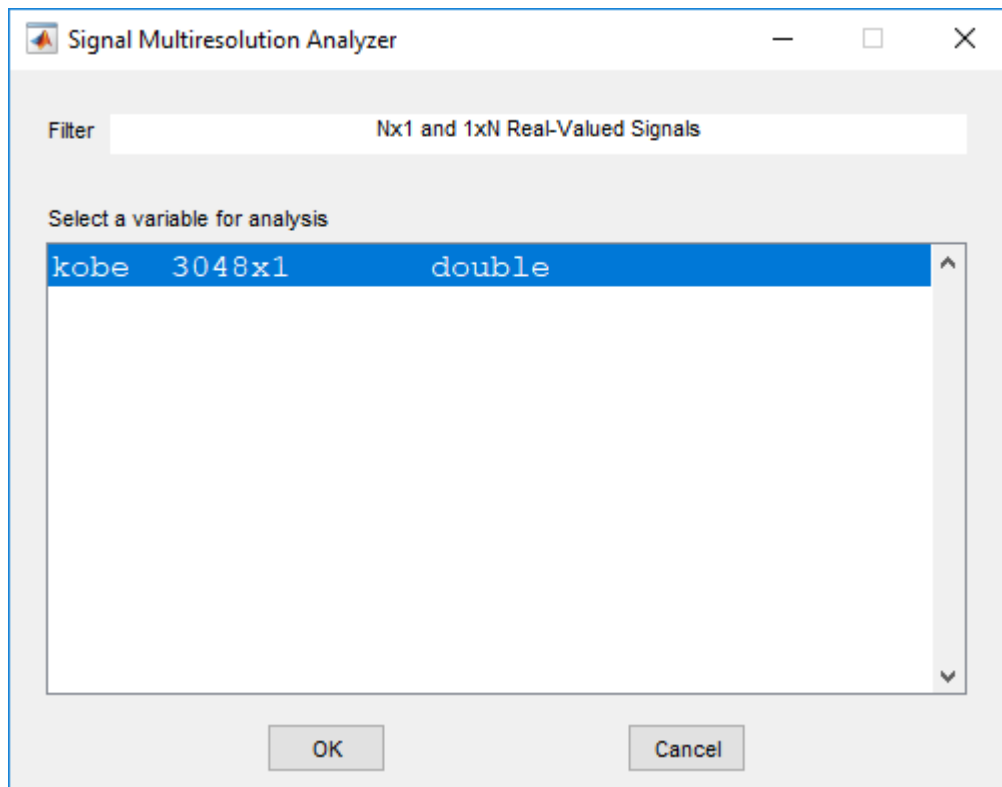
Examples

Visualize Time-Aligned Wavelet Decomposition

Load in the Kobe earthquake data. The data are seismograph measurements (vertical acceleration in nm/sec^2) recorded at Tasmania University, Hobart, Australia, on 16 January 1995, beginning at 20:56:51 (GMT) and continuing for 51 minutes at one second intervals.

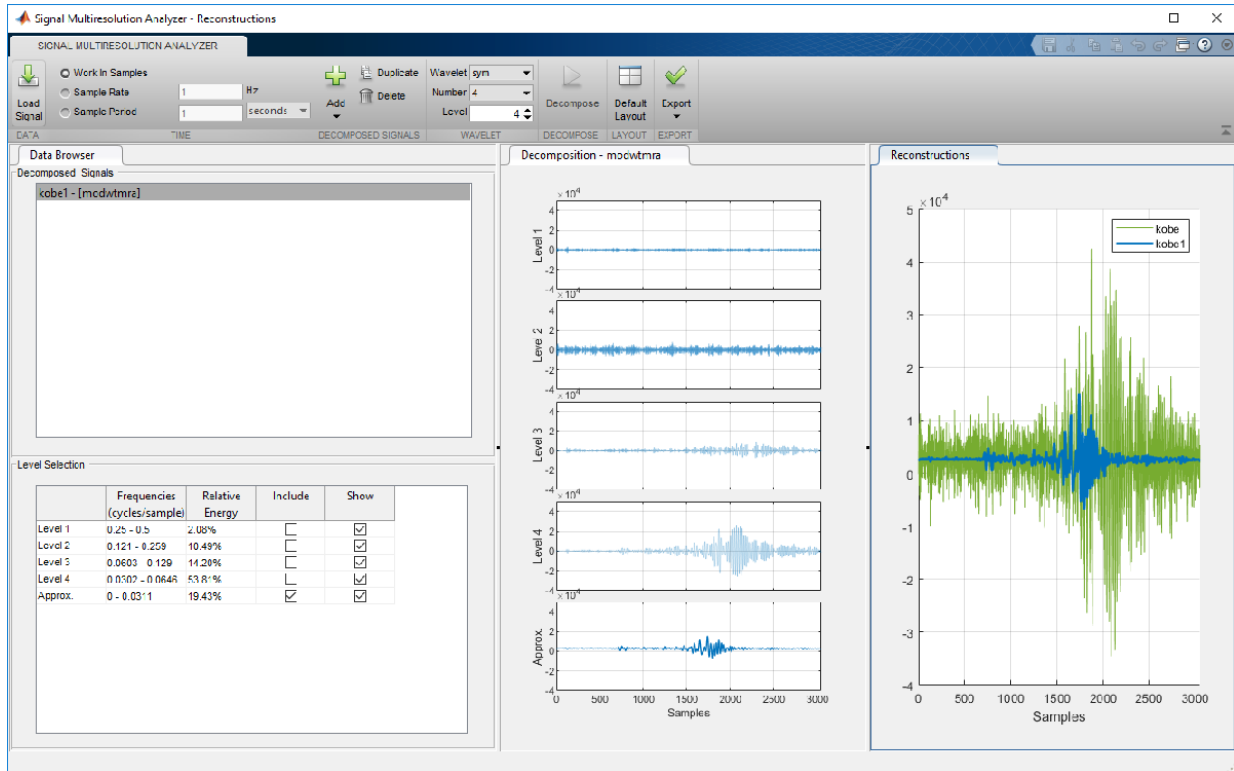
load `kobe`

Open **Signal Multiresolution Analyzer** and click **Load Signal**. A window appears listing all the workspace variables the app can process.



Select the Kobe data from the dialog box and click **OK**. A four-level MODWTMRA decomposition of the signal appears. The decomposed signal is named `kobe1` in the

Decomposed Signals pane. The suffix [modwtmra] identifies the decomposition as wavelet based.



The plots in the middle **Decomposition** pane are the projections of the wavelet decompositions of the signal at each scale on the original signal subspace. The original signal, kobe, and the reconstruction, kobe1, are plotted in the **Reconstruction** pane. The **Level Selection** pane shows the relative energies of the signal across scales, as well as the frequency bands.

Level Selection				
	Frequencies (cycles/sample)	Relative Energy	Include	Show
Level 1	0.25 - 0.5	2.08%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 2	0.121 - 0.259	10.49%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 3	0.0603 - 0.129	14.20%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 4	0.0302 - 0.0646	53.81%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Approx.	0 - 0.0311	19.43%	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

A check box in the **Show** column controls whether or not that level is displayed in the **Decomposition** pane. A check box in the **Include** column controls whether or not to include that level of the wavelet decomposition in the reconstruction. Clicking a plot in the **Decomposition** pane is another way to include or exclude that level in the signal reconstruction. To generate a new wavelet decomposition, change one of the wavelet parameters in the toolstrip and click **Decompose**.

- **Wavelet** - Wavelet family
- **Number** - Wavelet filter number
- **Level** - Wavelet decomposition level

Changing any setting in the toolstrip will enable the **Decompose** button.

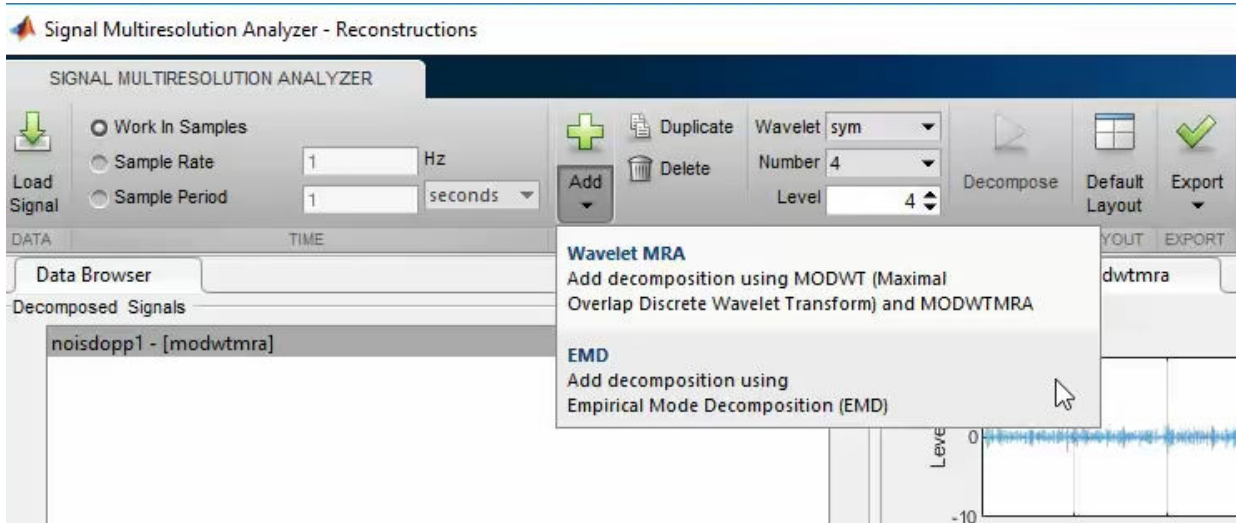
Compare MODWTMRA and EMD Decompositions

Load the noisy Doppler signal. The signal is noisy version of the Doppler test signal of Donoho and Johnstone [1].

```
load noisdopp
```

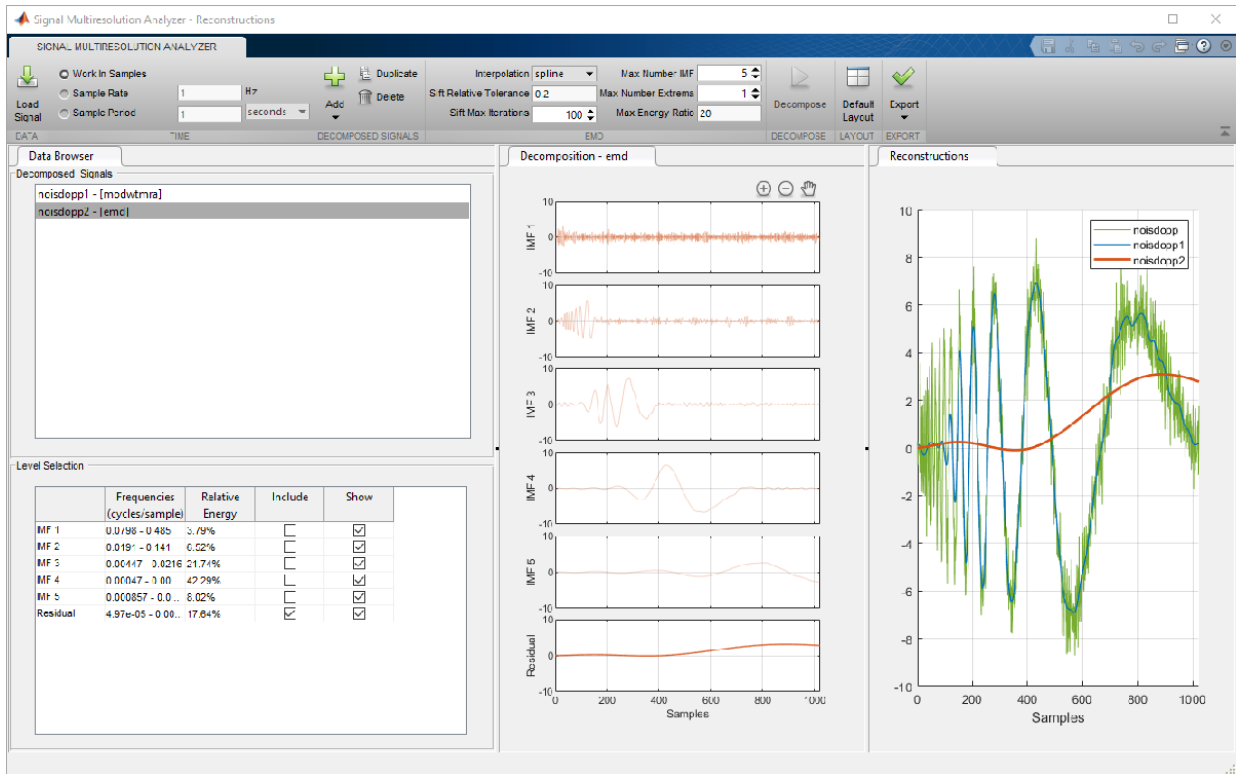
Open **Signal Multiresolution Analyzer** and load the signal into the app. By default, the app creates a four-level MODWTMRA decomposition of the signal. In the **Decomposed Signals** pane, the wavelet decomposition is named `noisdopp1`. The **Reconstruction** pane shows the original and reconstructed signals plotted in two different colors.

To add the EMD decomposition, click **Add ▼** and select **EMD**.

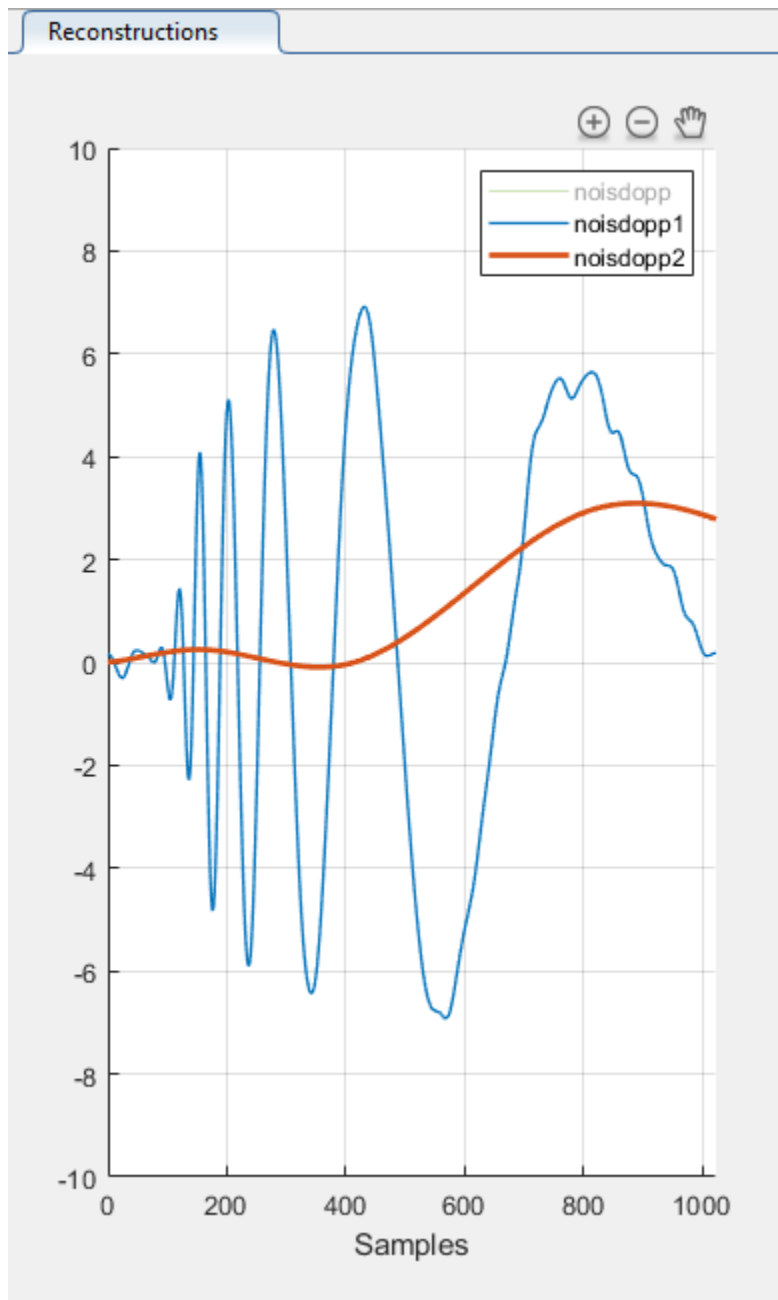


After a few moments the EMD decomposition `noisdopp2` appears in the app. Because the EMD decomposition is selected in the **Decomposed Signals** pane, the toolbar changes to show options related to EMD, and the residual is now the thickest plot in the **Reconstructions** pane.

1 Function Reference



To more easily see the differences between the two reconstructions, click `noisdopp` in the plot legend. The text fades, and the plot of the original signal is hidden. You can use the legend to hide any plot in the **Reconstruction** pane.



You can change the parameters in the toolstrip to generate a different EMD decomposition:

- **Interpolation** - Interpolation method for envelope construction
- **Sift Relative Tolerance** - Cauchy type convergence criterion
- **Sift Max Iterations** - Maximum number of sifting iterations
- **Max Number IMF** - Maximum number of IMFs extracted
- **Max Number Extrema** - Maximum number of extrema in the residual signal
- **Max Energy Ratio** - Signal to residual energy ratio

See `emd` for more information about the parameters.

Generate Decomposition Script

This example shows how to change the app default settings to duplicate a decomposition for modification, and then how to generate a script to recreate the decomposition in your workspace.

Load the Kobe earthquake data into your workspace. The data are seismograph measurements (vertical acceleration in nm/sec^2) recorded at Tasmania University, Hobart, Australia, on 16 January 1995, beginning at 20:56:51 (GMT) and continuing for 51 minutes at one second intervals.

```
load kobe
```

Open **Signal Multiresolution Analyzer** and load the earthquake data into the app. By default, the app creates a four-level MODWTMRA decomposition of the signal called `kobe1` using the order 4 Symlet `sym4`.

Create a new six-level decomposition using the order 1 Coiflet `coif1`. Click **Duplicate** in the toolstrip. Since `kobe1` is the currently selected item in **Decomposed Signals**, a duplicate of the first decomposition is created. The duplicate is called `kobe1Copy`. The plots in **Reconstruction** are updated to include the new decomposition. Except for the color, the duplicate will be identical with the first decomposition. You can change the name of the duplicate by right-clicking on the name in **Decomposed Signals**.

Change the settings in the toolstrip to the following values and then click **Decompose**.

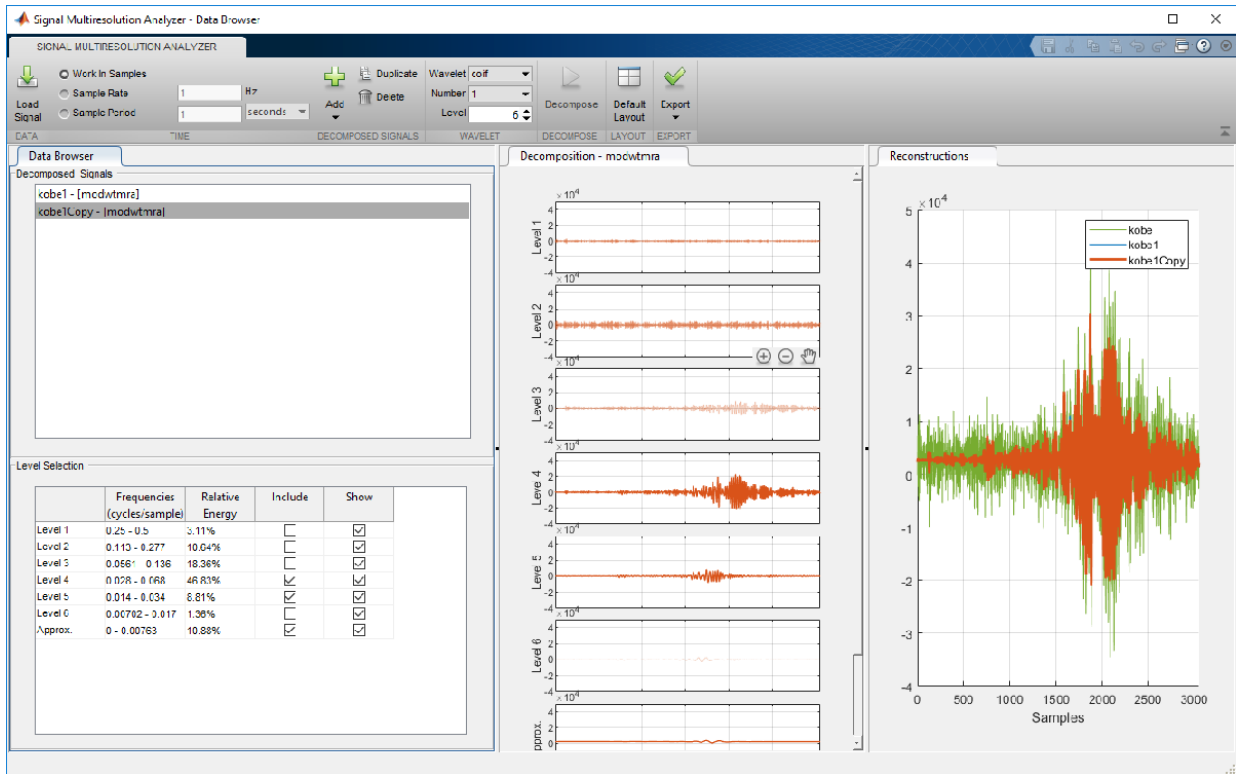
- **Wavelet:** coif
- **Number:** 1
- **Level:** 6

In **Level Selection**, note which components of the decomposition are included in the reconstruction: the approximation and the level 5 and 6 details. The signal does not have much energy at level 6, while level 4 has nearly one half the total energy. Remove Level 6 from the reconstruction, and instead include level 4.

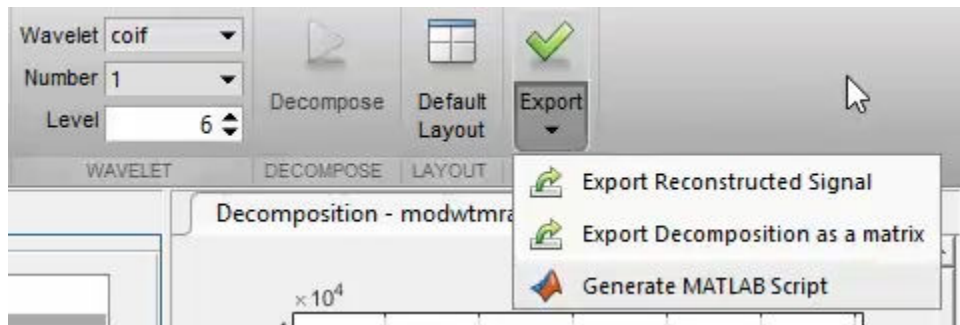
Level Selection				
	Frequencies (cycles/sample)	Relative Energy	Include	Show
Level 1	0.25 - 0.5	3.11%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 2	0.113 - 0.277	10.64%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 3	0.0561 - 0.136	18.36%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Level 4	0.028 - 0.068	46.83%	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Level 5	0.014 - 0.034	8.81%	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Level 6	0.00702 - 0.017	1.36%	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Approx.	0 - 0.00763	10.88%	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

The plot of the kobe1Copy reconstruction changes noticeably.

1 Function Reference



You have three export options. You can export the reconstruction or the entire decomposition to your workspace, or you can create a MATLAB™ script. To generate a script, click **Export > Generate MATLAB Script**.

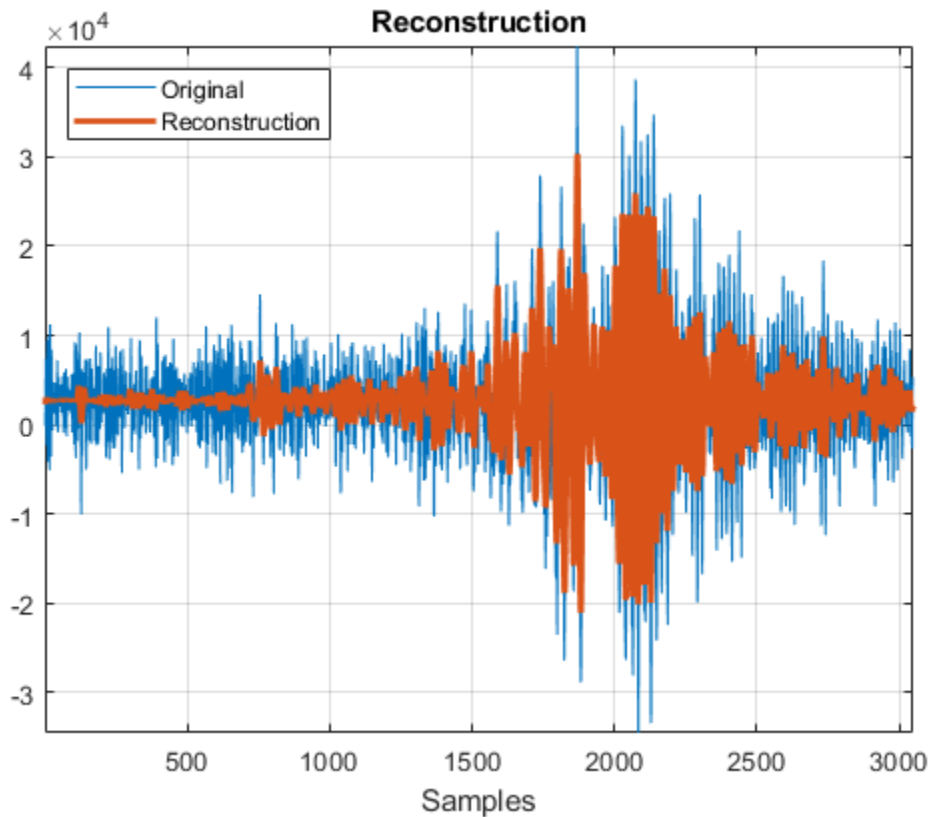


An untitled script appears in your MATLAB Editor.

```
% Decompose Signal using the MODWT
% Generated by MATLAB(R) 9.5 and Wavelet Toolbox 5.0.
% Generated on: 02-May-2018 18:41:50
% Logical array for selecting reconstruction elements
levelForReconstruction = [false, false, false, true, true, false, true];
% Perform the decomposition using modwt
wt = modwt(kobe, 'coif1', 6);
% Construct MRA matrix using modwtmra
mra = modwtmra(wt, 'coif1');
% Sum along selected multiresolution signals
kobe1Copy = sum(mra(levelForReconstruction,:),1);
```

The true-false values in `levelForReconstruction` correspond to which **Include** boxes are checked in **Level Selection**. You can save the script as is, or modify it to apply the same decomposition settings to other signals. Run the script and plot the original signal and reconstruction. Except for possibly the colors, the plot will match the `kobe1Copy` reconstruction shown in the app.

```
plot(kobe)
grid on
hold on
plot(kobe1Copy, 'LineWidth',2)
xlabel('Samples')
title('Reconstruction')
legend('Original', 'Reconstruction', 'Location', 'northwest')
axis tight
```



Parameters

Wavelet — Orthogonal wavelet family

`sym` (default) | `coif` | `db` | `fk`

Orthogonal wavelet family to use to generate the multiresolution analysis (default), specified as:

- `sym` — Symlets
- `coif` — Coiflets

- `db` — Daubechies wavelets
- `fk` — Fejér-Korovkin wavelets

The `Wavelet` parameter is applicable only for generating a multiresolution analysis.

For more information about the wavelets, use the `waveinfo` function. For example, to learn more about Daubechies wavelets, enter `waveinfo('db')`.

Interpolation — Interpolation method

`spline` (default) | `pchip`

Interpolation method to use for envelope construction in empirical mode decomposition, specified as one of the following:

- `spline` — Cubic spline interpolation
- `pchip` — Piecewise cubic Hermite interpolating polynomial method

The `Interpolation` parameter is applicable only for generating an empirical mode decomposition. You can change other options with the app when creating empirical mode decompositions. For more information, see `emd`.

Programmatic Use

`signalMultiresolutionAnalyzer` opens the **Signal Multiresolution Analyzer** app. Once the app initializes, import a signal for analysis by clicking **Load Signal**.

`signalMultiresolutionAnalyzer(sig)` opens the **Signal Multiresolution Analyzer** app and imports, decomposes, and plots the multiresolution analysis of `sig` using `modwtmra` and `modwt` with the `sym4` wavelet and default settings. `sig` is a real-valued vector.

By default, the app plots the decomposition levels as functions of sample index. To plot with respect to time, you can set a sample rate or sample period using the app.

Tips

To decompose more than one signal simultaneously, you can run multiple instances of the **Signal Multiresolution Analyzer** app.

Algorithms

The **Signal Multiresolution Analyzer** uses `modwt` and `modwtmra` to generate the multiresolution analysis and `emd` to generate the empirical mode decompositions.

References

- [1] Donoho, D. L., and I. M. Johnstone. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*. Vol. 81, 1994, pp. 425-455 1994.

See Also

Apps

Wavelet Analyzer | Wavelet Signal Denoiser

Functions

`emd` | `modwt` | `modwtmra`

Topics

"Choose a Wavelet"

"Comparing MODWT and MODWTMRA" on page 1-845

"Code Generation"

Introduced in R2018b

subset

Get new labeled signal set with subset of members

Syntax

```
lssnew = subset(lss,midxvect)
```

Description

`lssnew = subset(lss,midxvect)` returns a new labeled signal set containing the members specified in `midxvect`.

Examples

Labeled Subset

Load a labeled signal set of whale songs.

```
load whales  
lss
```

```
lss =  
  labeledSignalSet with properties:  
      Source: {2x1 cell}  
      NumMembers: 2  
      TimeInformation: "sampleRate"  
      SampleRate: 4000  
      Labels: [2x3 table]  
      Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Create a new labeled signal set consisting of the second member of the original set.

```
lssnew = subset(lss,2)
```

```
lssnew =  
  labeledSignalSet with properties:  
  
      Source: {[76579x1 double]}  
    NumMembers: 1  
  TimeInformation: "sampleRate"  
    SampleRate: 4000  
      Labels: [1x3 table]  
  Description: "Characterize wave song regions"
```

Use `labelDefinitionsHierarchy` to see a list of labels and sublabels.
Use `setLabelValue` to add data to the set.

Input Arguments

lss — Labeled signal set

`labeledSignalSet` object

Labeled signal set, specified as a `labeledSignalSet` object.

Example: `labeledSignalSet({randn(100,1)
randn(10,1)},signalLabelDefinition('female'))` specifies a two-member set of random signals containing the attribute 'female'.

midxvect — Subset member row numbers

vector of positive integers

Subset member row numbers, specified as a vector of positive integers. Each element of `midxvect` specifies a member row number as it appears in the “Labels” on page 1-0 table of the `labeledSignalSet` object `lss`.

Example: `[2 3 5 7 11 13 17]` chooses a subset of signals indexed by prime numbers.

Output Arguments

lssnew — New labeled signal set

labeledSignalSet object

New labeled signal set, returned as a labeledSignalSet object.

See Also

labeledSignalSet | signalLabelDefinition

Introduced in R2018b

swt

Discrete stationary wavelet transform 1-D

Syntax

```
SWC = swt(X,N,'wname')  
SWC = swt(X,N,Lo_D,Hi_D)  
[SWA,SWD] = swt(____)
```

Description

swt performs a multilevel 1-D stationary wavelet decomposition using either an orthogonal or a biorthogonal wavelet. Specify the wavelet using its name ('wname', see `wfilters` for more information) or its decomposition filters.

`SWC = swt(X,N,'wname')` computes the stationary wavelet decomposition of the signal `X` at level `N`, using 'wname'.

`N` must be a strictly positive integer (see `wmaxlev` for more information) and `length(X)` must be a multiple of 2^N .

`SWC = swt(X,N,Lo_D,Hi_D)` computes the stationary wavelet decomposition as above, given these filters as input:

- `Lo_D` is the decomposition low-pass filter.
- `Hi_D` is the decomposition high-pass filter.

`Lo_D` and `Hi_D` must be the same length.

The output matrix `SWC` contains the vectors of coefficients stored row-wise:

For $1 \leq i \leq N$, the output matrix `SWC(i,:)` contains the detail coefficients of level `i` and `SWC(N+1,:)` contains the approximation coefficients of level `N`.

`[SWA,SWD] = swt(____)` computes approximations, `SWA`, and details, `SWD`, stationary wavelet coefficients.

The vectors of coefficients are stored row-wise:

For $1 \leq i \leq N$, the output matrix $SWA(i, :)$ contains the approximation coefficients of level i and the output matrix $SWD(i, :)$ contains the detail coefficients of level i .

Note `swt` is defined using periodic extension. The length of the approximation and detail coefficients computed at each level equals the length of the signal.

Examples

Multilevel Stationary Wavelet Decomposition

Perform a multilevel stationary wavelet decomposition of a signal.

Load a one-dimensional signal and acquire its length.

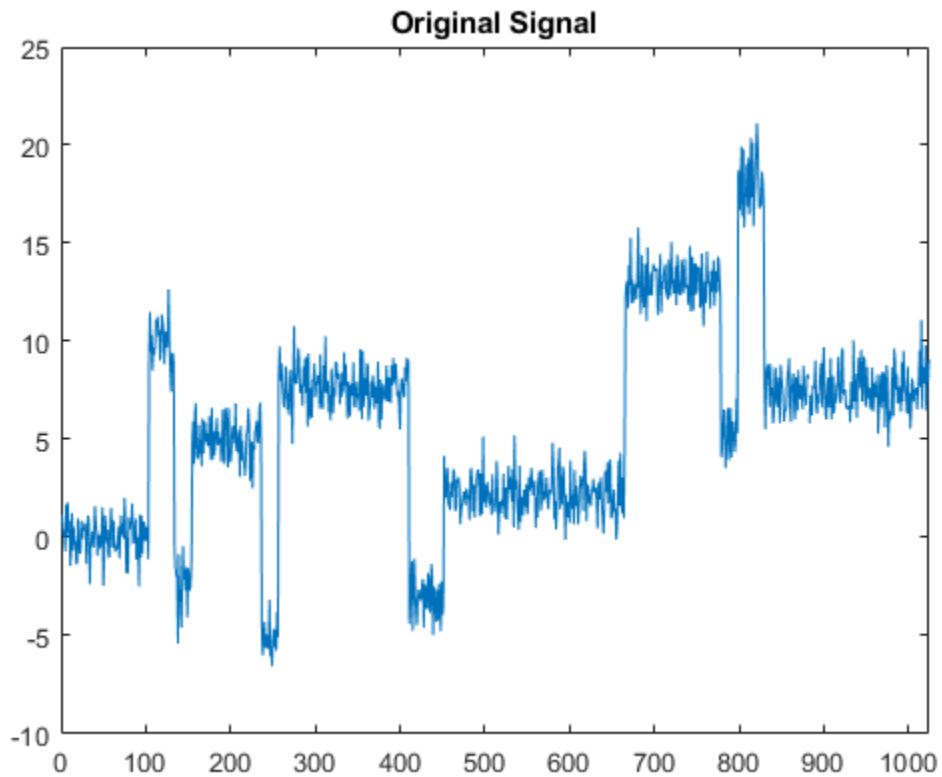
```
load noisbloc
s = noisbloc;
sLen = length(s);
```

Perform a stationary wavelet decomposition at level 3 of the signal using 'db1'. Extract the detail and approximation coefficients at level 3.

```
[swa,swd] = swt(s,3,'db1');
swd3 = swd(3,:);
swa3 = swa(3,:);
```

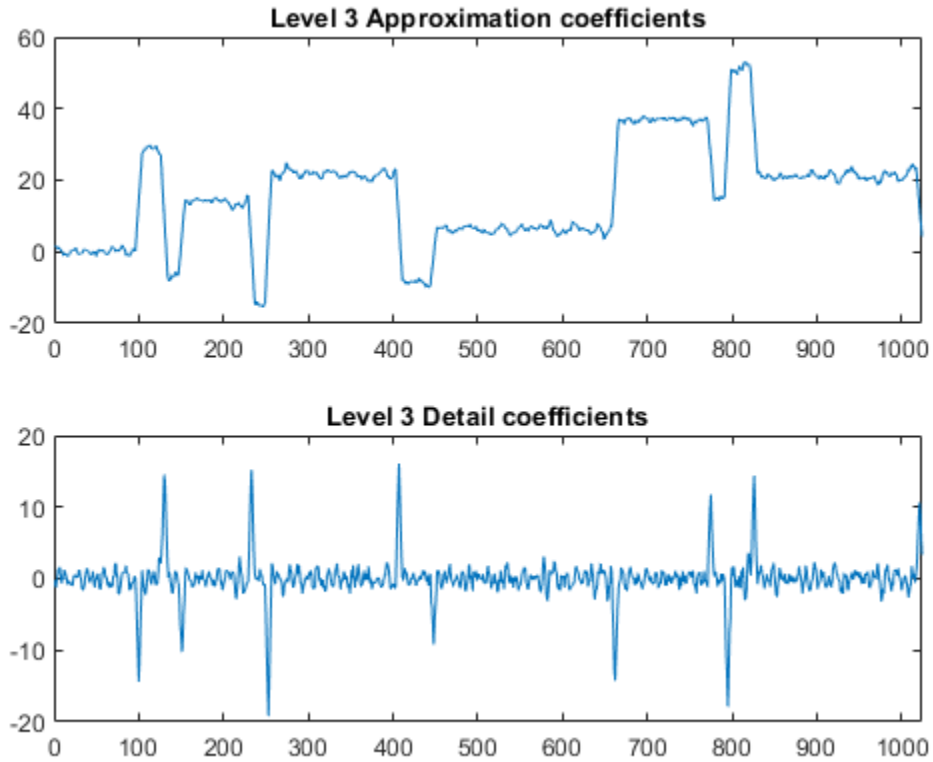
Plot the output of the decomposition.

```
plot(s)
xlim([0 sLen])
title('Original Signal')
```



Plot the level 3 approximation and detail coefficients.

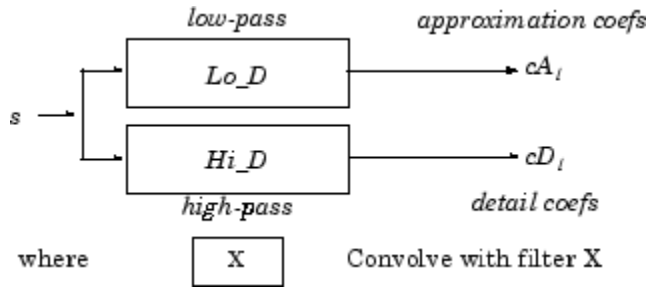
```
subplot(2,1,1)
plot(swa3)
xlim([0 sLen])
title('Level 3 Approximation coefficients')
subplot(2,1,2)
plot(swd3)
xlim([0 sLen])
title('Level 3 Detail coefficients')
```

Algorithms

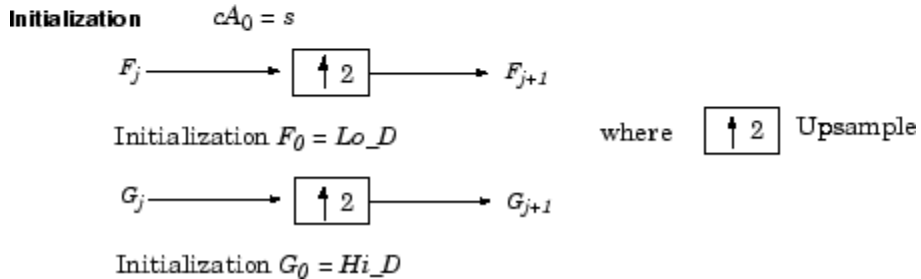
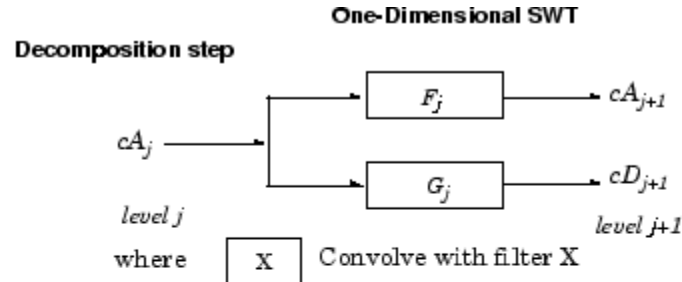
Given a signal s of length N , the first step of the SWT produces, starting from s , two sets of coefficients: approximation coefficients cA_1 and detail coefficients cD_1 . These vectors are obtained by convolving s with the low-pass filter `Lo_D` for approximation, and with the high-pass filter `Hi_D` for detail.

More precisely, the first step is



Note cA_1 and cD_1 are of length N instead of $N/2$ as in the DWT case.

The next step splits the approximation coefficients cA_1 in two parts using the same scheme, but with modified filters obtained by upsampling the filters used for the previous step and replacing s by cA_1 . Then, the SWT produces cA_2 and cD_2 . More generally,



References

Nason, G.P.; B.W. Silverman (1995), "The stationary wavelet transform and some statistical applications," *Lecture Notes in Statistics*, 103, pp. 281-299.

Coifman, R.R.; Donoho, D.L. (1995), "Translation invariant de-noising," *Lecture Notes in Statistics*, 103, pp. 125-150.

Pesquet, J.C.; H. Krim, H. Carfatan (1996), "Time-invariant orthonormal wavelet representations," *IEEE Trans. Sign. Proc.*, vol. 44, 8, pp. 1964-1970.

See Also

dwt | iswt | modwt | wavedec

Introduced before R2006a

swt2

Discrete stationary wavelet transform 2-D

Syntax

```
SWC = swt2(X,N,'wname')  
[A,H,V,D] = swt2(X,N,'wname')  
SWC = swt2(X,N,Lo_D,Hi_D)  
[A,H,V,D] = swt2(X,N,Lo_D,Hi_D)
```

Description

swt2 performs a multilevel 2-D stationary wavelet decomposition using either an orthogonal or a biorthogonal wavelet. Specify the wavelet using its name('wname', see wfilters for more information) or its decomposition filters.

SWC = swt2(X,N,'wname') or [A,H,V,D] = swt2(X,N,'wname') compute the stationary wavelet decomposition of the real-valued 2-D or 3-D matrix X at level N, using 'wname'.

If X is a 3-D matrix, the third dimension of X must equal 3.

N must be a strictly positive integer (see wmaxlev for more information), and 2^N must divide size(X,1) and size(X,2).

The dimension of X and level N determine the dimensions of the outputs.

- If X is a 2-D matrix and N is greater than 1, the outputs [A,H,V,D] are 3-D arrays, which contain the coefficients:
 - For $1 \leq i \leq N$, the output matrix $A(:, :, i)$ contains the coefficients of approximation of level i .
 - The output matrices $H(:, :, i)$, $V(:, :, i)$ and $D(:, :, i)$ contain the coefficients of details of level i (horizontal, vertical, and diagonal):

```
SWC = [H(:, :, 1:N) ; V(:, :, 1:N) ; D(:, :, 1:N) ; A(:, :, N)]
```

- If X is a 2-D matrix and N is equal to 1, the outputs $[A, H, V, D]$ are 2-D arrays where A contains the approximation coefficients, and H , V , and D contain the horizontal, vertical, and diagonal detail coefficients, respectively.
- If X is a 3-D matrix of dimension m -by- n -by-3, and N is greater than 1, the outputs $[A, H, V, D]$ are 4-D arrays of dimension m -by- n -by-3-by- N , which contain the coefficients:
 - For $1 \leq i \leq N$ and $j = 1, 2, 3$, the output matrix $A(:, :, j, i)$ contains the coefficients of approximation of level i .
 - The output matrices $H(:, :, j, i)$, $V(:, :, j, i)$ and $D(:, :, j, i)$ contain the coefficients of details of level i (horizontal, vertical, and diagonal):

$$SWC = [H(:, :, 1:3, 1:N) ; V(:, :, 1:3, 1:N) ; D(:, :, 1:3, 1:N) ; A(:, :, 1:3, N)]$$

- If X is a 3-D matrix of dimension m -by- n -by-3, and N is equal to 1, the outputs $[A, H, V, D]$ are 4-D arrays of dimension m -by- n -by-1-by-3, which contain the coefficients:
 - For $j = 1, 2, 3$, the output matrix $A(:, :, 1, j)$ contains the approximation coefficients.
 - The output matrices $H(:, :, 1, j)$, $V(:, :, 1, j)$ and $D(:, :, 1, j)$ contain the horizontal, vertical, and diagonal detail coefficients, respectively.

$$SWC = [H(:, :, 1, 1:3) ; V(:, :, 1, 1:3) ; D(:, :, 1, 1:3) ; A(:, :, 1, 1:3)]$$

Note

- `swt2` uses double-precision arithmetic internally and returns double-precision coefficient matrices. `swt2` warns if there is a loss of precision when converting to double.
- To distinguish a single-level decomposition of a truecolor image from a multilevel decomposition of an indexed image, the approximation and detail coefficient arrays of truecolor images are 4-D arrays. See “Distinguish Single-Level Truecolor Image from Multilevel Indexed Image Decompositions” on page 1-1151. Also see examples “Stationary Wavelet Transform of an Image” on page 1-1140 and “Inverse Stationary Wavelet Transform of an Image” on page 1-1145.

If an K -level decomposition is performed, the dimensions of the A , H , V , and D coefficient arrays are m -by- n -by-3-by- K .

If a single-level decomposition is performed, the dimensions of the A, H, V, and D coefficient arrays are m-by-n-by-1-by-3. Since MATLAB removes singleton last dimensions by default, the third dimension of the coefficient arrays is singleton.

`SWC = swt2(X,N,Lo_D,Hi_D)` or `[A,H,V,D] = swt2(X,N,Lo_D,Hi_D)`, computes the stationary wavelet decomposition as in the previous syntax, given these filters as input:

- `Lo_D` is the decomposition low-pass filter.
- `Hi_D` is the decomposition high-pass filter.

`Lo_D` and `Hi_D` must be the same length.

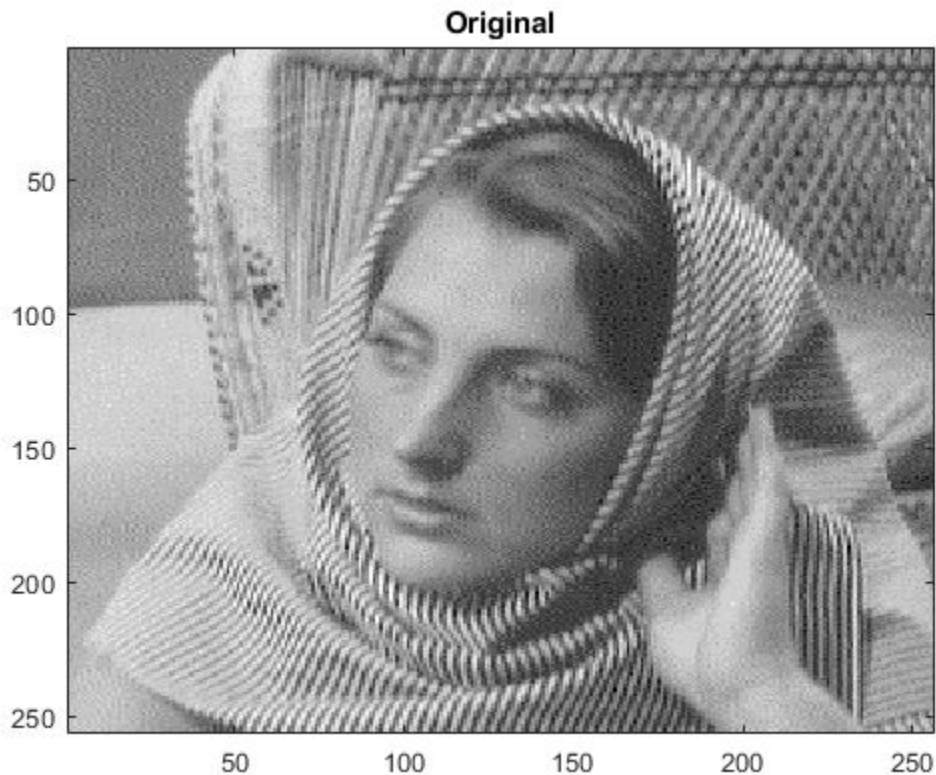
Note `swt2` is defined using periodic extension. The size of the approximation and details coefficients computed at each level equals the size of the input data.

Examples

Extract and Display Two-Dimensional Stationary Wavelet Decomposition

Extract and display images of the stationary wavelet decomposition level coefficients. First load and display the original image. Then perform the stationary wavelet decomposition of the image at level 2 using `db6`.

```
load woman
imagesc(X)
colormap(map)
title('Original')
```



```
[ca, chd, cvd, cdd] = swt2(X, 2, 'db6');
```

Extract the Level 1 and Level 2 approximation and detail coefficients from the decomposition.

```
A1 = wcodemat(ca(:, :, 1), 255);
H1 = wcodemat(chd(:, :, 1), 255);
V1 = wcodemat(cvd(:, :, 1), 255);
D1 = wcodemat(cdd(:, :, 1), 255);
```

```
A2 = wcodemat(ca(:, :, 2), 255);
H2 = wcodemat(chd(:, :, 2), 255);
V2 = wcodemat(cvd(:, :, 2), 255);
D2 = wcodemat(cdd(:, :, 2), 255);
```

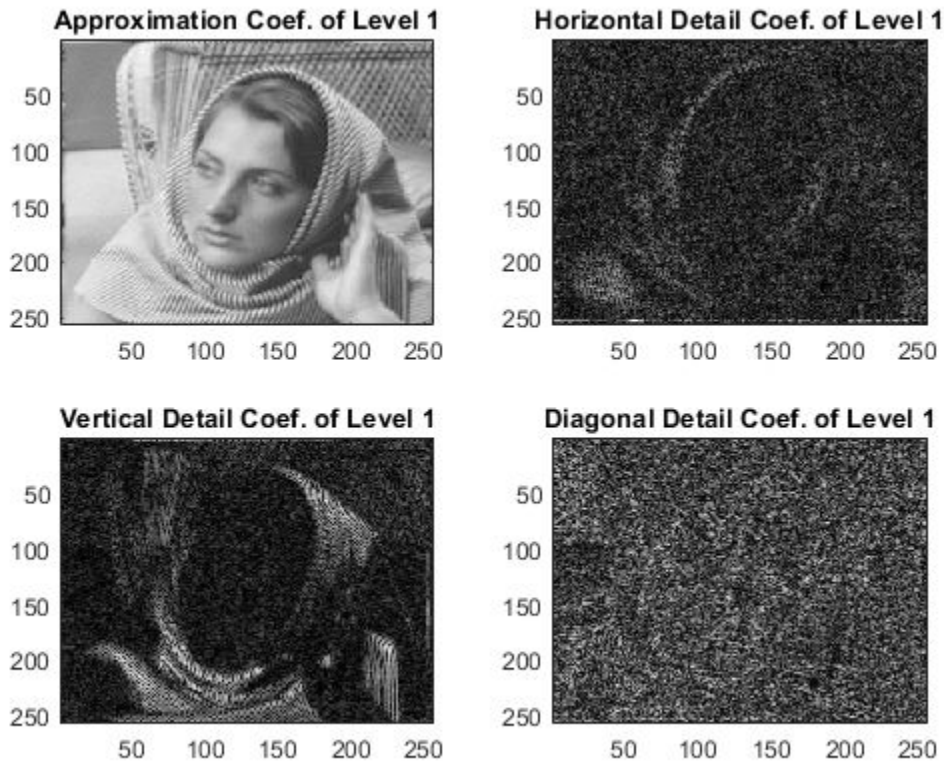
Display the approximation and detail coefficients from the two levels.

```
subplot(2,2,1)
imagesc(A1)
title('Approximation Coef. of Level 1')

subplot(2,2,2)
imagesc(H1)
title('Horizontal Detail Coef. of Level 1')

subplot(2,2,3)
imagesc(V1)
title('Vertical Detail Coef. of Level 1')

subplot(2,2,4)
imagesc(D1)
title('Diagonal Detail Coef. of Level 1')
```

```

subplot(2,2,1)
imagesc(A2)
title('Approximation Coef. of Level 2')

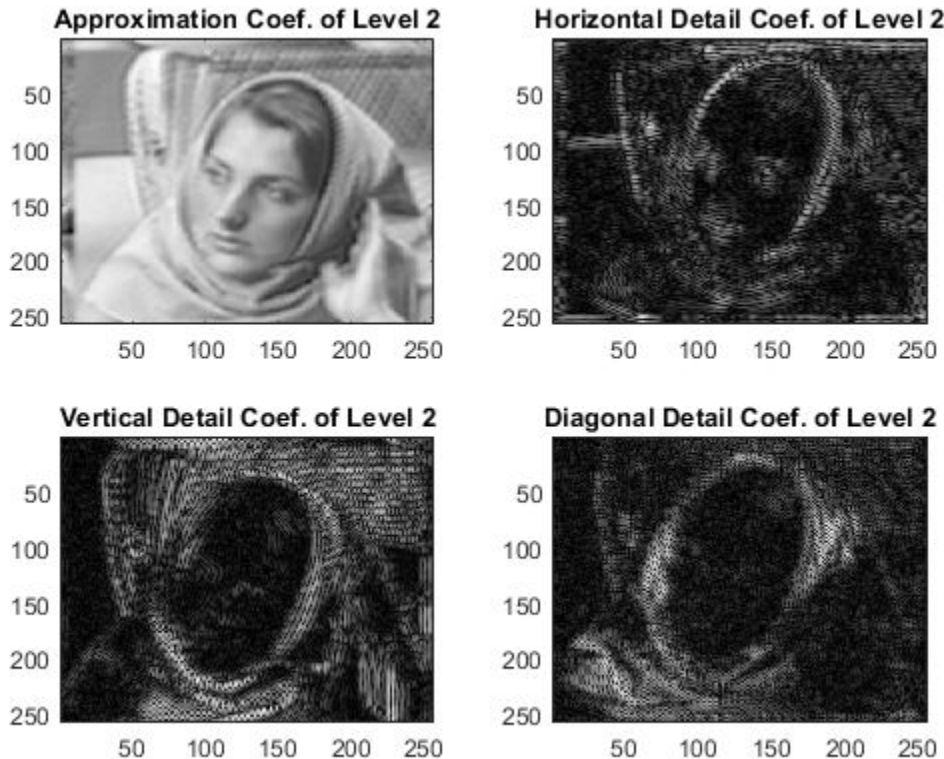
subplot(2,2,2)
imagesc(H2)
title('Horizontal Detail Coef. of Level 2')

subplot(2,2,3)
imagesc(V2)
title('Vertical Detail Coef. of Level 2')

subplot(2,2,4)

```

```
imagesc(D2)  
title('Diagonal Detail Coef. of Level 2')
```

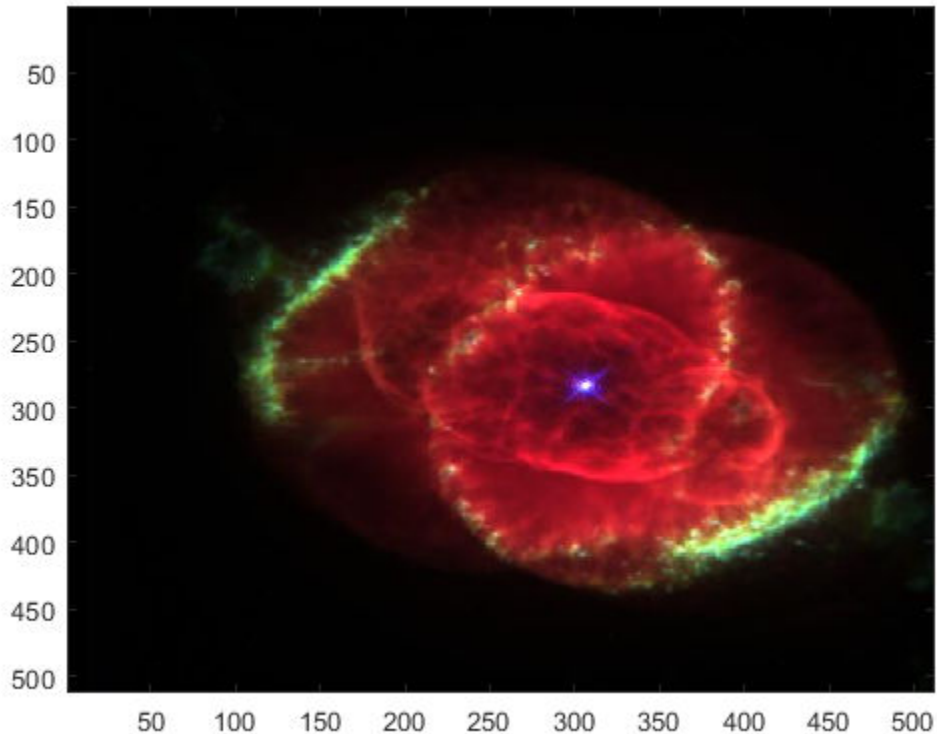


Stationary Wavelet Transform of an Image

In this example you obtain single-level and multilevel stationary wavelet decompositions of a truecolor image. You view results of each decomposition.

Load in a truecolor image. The image is a 3-D array of type `uint8`. Since `swt2` requires the first and second dimensions both be divisible by a power of 2, extract a portion of the image and view it.

```
imdata = imread('ngc6543a.jpg');  
x = imdata(1:512,1:512,:);  
imagesc(x)
```



Obtain the 4-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and the horizontal, vertical, and detail coefficients as separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,4,'db4');  
size(a)
```

```
ans = 1x4
```

```
512 512 3 4

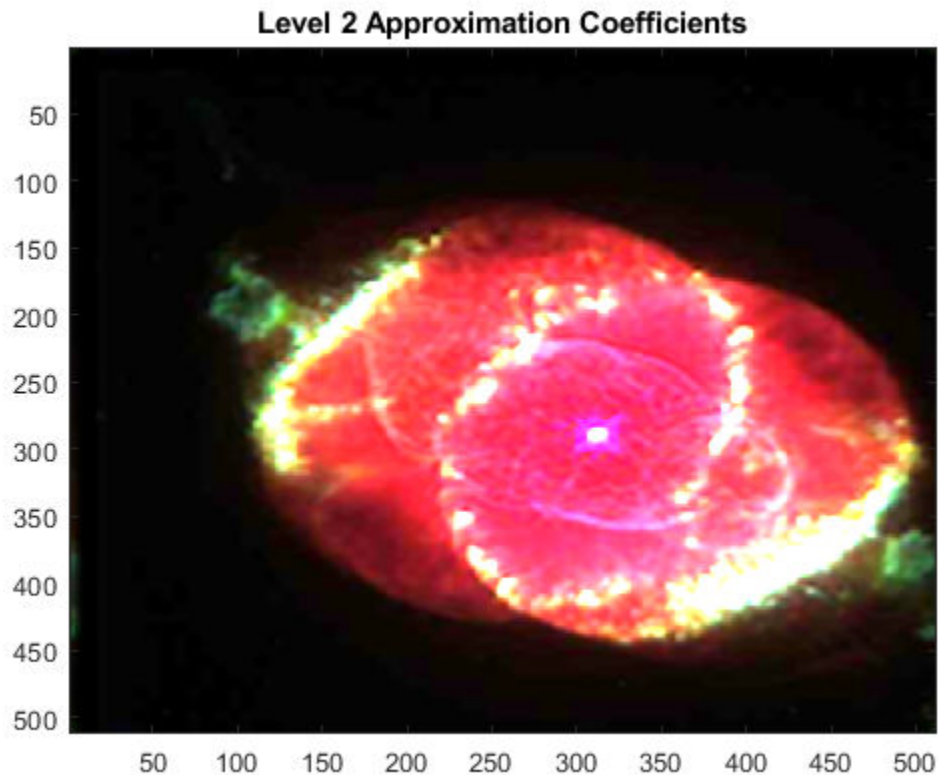
size(h)
ans = 1x4
512 512 3 4

size(v)
ans = 1x4
512 512 3 4

size(d)
ans = 1x4
512 512 3 4
```

The output arrays are all of type `double`. View the level 2 approximation coefficients. Since the approximation coefficients are of type `double`, cast them as `uint8`, which is the datatype of the image.

```
figure
imagesc(uint8(a(:,:, :, 2)))
title('Level 2 Approximation Coefficients')
```



Reconstruct the image by performing the inverse transform. Compute the difference between the original image and reconstruction. Since the reconstruction is of type double, cast the reconstruction as type uint8 before computing the difference.

```
rec = iswt2(a,h,v,d,'db4');
maxdiff = max(abs(uint8(rec(:))-x(:)));
disp(['maximum difference = ' num2str(maxdiff)])
```

```
maximum difference = 0
```

Obtain the single-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and horizontal, vertical, and detail coefficients in separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,1, 'db4');  
size(a)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(h)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(v)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(d)
```

```
ans = 1×4
```

```
512 512 1 3
```

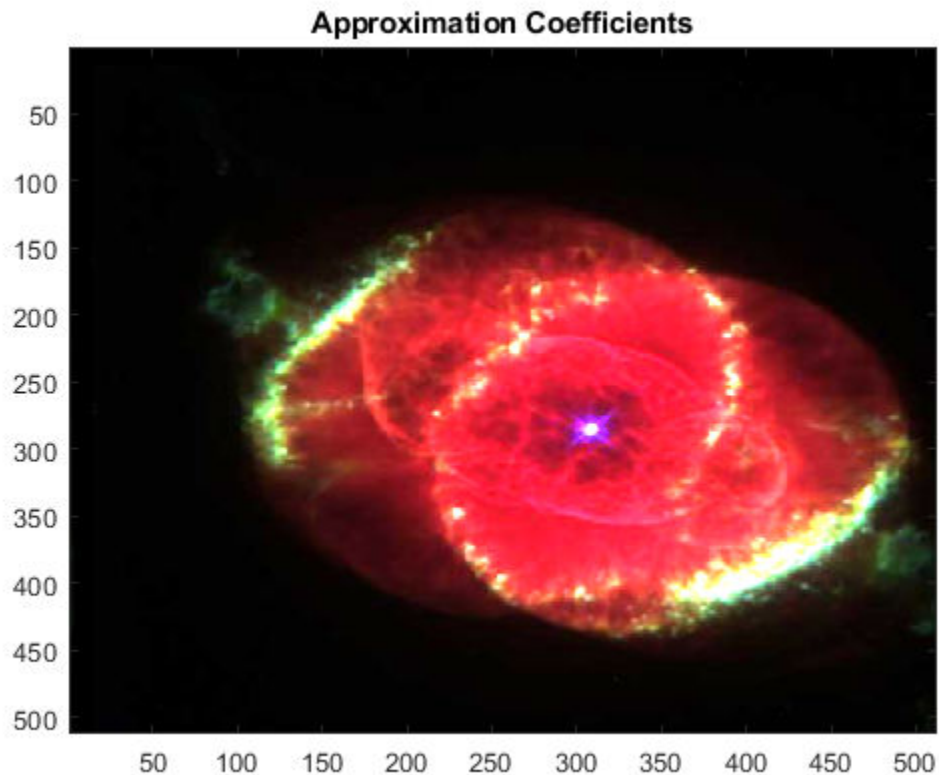
View the approximation coefficients. To prevent an error when using `imagesc`, squeeze the approximation coefficients array to remove the singleton dimension.

```
asqueeze = squeeze(a);  
size(asqueeze)
```

```
ans = 1×3
```

```
512 512 3
```

```
figure  
imagesc(uint8(asqueeze))  
title('Approximation Coefficients')
```



Reconstruct the image by performing the inverse transform. Compute the difference between the original image and reconstruction. Since the reconstruction is of type double, cast the reconstruction as type uint8 before computing the difference.

```
rec = iswt2(a,h,v,d,'db4');  
maxdiff = max(abs(uint8(rec(:))-x(:)));  
disp(['maximum difference = ' num2str(maxdiff)])
```

```
maximum difference = 0
```

Inverse Stationary Wavelet Transform of an Image

This example shows how to reconstruct a truecolor image from a single-level stationary wavelet decomposition using 3-D approximation and detail coefficient arrays.

Load in a truecolor image. The image is a 3-D array of type `uint8`. Since `swt2` requires the first and second dimensions both be divisible by a power of 2, extract a portion of the image and view it.

```
imdata = imread('ngc6543a.jpg');  
x = imdata(1:512,1:512,:);  
imagesc(x)
```



Obtain the single-level stationary wavelet decomposition of the image using the db4 wavelet. Return the approximation coefficients and horizontal, vertical, and detail coefficients in separate arrays. Note the dimensions of the output arrays.

```
[a,h,v,d] = swt2(x,1,'db4');  
size(a)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(h)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(v)
```

```
ans = 1×4
```

```
512 512 1 3
```

```
size(d)
```

```
ans = 1×4
```

```
512 512 1 3
```

Squeeze the coefficient arrays to remove their singleton dimensions. Note the dimensions of the squeezed arrays.

```
asq = squeeze(a);  
hsq = squeeze(h);  
vsq = squeeze(v);  
dsq = squeeze(d);  
size(asq)
```

```
ans = 1×3
```

```
512 512 3
```

```
size(hsq)
ans = 1×3
    512    512     3

size(vsq)
ans = 1×3
    512    512     3

size(dsq)
ans = 1×3
    512    512     3
```

So that `iswt2` can properly reconstruct the trueimage from the new coefficient arrays, insert a singleton dimension by reshaping the squeezed arrays. Reconstruct the image with the reshaped coefficient arrays.

```
a2 = reshape(asq,[512,512,1,3]);
h2 = reshape(hsq,[512,512,1,3]);
v2 = reshape(vsq,[512,512,1,3]);
d2 = reshape(dsq,[512,512,1,3]);
rec = iswt2(a2,h2,v2,d2,'db4');
```

Compute the difference between the original image and reconstruction. Since the reconstruction is of type `double`, cast the reconstruction as type `uint8` before computing the difference.

```
maxdiff = max(abs(uint8(rec(:))-x(:)));
disp(['maximum difference = ' num2str(maxdiff)])

maximum difference = 0
```

Tips

When `X` represents an indexed image, `X` is an `m`-by-`n` matrix. If the level of decomposition `N` is greater than 1, the output arrays `SWC` or `cA`, `cH`, `cV`, and `cD` are `m`-by-`n`-by-`N` arrays. If

the level of decomposition N is equal to 1, the output arrays SWC or cA, cH, cV, and cD are m -by- n arrays.

When X represents a truecolor image, it becomes an m -by- n -by-3 array. This array is an m -by- n -by-3 array, where each m -by- n matrix represents a red, green, or blue color plane concatenated along the third dimension. If the level of decomposition N is greater than 1, the output arrays SWC or cA, cH, cV, and cD are m -by- n -by-3-by- N . If the level of decomposition N is equal to 1, the output arrays SWC or cA, cH, cV, and cD are m -by- n -by-1-by-3.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Algorithms

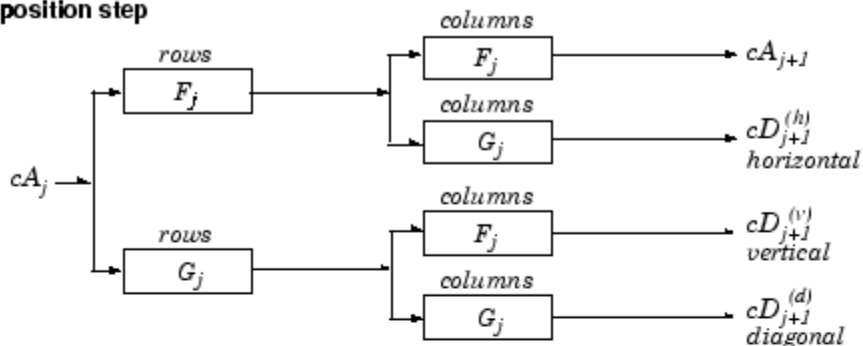
For images, an algorithm similar to the one-dimensional case is possible for two-dimensional wavelets and scaling functions obtained from one-dimensional ones by tensor product.

This kind of two-dimensional SWT leads to a decomposition of approximation coefficients at level j in four components: the approximation at level $j+1$, and the details in three orientations (horizontal, vertical, and diagonal).

The following chart describes the basic decomposition step for images:

Two-Dimensional SWT

Decomposition step



where

$\begin{matrix} \text{rows} \\ \boxed{X} \end{matrix}$ Convolve with filter X the rows of the entry

$\begin{matrix} \text{columns} \\ \boxed{X} \end{matrix}$ Convolve with filter X the columns of the entry

Initialization $cA_0 = s$ for the decomposition initialization

$$F_j \longrightarrow \boxed{\uparrow 2} \longrightarrow F_{j+1}$$

Initialization $F_0 = Lo_D$

where $\boxed{\uparrow 2}$ Upsample

$$G_j \longrightarrow \boxed{\uparrow 2} \longrightarrow G_{j+1}$$

Initialization $G_0 = Hi_D$

Note $size(cA_j) = size(cD_j^{(h)}) = size(cD_j^{(v)}) = size(cD_j^{(d)}) = s$

where $s = size\ of\ the\ analyzed\ image$

Compatibility Considerations

Distinguish Single-Level Truecolor Image from Multilevel Indexed Image Decompositions

Behavior changed in R2017b

To distinguish a single-level decomposition of a truecolor image from a multilevel decomposition of an indexed image, the approximation and detail coefficient arrays of truecolor images are 4-D arrays.

Migrate from Previous Releases to R2017b

Depending on the original input data type and level of wavelet decomposition, you might have to take different steps to make swt2 coefficient arrays from previous releases compatible with R2017b coefficient arrays. The steps depend on whether you have a single coefficient array or separate approximation and detail coefficient arrays.

Single Coefficient Array	Multiple Coefficient Arrays
Input: Index image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues 	Input: Index image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues

Single Coefficient Array	Multiple Coefficient Arrays
<p>Input: Truecolor image</p> <ul style="list-style-type: none"> Single-level: If <code>swc</code> is the output of <code>swt2</code> from a previous release, execute: <code>swc1 = double(swc);</code> Multi-level: If <code>swc</code> is the output of <code>swt2</code> from a previous release, execute: <code>swc1 = double(swc);</code> 	<p>Input: Truecolor image</p> <ul style="list-style-type: none"> Single-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from a previous release, execute: <code>ca1 = double(ca);</code> <code>chd1 = double(chd);</code> <code>cvd1 = double(cvd);</code> <code>cdd1 = double(cdd);</code> <code>ca2 = reshape(ca1, [m,n,1,3]);</code> <code>chd2 = reshape(chd1, [m,n,1,3]);</code> <code>cvd2 = reshape(cvd1, [m,n,1,3]);</code> <code>cdd2 = reshape(cdd1, [m,n,1,3]);</code> Multi-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from a previous release, execute: <code>ca1 = double(ca);</code> <code>chd1 = double(chd);</code> <code>cvd1 = double(cvd);</code> <code>cdd1 = double(cdd);</code>

Migrate from R2017b to Previous Releases

Depending on the original input data type and level of wavelet decomposition, you might have to take different steps to make R2017b `swt2` coefficient arrays compatible with the coefficient arrays from previous releases. The steps depend on whether you have a single coefficient array or separate approximation and detail coefficient arrays.

Single Coefficient Array	Multiple Coefficient Arrays
<p>Input: Index image</p> <ul style="list-style-type: none"> Single-level: No compatibility issues Multi-level: No compatibility issues 	<p>Input: Index image</p> <ul style="list-style-type: none"> Single-level: No compatibility issues Multi-level: No compatibility issues

Single Coefficient Array	Multiple Coefficient Arrays
Input: Truecolor image <ul style="list-style-type: none"> • Single-level: No compatibility issues • Multi-level: No compatibility issues 	Input: Truecolor image <ul style="list-style-type: none"> • Single-level: If <code>ca</code>, <code>chd</code>, <code>cvd</code>, and <code>cdd</code> are outputs of <code>swt2</code> from R2017b, execute: <pre>ca1 = single(squeeze(ca)); chd1 = single(squeeze(chd)); cvd1 = single(squeeze(cvd)); cdd1 = single(squeeze(cdd));</pre> • Multi-level: No compatibility issues

References

Nason, G.P.; B.W. Silverman (1995), "The stationary wavelet transform and some statistical applications," *Lecture Notes in Statistics*, 103, pp. 281-299.

Coifman, R.R.; Donoho, D.L. (1995), "Translation invariant de-noising," *Lecture Notes in Statistics*, 103, pp. 125-150.

Pesquet, J.C.; H. Krim, H. Carfatan (1996), "Time-invariant orthonormal wavelet representations," *IEEE Trans. Sign. Proc.*, vol. 44, 8, pp. 1964-1970.

See Also

`dwt2` | `iswt2` | `wavedec2`

Introduced before R2006a

symaux

Symlet wavelet filter computation

The `symaux` function generates the scaling filter coefficients for the "least asymmetric" Daubechies wavelets.

Syntax

```
w = symaux(n)
w = symaux( ____, sumw)
```

Description

`w = symaux(n)` is the order n Symlet scaling filter such that $\text{sum}(w) = 1$.

Note

- Instability may occur when n is too large. Starting with values of n in the 30s range, function output will no longer accurately represent scaling filter coefficients.
 - As n increases, the time required to compute the filter coefficients rapidly grows.
-

`w = symaux(____, sumw)` is the order n Symlet scaling filter such that $\text{sum}(w) = \text{sumw}$.

`w = symaux(n, 0)` is equivalent to `w = symaux(n, 1)`.

Examples

Unit Norm Scaling Filter Coefficients

In this example you will generate symlet scaling filter coefficients whose norm is equal to 1. You will also confirm the coefficients satisfy a necessary relation. This example requires the Signal Processing Toolbox™.

Compute the scaling filter coefficients of the order 10 symlet whose sum equals $\sqrt{2}$.

```
n = 10;  
w = symaux(n,sqrt(2));
```

Confirm the sum of the coefficients is equal to $\sqrt{2}$ and the norm is equal to 1.

```
sqrt(2)-sum(w)
```

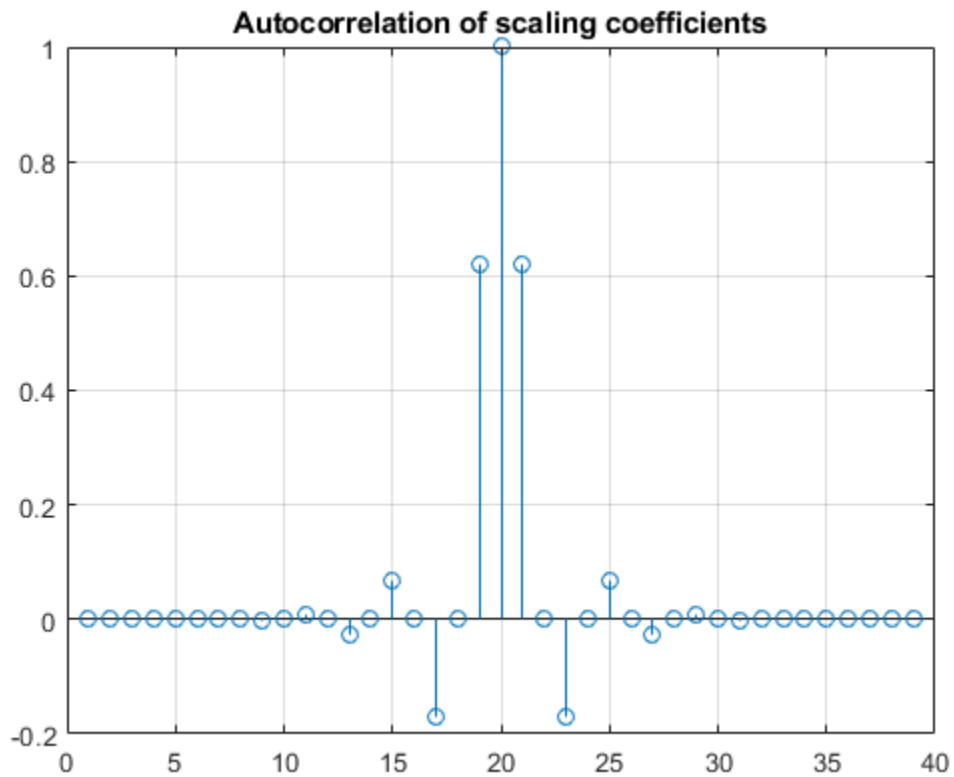
```
ans = 0
```

```
1-sum(w.^2)
```

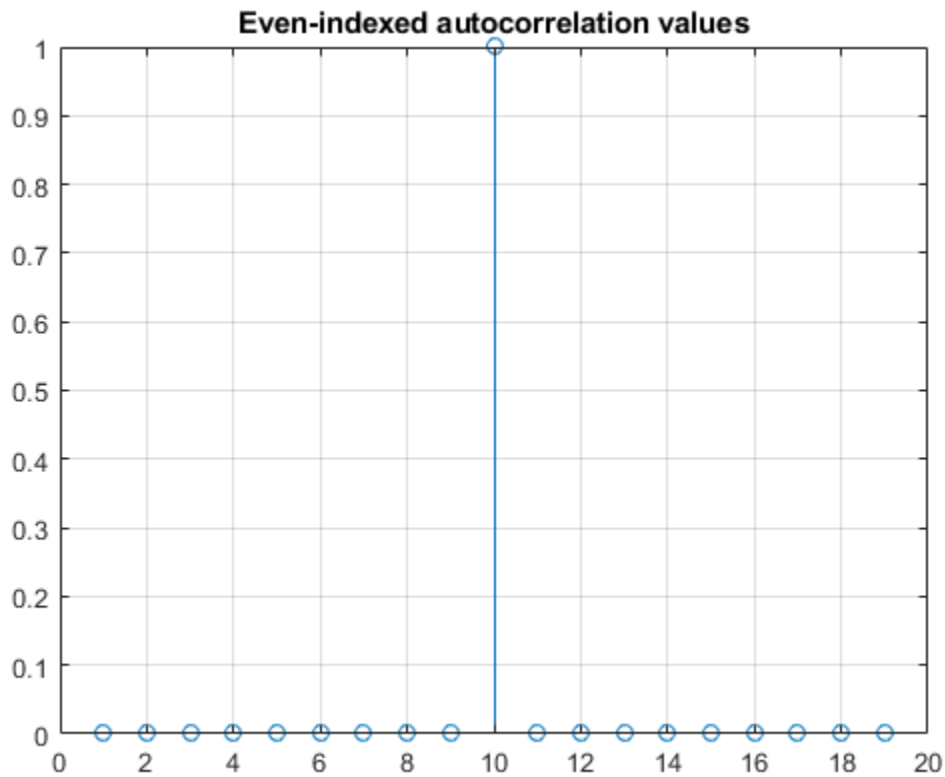
```
ans = 1.1324e-14
```

Since integer translations of the scaling function form an orthogonal basis, the coefficients satisfy the relation $\sum_n w(n)w(n-2k) = \delta(k)$. Confirm this by taking the autocorrelation of the coefficients and plotting the result.

```
corrw = xcorr(w,w);  
stem(corrw)  
grid on  
title('Autocorrelation of scaling coefficients')
```



```
stem(corrw(2:2:end))  
grid on  
title('Even-indexed autocorrelation values')
```



Symlet and Daubechies Scaling Filters

This example shows that symlet and Daubechies scaling filters of the same order are both solutions of the same polynomial equation.

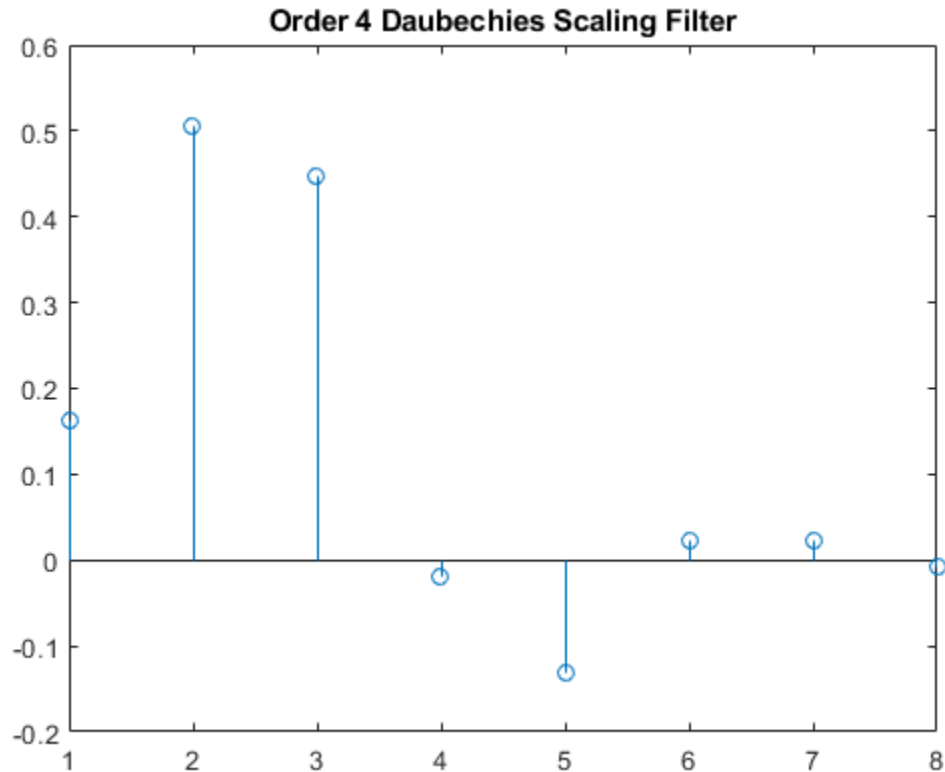
Generate the order 4 Daubechies scaling filter and plot it.

```
wdb4 = dbaux(4)
```

```
wdb4 = 1×8
```

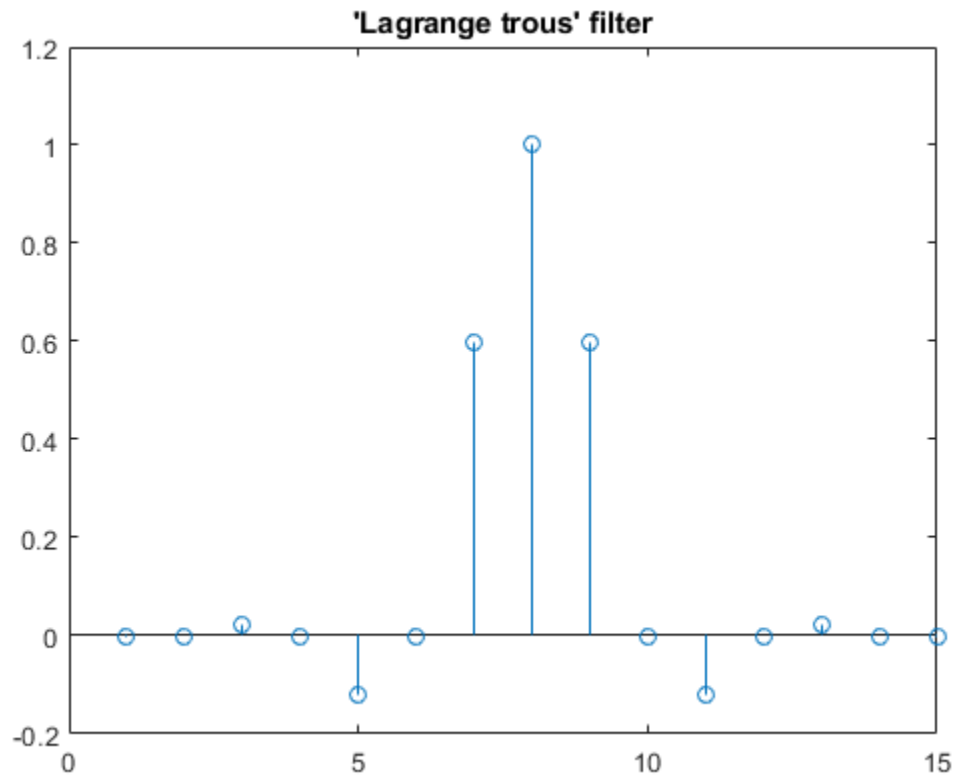
```
0.1629 0.5055 0.4461 -0.0198 -0.1323 0.0218 0.0233 -0.0075
```

```
stem(wdb4)  
title('Order 4 Daubechies Scaling Filter')
```



wdb4 is a solution of the equation: $P = \text{conv}(\text{wrev}(w), w) * 2$, where P is the "Lagrange trous" filter for $N = 4$. Evaluate P and plot it. P is a symmetric filter and wdb4 is a minimum phase solution of the previous equation based on the roots of P.

```
P = conv(wrev(wdb4), wdb4) * 2;  
stem(P)  
title('''Lagrange trous'' filter')
```



Generate `wsym4`, the order 4 symlet scaling filter and plot it. The Symlets are the "least asymmetric" Daubechies' wavelets obtained from another choice between the roots of P .

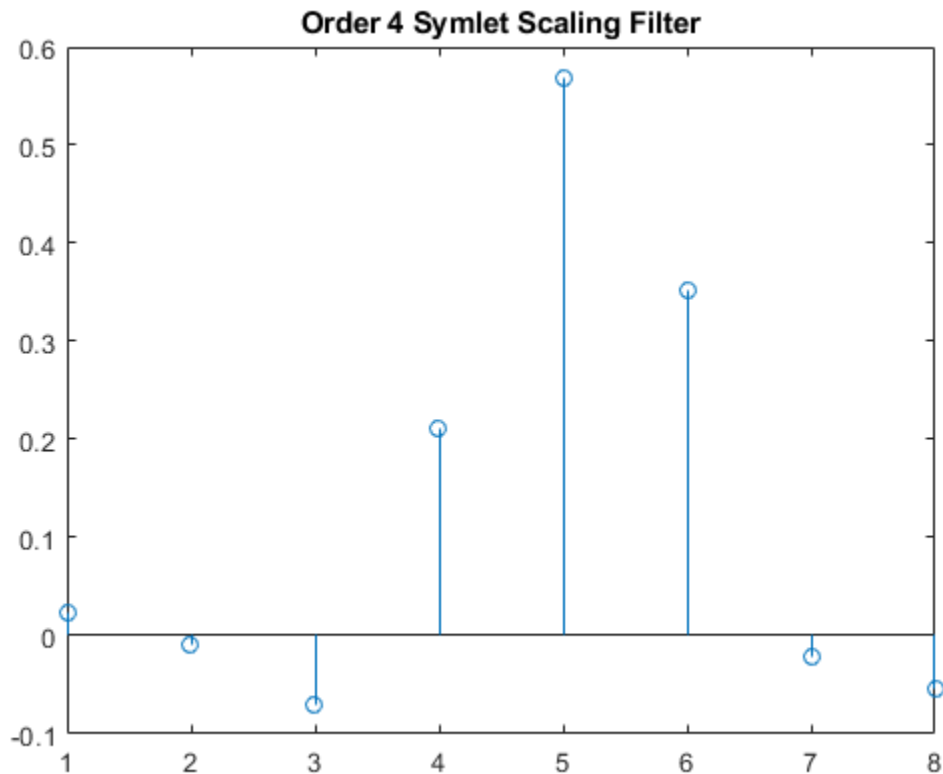
```
wsym4 = symaux(4)
```

```
wsym4 = 1×8
```

```
0.0228 -0.0089 -0.0702 0.2106 0.5683 0.3519 -0.0210 -0.0536
```

```
stem(wsym4)
```

```
title('Order 4 Symlet Scaling Filter')
```



Compute $\text{conv}(\text{wrev}(\text{wsym4}), \text{wsym4}) * 2$ and confirm that wsym4 is another solution of the equation $P = \text{conv}(\text{wrev}(w), w) * 2$.

```
P_sym = conv(wrev(wsym4), wsym4) * 2;
err = norm(P_sym - P)
```

```
err = 1.8677e-15
```

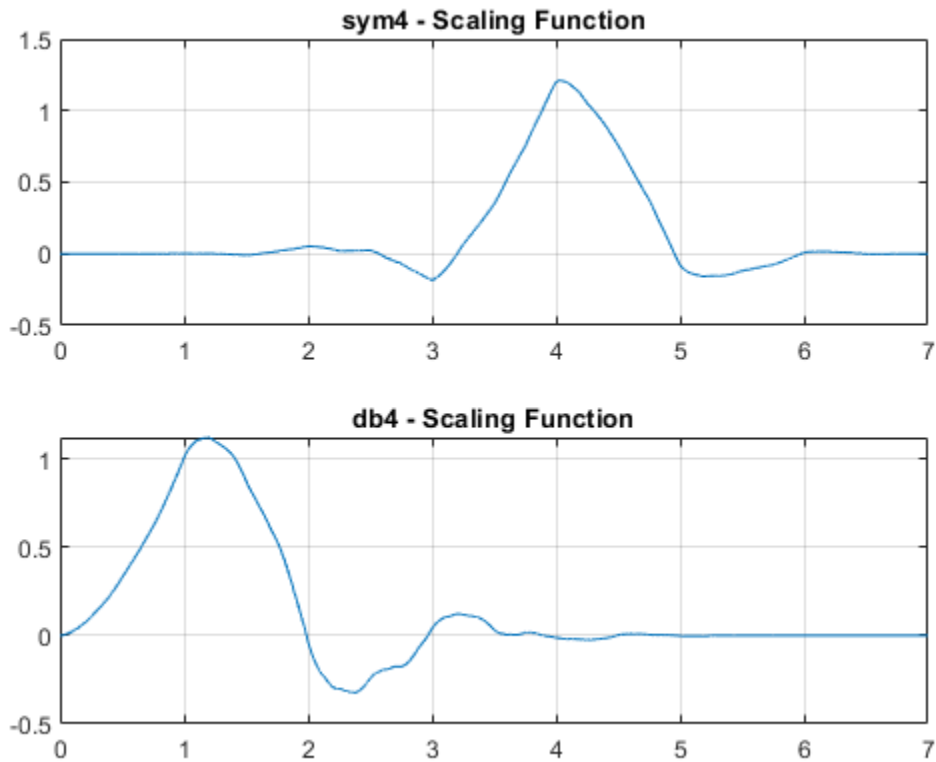
Least Asymmetric Wavelet and Phase

For a given support, the orthogonal wavelet with a phase response that most closely resembles a linear phase filter is called least asymmetric. Symlets are examples of least

asymmetric wavelets. They are modified versions of the classic Daubechies db wavelets. In this example you will show that the order 4 symlet has a nearly linear phase response, while the order 4 Daubechies wavelet does not.

First plot the order 4 symlet and order 4 Daubechies scaling functions. While neither is perfectly symmetric, note how much more symmetric the symlet is.

```
[phi_sym,~,xval_sym]=wavefun('sym4',10);  
[phi_db,~,xval_db]=wavefun('db4',10);  
subplot(2,1,1)  
plot(xval_sym,phi_sym)  
title('sym4 - Scaling Function')  
grid on  
subplot(2,1,2)  
plot(xval_db,phi_db)  
title('db4 - Scaling Function')  
grid on
```



Generate the filters associated with the order 4 symlet and Daubechies wavelets.

```
scal_sym = symaux(4,sqrt(2));  
scal_db = dbaux(4,sqrt(2));
```

Compute the frequency response of the scaling synthesis filters.

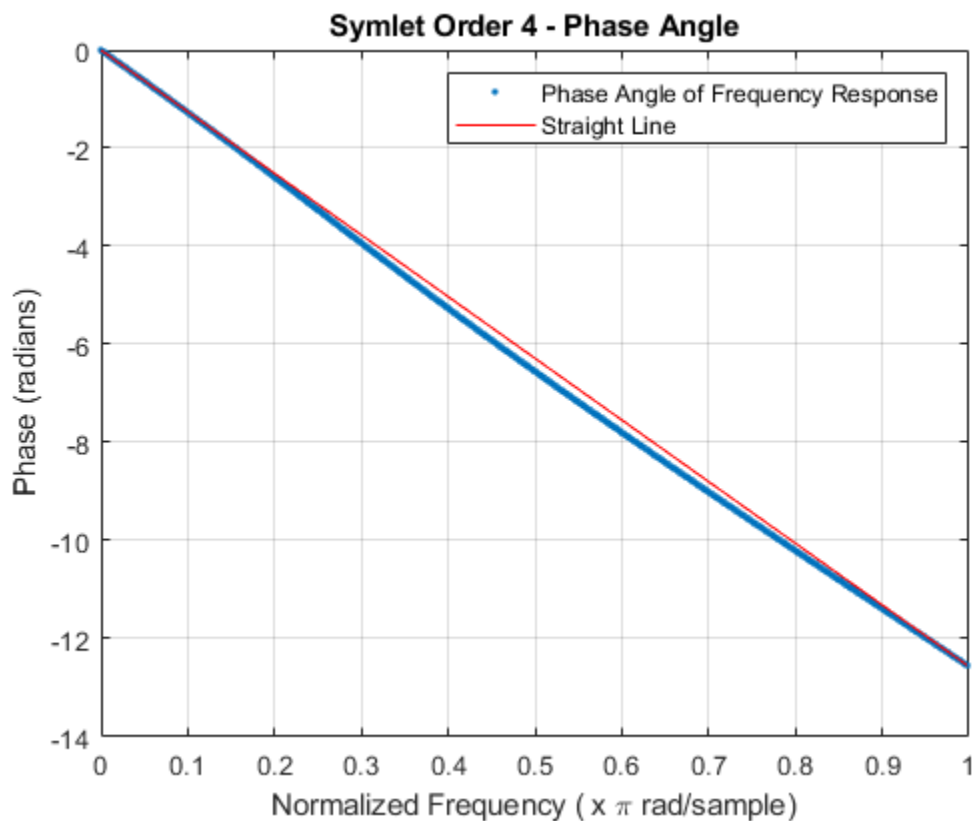
```
[h_sym,w_sym] = freqz(scal_sym);  
[h_db,w_db] = freqz(scal_db);
```

To avoid visual discontinuities, unwrap the phase angles of the frequency responses and plot them. Note how well the phase angle of the symlet filter approximates a straight line.

```
h_sym_u = unwrap(angle(h_sym));  
h_db_u = unwrap(angle(h_db));
```



```
figure
plot(w_sym/pi,h_sym_u, '.')
hold on
plot(w_sym([1 end])/pi,h_sym_u([1 end]), 'r')
grid on
xlabel('Normalized Frequency ( x \pi rad/sample)')
ylabel('Phase (radians)')
legend('Phase Angle of Frequency Response', 'Straight Line')
title('Symlet Order 4 - Phase Angle')
```

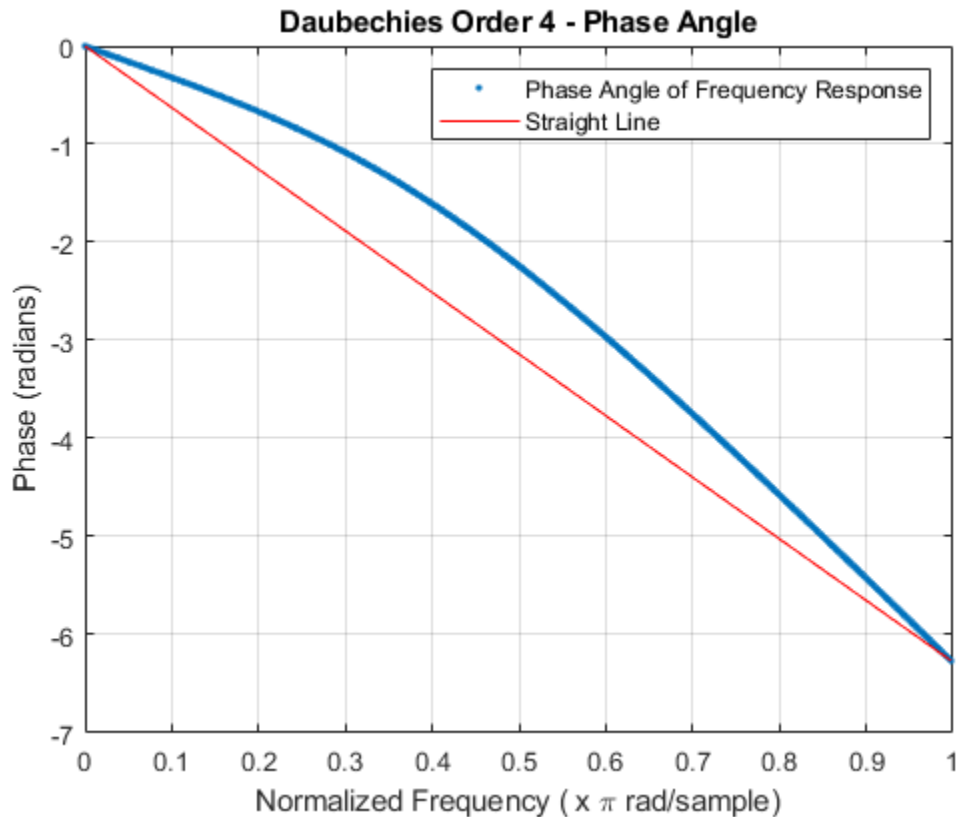


```
figure
plot(w_db/pi,h_db_u, '.')
hold on
plot(w_db([1 end])/pi,h_db_u([1 end]), 'r')
```

```

grid on
xlabel('Normalized Frequency ( x \pi rad/sample)')
ylabel('Phase (radians)')
legend('Phase Angle of Frequency Response','Straight Line')
title('Daubechies Order 4 - Phase Angle')

```



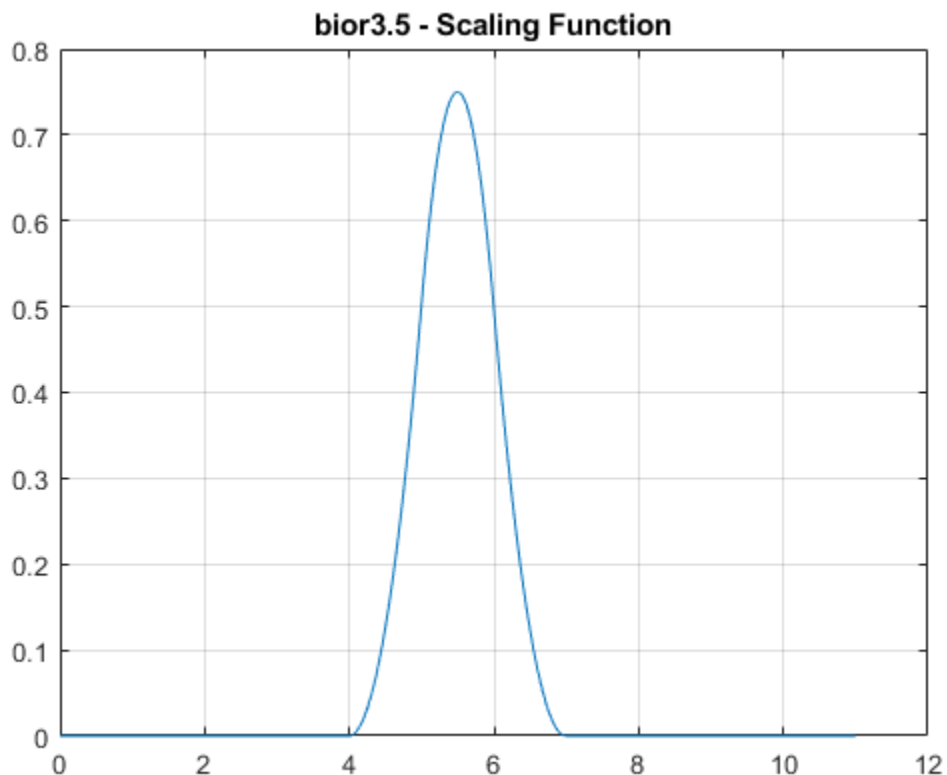
The `sym4` and `db4` wavelets are not symmetric, but the biorthogonal wavelet is. Plot the scaling function associated with the `bior3.5` wavelet. Compute the frequency response of the synthesis scaling filter for the wavelet and verify that it has linear phase.

```

[~,~,phi_bior_r,~,xval_bior]=wavefun('bior3.5',10);
figure
plot(xval_bior,phi_bior_r)

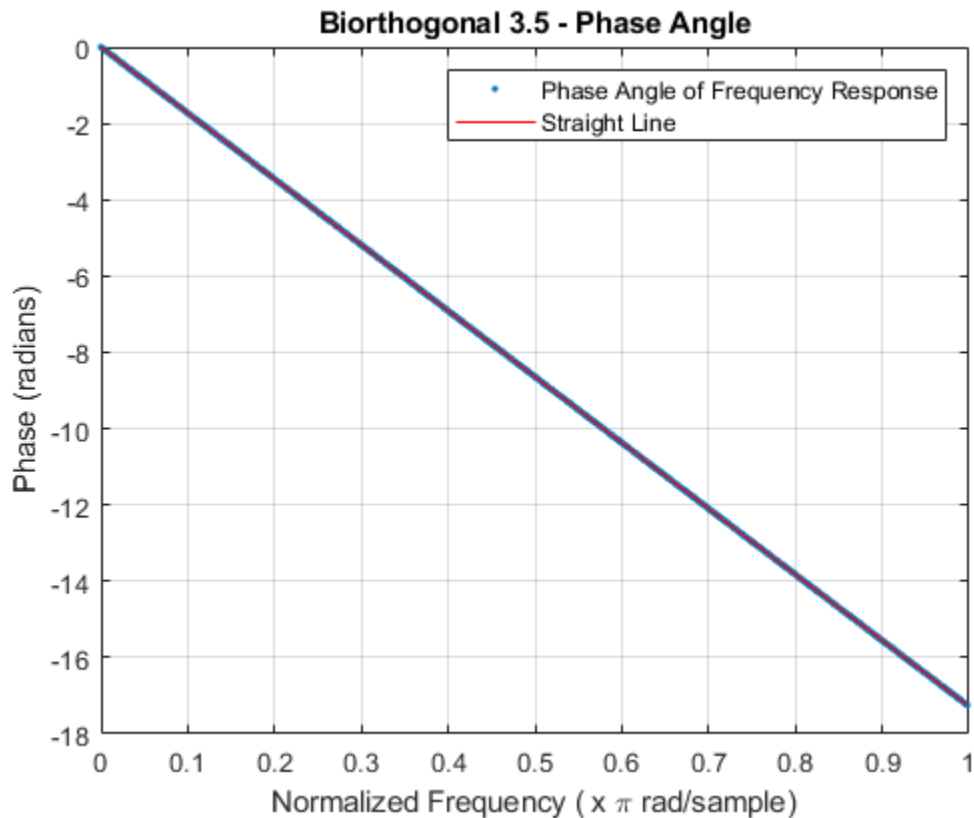
```

```
title('bior3.5 - Scaling Function')
grid on
```



```
[LoD_bior,HiD_bior,LoR_bior,HiR_bior] = wfilters('bior3.5');
[h_bior,w_bior] = freqz(LoR_bior);
h_bior_u = unwrap(angle(h_bior));
figure
plot(w_bior/pi,h_bior_u, '.')
hold on
plot(w_bior([1 end])/pi,h_bior_u([1 end]), 'r')
grid on
xlabel('Normalized Frequency ( x \pi rad/sample)')
ylabel('Phase (radians)')
```

```
legend('Phase Angle of Frequency Response','Straight Line')  
title('Biorthogonal 3.5 - Phase Angle')
```



Extremal Phase

This example demonstrates that for a given support, the cumulative sum of the squared coefficients of a scaling filter increase more rapidly for an extremal phase wavelet than other wavelets.

First, set the order to 15 and generate the scaling filter coefficients for the Daubechies wavelet and Symlet. Both wavelets have support of length 29.

```
n = 15;
[~,~,LoR_db,~] = wfilters('db15');
[~,~,LoR_sym,~] = wfilters('sym15');
```

Next, generate the scaling filter coefficients for the order 5 Coiflet. This wavelet also has support of length 29.

```
[~,~,LoR_coif,~] = wfilters('coif5');
```

Confirm the sum of the coefficients for all three wavelets equals $\sqrt{2}$.

```
sqrt(2)-sum(LoR_db)
```

```
ans = 2.2204e-16
```

```
sqrt(2)-sum(LoR_sym)
```

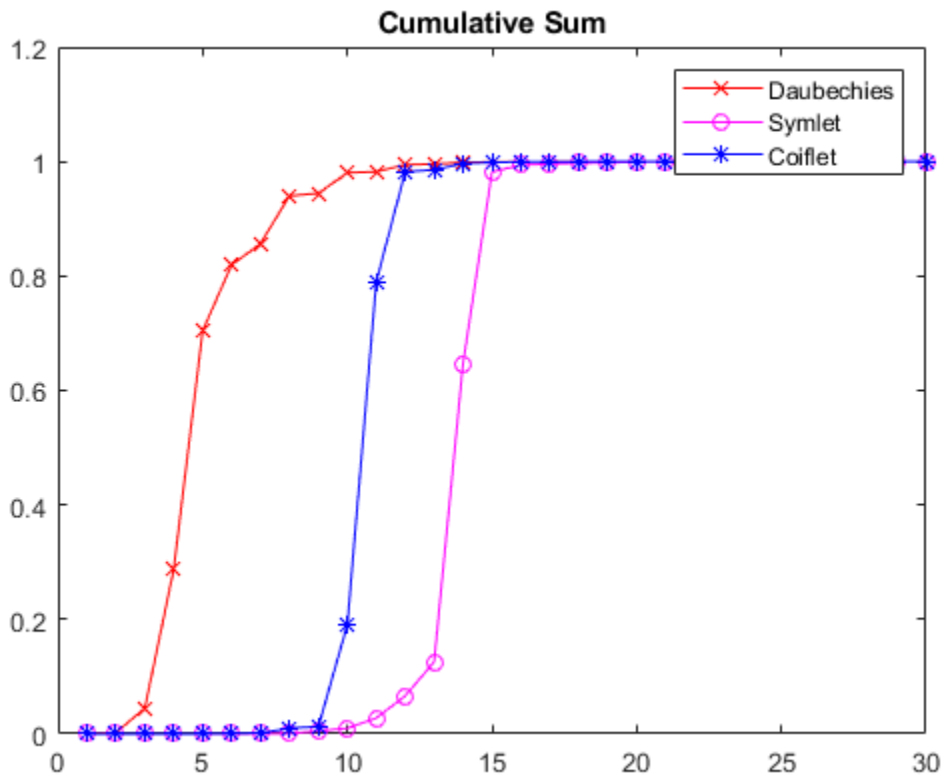
```
ans = 0
```

```
sqrt(2)-sum(LoR_coif)
```

```
ans = 2.2204e-16
```

Plot the cumulative sums of the squared coefficients. Note how rapidly the Daubechies sum increases. This is because its energy is concentrated at small abscissas. Since the Daubechies wavelet has extremal phase, the cumulative sum of its squared coefficients increases more rapidly than the other two wavelets.

```
plot(cumsum(LoR_db.^2), 'rx-')
hold on
plot(cumsum(LoR_sym.^2), 'mo-')
plot(cumsum(LoR_coif.^2), 'b*-')
legend('Daubechies', 'Symlet', 'Coiflet')
title('Cumulative Sum')
```



Input Arguments

n — Order of symlet

positive integer

Order of the symlet, specified as a positive integer.

sumw — Sum of coefficients

1 (default) | positive real number

Sum of the scaling filter coefficients, specified as a positive real number. Set to $\sqrt{2}$ to generate vector of coefficients whose norm is 1.

Output Arguments

w — Filter coefficients

row vector

Vector of scaling filter coefficients of the order n symlet.

The scaling filter coefficients satisfy a number of properties. You can use these properties to check your results. See “Unit Norm Scaling Filter Coefficients” on page 1-1154 for additional information.

Definitions

Least Asymmetric Wavelet

The Haar wavelet, also known as the Daubechies wavelet of order 1, **db1**, is the only compactly supported orthogonal wavelet that is symmetric, or equivalently has linear phase. No other compactly supported orthogonal wavelet can be symmetric. However, it is possible to derive wavelets which are minimally asymmetric, meaning that their phase will be very nearly linear. For a given support, the orthogonal wavelet with a phase response that most closely resembles a linear phase filter is called least asymmetric.

Constructing a compactly supported orthogonal wavelet basis involves choosing roots of a particular polynomial equation. Different choices of roots will result in wavelets whose phases are different. The example “Least Asymmetric Wavelet and Phase” on page 1-1160 compares wavelets with different phases. The example “Symlet and Daubechies Scaling Filters” on page 1-1157 shows that two different scaling filters can satisfy the same polynomial equation. For additional information, see Daubechies [1].

Extremal Phase

As mentioned in “Least Asymmetric Wavelet” on page 1-1169, when constructing a wavelet, you must choose among a set of roots of a particular equation. Choosing roots that lie within the unit circle in the complex plane results in a filter with highly nonlinear phase. Such a wavelet is said to have extremal phase, and has energy concentrated at small abscissas. Let $\{h_k\}$ denote the set of scaling coefficients associated with an extremal phase wavelet, where $k = 1, \dots, N$. Then for any other set of scaling coefficients

$\{g_k\}$ resulting from a different choice of roots, the following inequality will hold for all $J = 1, \dots, N$:

$$\sum_{k=1}^J g_k^2 \leq \sum_{k=1}^J h_k^2$$

The inequality is illustrated in the example “Extremal Phase” on page 1-1166. The $\{h_k\}$ are sometimes called a *minimal delay filter* [2].

References

- [1] Daubechies, I. (1992), *Ten Lectures on Wavelets*, CBMS-NSF conference series in applied mathematics, SIAM Ed.
- [2] Oppenheim, Alan V., and Ronald W. Schafer. *Discrete-Time Signal Processing*. Englewood Cliffs, NJ: Prentice Hall, 1989.

See Also

dbaux | symwavf | wfilters

Introduced before R2006a

symwavf

Symlet wavelet filter

Syntax

```
f = symwavf(wname)
```

Description

`f = symwavf(wname)` returns the scaling filter associated with the Symlet wavelet specified by `wname`. `f` is a real-valued vector.

Examples

Scaling Filter Associated With the Symlet Wavelet

Specify the order 4 Symlet wavelet.

```
wname = 'sym4';
```

Compute the corresponding scaling filter.

```
f = symwavf(wname);  
f'
```

```
ans = 8×1
```

```
 0.0228  
-0.0089  
-0.0702  
 0.2106  
 0.5683  
 0.3519  
-0.0210  
-0.0536
```

Input Arguments

wname — Symlet wavelet

'symN'

Symlet wavelet with N vanishing moments, where N is a positive integer in the closed interval $[1, 45]$.

See Also

[symaux](#) | [waveinfo](#)

Introduced before R2006a

thselect

Threshold selection for denoising

Syntax

```
THR = thselect(X,TPTR)
```

Description

`THR = thselect(X,TPTR)` returns the threshold value adapted to the 1-D signal `X` using the selection rule specified by `TPTR`. Available selection rules are:

- `'rigrsure'` — Adaptive threshold selection using the principle of Stein's Unbiased Risk Estimate (SURE).
- `'sqrtwolog'` — Fixed-form threshold is $\sqrt{2 \cdot \log(\text{length}(X))}$.
- `'heursure'` — Heuristic variant of `'rigrsure'` and `'sqrtwolog'`.
- `'minimaxi'` — Minimax thresholding.

Examples

Threshold Selection Rules

Generate a Gaussian white noise signal. For reproducibility, set the random seed to the default value.

```
rng default  
x = randn(1,1000);
```

Find the threshold for each selection rule.

```
thrRig = thselect(x,'rigrsure');  
disp(['SURE ('rigrsure') threshold: ',num2str(thrRig)]);
```

```
SURE ('rigrsure') threshold: 2.0518
```

```
thrSqt = thselect(x,'sqrtwolog');  
disp(['Universal (''sqrtwolog'') threshold: ',num2str(thrSqt)]);  
  
Universal ('sqrtwolog') threshold: 3.7169  
  
thrHeu = thselect(x,'heursure');  
disp(['Heuristic variant (''heursure'') threshold: ',num2str(thrHeu)]);  
  
Heuristic variant ('heursure') threshold: 3.7169  
  
thrMin = thselect(x,'minimaxi');  
disp(['Minimax (''minimaxi'') threshold: ',num2str(thrMin)]);  
  
Minimax ('minimaxi') threshold: 2.2163
```

Minimax and SURE threshold selection rules are more conservative and would be more convenient when small details of the signal lie near the noise range.

Input Arguments

X — Input data

real-valued vector

Input data, specified as a real-valued vector.

Data Types: double

TPTR — Threshold selection rule

'rigrsure' | 'heursure' | 'sqrtwolog' | 'minimaxi'

Threshold selection rule, specified:

- 'rigrsure' — A threshold selection rule based on SURE (a quadratic loss function) for the soft threshold estimator. Starting with an estimate of risk for a particular threshold value, t , the algorithm minimizes the risks in t to yield a threshold value.
- 'heursure' — A mixture of 'rigrsure' and 'sqrtwolog'. If the signal-to-noise ratio is small, the SURE estimate is noisy. In that case, the fixed-form threshold is used.
- 'sqrtwolog' — A fixed-form (universal) threshold yielding minimax performance multiplied by a small factor proportional to $\log(\text{length}(X))$.

- 'minimaxi' — A fixed threshold chosen to yield minimax performance for mean square error against an ideal procedure. The minimax principle is used in statistics to design estimators. The denoised signal can be assimilated to the estimator of the unknown regression function. Therefore, the minimax estimator realizes the minimum of the maximum mean square error obtained for the worst function in a given set.

Threshold selection rules are based on the underlying model $y = f(t) + e$, where e is an $N(0,1)$ white noise. Use level-dependent noise estimates for unscaled or nonwhite noise. (See `NoiseEstimate` parameter in `wdenoise` for more information.)

Output Arguments

THR — Threshold

positive real number

Threshold value adapted to X , returned as a positive real number.

References

- [1] Donoho, D. L. "Progress in Wavelet Analysis and WVD: A Ten Minute Tour." *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.
- [2] Donoho, D. L., and Johnstone, I. M. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*, Vol. 81, pp. 425-455, 1994.
- [3] Donoho, D. L. "De-noising by Soft-Thresholding." *IEEE Transactions on Information Theory*, Vol. 42, Number 3, pp. 613-627, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

Functions

wdenoise

Apps

Wavelet Signal Denoiser

Introduced before R2006a

tnodes

Determine terminal nodes

Syntax

```
N = tnodes(T)
N = tnodes(T, 'deppos')
[N,K] = tnodes(T)
[N,K] = tnodes(T, 'deppos'), M = N(K)
```

Description

tnodes is a tree-management utility.

`N = tnodes(T)` returns the indices of terminal nodes of the tree T . N is a column vector.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

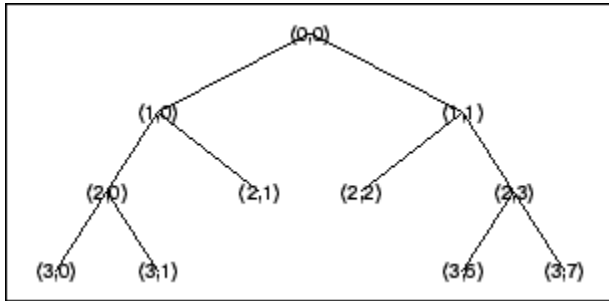
`N = tnodes(T, 'deppos')` returns a matrix N , which contains the depths and positions of terminal nodes.

$N(i, 1)$ is the depth of the i -th terminal node. $N(i, 2)$ is the position of the i -th terminal node.

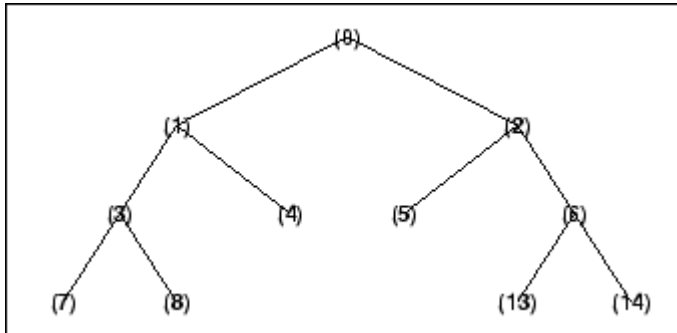
For `[N,K] = tnodes(T)` or `[N,K] = tnodes(T, 'deppos')`, $M = N(K)$ are the indices reordered as in tree T , from left to right.

Examples

```
% Create initial tree.
ord = 2;
t = ntree(ord,3);      % Binary tree of depth 3.
t = nodejoin(t,5);
t = nodejoin(t,4);
plot(t)
```



```
% Change Node Label from Depth_Position to Index
% (see the plot function).
```



```
% List terminal nodes (index).
tnodes(t)
```

```
ans =
     4
     5
     7
     8
    13
    14
```

```
% List terminal nodes (Depth_Position).
tnodes(t,'deppos')
```

```
ans =
     2     1
     2     2
     3     0
     3     1
```


3 6
3 7

See Also

leaves | noleaves | wtreemgr

Introduced before R2006a

treedpth

Tree depth

Syntax

```
D = treedpth(T)
```

Description

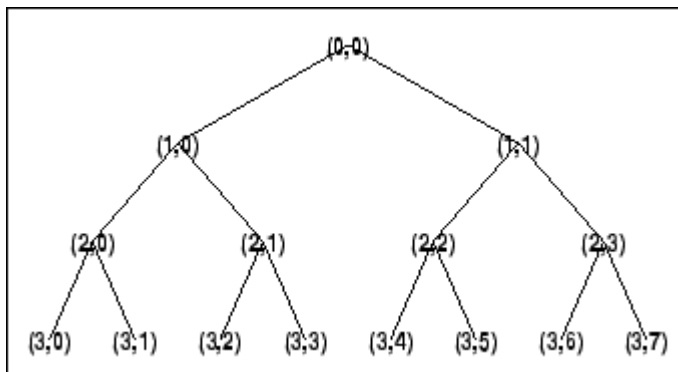
`treedpth` is a tree-management utility.

`D = treedpth(T)` returns the depth `D` of the tree `T`.

Examples

```
% Create binary tree (tree of order 2) of depth 3.  
t = ntree(2,3);
```

```
% Plot tree t.  
plot(t)
```



```
% Tree depth.  
treedpth(t)
```

```
ans =  
  3
```

See Also

wtreemgr

Introduced before R2006a

treeord

Tree order

Syntax

```
ORD = treeord(T)
```

Description

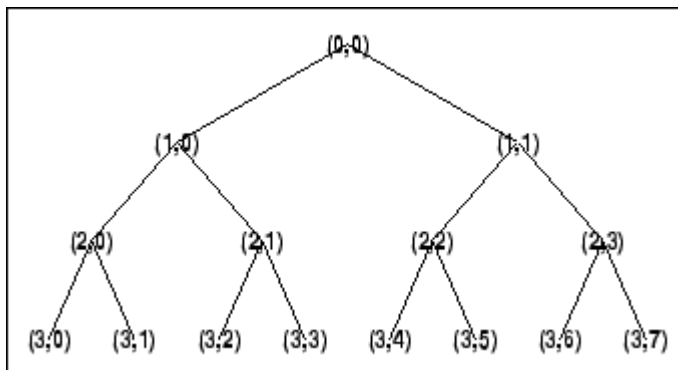
`treeord` is a tree-management utility.

`ORD = treeord(T)` returns the order `ORD` of the tree `T`.

Examples

```
% Create binary tree (tree of order 2) of depth 3.  
t = ntree(2,3);
```

```
% Plot tree t.  
plot(t)
```



```
% Tree order.  
treeord(t)
```

```
ans =  
  2
```

See Also

wtreemgr

Introduced before R2006a

upcoef

Direct reconstruction from 1-D wavelet coefficients

Syntax

```
Y = upcoef(0,X,wname,N)
Y = upcoef(0,X,wname,N,L)
Y = upcoef(0,X,Lo_R,Hi_R,N)
Y = upcoef(0,X,Lo_R,Hi_R,N,L)
Y = upcoef(0,X,wname)
Y = upcoef(0,X,wname,1)
Y = upcoef(0,X,Lo_R,Hi_R)
Y = upcoef(0,X,Lo_R,Hi_R,1)
```

Description

`upcoef` is a one-dimensional wavelet analysis function.

`Y = upcoef(0,X,wname,N)` computes the N -step reconstructed coefficients of vector X .

`wname` is a character vector or string scalar specifying the wavelet. See `wfilters` for more information.

N must be a strictly positive integer.

If `0 = 'a'`, approximation coefficients are reconstructed.

If `0 = 'd'`, detail coefficients are reconstructed.

`Y = upcoef(0,X,wname,N,L)` computes the N -step reconstructed coefficients of vector X and takes the length- L central portion of the result.

Instead of giving the wavelet name, you can give the filters.

For `Y = upcoef(0,X,Lo_R,Hi_R,N)` or `Y = upcoef(0,X,Lo_R,Hi_R,N,L)`, `Lo_R` is the reconstruction low-pass filter and `Hi_R` is the reconstruction high-pass filter.

$Y = \text{upcoef}(0, X, \text{wname})$ is equivalent to $Y = \text{upcoef}(0, X, \text{wname}, 1)$.

$Y = \text{upcoef}(0, X, \text{Lo_R}, \text{Hi_R})$ is equivalent to $Y = \text{upcoef}(0, X, \text{Lo_R}, \text{Hi_R}, 1)$.

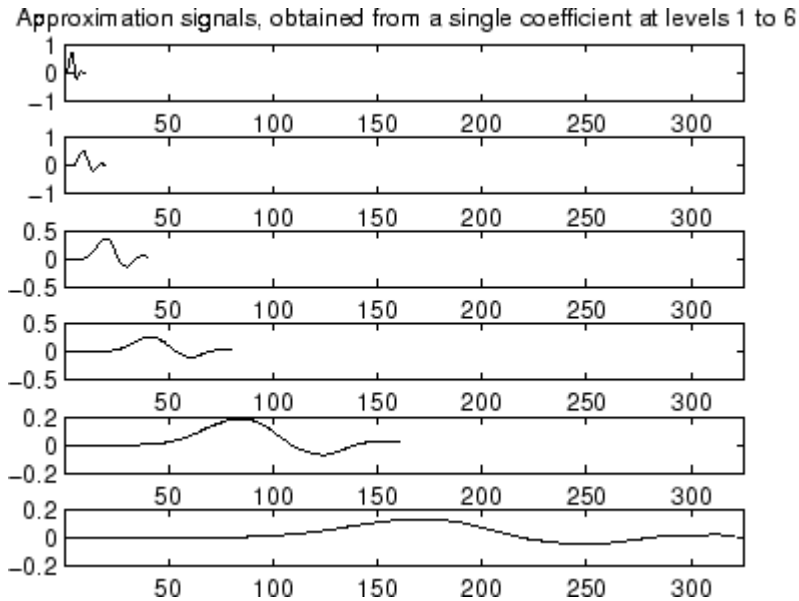
Examples

```
% The current extension mode is zero-padding (see dwtmode).

% Approximation signals, obtained from a single coefficient
% at levels 1 to 6.
cfs = [1]; % Decomposition reduced a single coefficient.
essup = 10; % Essential support of the scaling filter db6.
figure(1)
for i=1:6
    % Reconstruct at the top level an approximation
    % which is equal to zero except at level i where only
    % one coefficient is equal to 1.
    rec = upcoef('a',cfs,'db6',i);

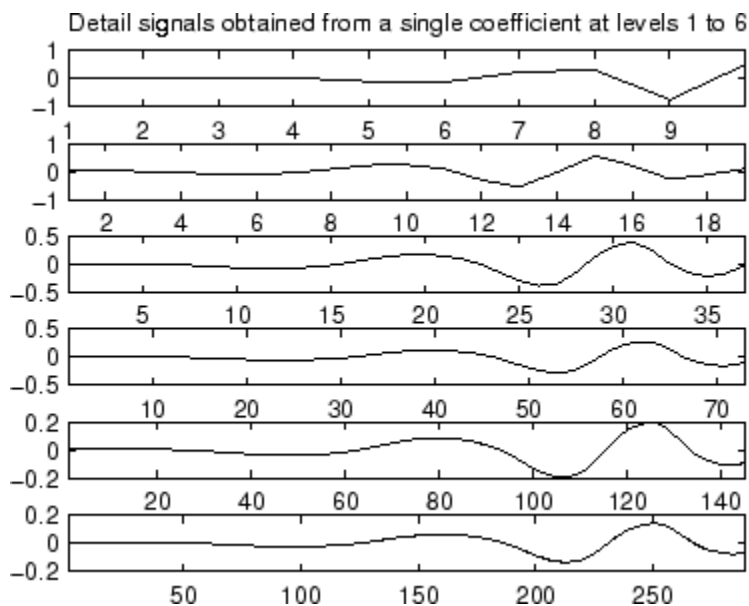
    % essup is the essential support of the
    % reconstructed signal.
    % rec(j) is very small when j is  $\geq$  essup.
    ax = subplot(6,1,i),h = plot(rec(1:essup));
    set(ax,'xlim',[1 325]);
    essup = essup*2;
end
subplot(6,1,1)
title(['Approximation signals, obtained from a single ' ...
      'coefficient at levels 1 to 6'])

% Editing some graphical properties,
% the following figure is generated.
```



```
% The same can be done for details.
% Details signals, obtained from a single coefficient
% at levels 1 to 6.
```

```
cfs = [1];
mi = 12; ma = 30; % Essential support of
                 % the wavelet filter db6.
rec = upcoef('d',cfs,'db6',1);
figure(2)
subplot(611), plot(rec(3:12))
for i=2:6
    % Reconstruct at top level a single detail
    % coefficient at level i.
    rec = upcoef('d',cfs,'db6',i);
    subplot(6,1,i), plot(rec(mi*2^(i-2):ma*2^(i-2)))
end
subplot(611)
title(['Detail signals obtained from a single ' ...
      'coefficient at levels 1 to 6'])
% Editing some graphical properties,
% the following figure is generated.
```

Algorithms

upcoef is equivalent to an N time repeated use of the inverse wavelet transform.

See Also

idwt

Introduced before R2006a

upcoef2

Direct reconstruction from 2-D wavelet coefficients

Syntax

```
Y = upcoef2(0,X,wname,N,S)
Y = upcoef2(0,X,Lo_R,Hi_R,N,S)
Y = upcoef2(0,X,wname,N)
Y = upcoef2(0,X,Lo_R,Hi_R,N)
Y = upcoef2(0,X,wname)
Y = upcoef2(0,X,wname,1)
Y = upcoef2(0,X,Lo_R,Hi_R)
Y = upcoef2(0,X,Lo_R,Hi_R,1)
```

Description

upcoef2 is a two-dimensional wavelet analysis function.

$Y = \text{upcoef2}(0,X,wname,N,S)$ computes the N -step reconstructed coefficients of matrix X and takes the central part of size S . $wname$ is a character vector or string scalar specifying the wavelet. See `wfilters` for more information.

If $0 = 'a'$, approximation coefficients are reconstructed; otherwise if $0 = 'h'$ ('v' or 'd', respectively), horizontal (vertical or diagonal, respectively) detail coefficients are reconstructed. N must be a strictly positive integer.

Instead of giving the wavelet name, you can give the filters.

For $Y = \text{upcoef2}(0,X,Lo_R,Hi_R,N,S)$ is the reconstruction low-pass filter and Hi_R is the reconstruction high-pass filter.

$Y = \text{upcoef2}(0,X,wname,N)$ or $Y = \text{upcoef2}(0,X,Lo_R,Hi_R,N)$ returns the computed result without any truncation.

$Y = \text{upcoef2}(0,X,wname)$ is equivalent to $Y = \text{upcoef2}(0,X,wname,1)$.

$Y = \text{upcoef2}(0,X,Lo_R,Hi_R)$ is equivalent to

```
Y = upcoef2(0,X,Lo_R,Hi_R,1).
```

Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load original image.
```

```
load woman;
```

```
% X contains the loaded image.
```

```
% Perform decomposition at level 2
```

```
% of X using db4.
```

```
[c,s] = wavedec2(X,2,'db4');
```

```
% Reconstruct approximation and details
```

```
% at level 1, from coefficients.
```

```
% This can be done using wrcoef2, or
```

```
% equivalently using:
```

```
%
```

```
% Step 1: Extract coefficients from the
```

```
% decomposition structure [c,s].
```

```
%
```

```
% Step 2: Reconstruct using upcoef2.
```

```
siz = s(size(s,1),:);
```

```
ca1 = appcoef2(c,s,'db4',1);
```

```
a1 = upcoef2('a',ca1,'db4',1,siz);
```

```
chd1 = detcoef2('h',c,s,1);
```

```
hd1 = upcoef2('h',chd1,'db4',1,siz);
```

```
cvd1 = detcoef2('v',c,s,1);
```

```
vd1 = upcoef2('v',cvd1,'db4',1,siz);
```

```
cdd1 = detcoef2('d',c,s,1);
```

```
dd1 = upcoef2('d',cdd1,'db4',1,siz);
```

Algorithms

See upcoef.

See Also

idwt2

Introduced before R2006a

upwlev

Single-level reconstruction of 1-D wavelet decomposition

Syntax

```
[NC,NL,cA] = upwlev(C,L,wname)
[NC,NL,cA] = upwlev(C,L,Lo_R,Hi_R)
```

Description

`upwlev` is a one-dimensional wavelet analysis function.

`[NC,NL,cA] = upwlev(C,L,wname)` performs the single-level reconstruction of the wavelet decomposition structure `[C,L]` giving the new one `[NC,NL]`, and extracts the last approximation coefficients vector `cA`.

`[C,L]` is a decomposition at level `n = length(L) - 2`, so `[NC,NL]` is the same decomposition at level `n-1` and `cA` is the approximation coefficients vector at level `n`.

`wname` is a character vector or string scalar specifying the wavelet, `C` is the original wavelet decomposition vector, and `L` the corresponding bookkeeping vector (for detailed storage information, see `wavedec`).

Instead of giving the wavelet name, you can give the filters.

For `[NC,NL,cA] = upwlev(C,L,Lo_R,Hi_R)`, `Lo_R` is the reconstruction low-pass filter and `Hi_R` is the reconstruction high-pass filter.

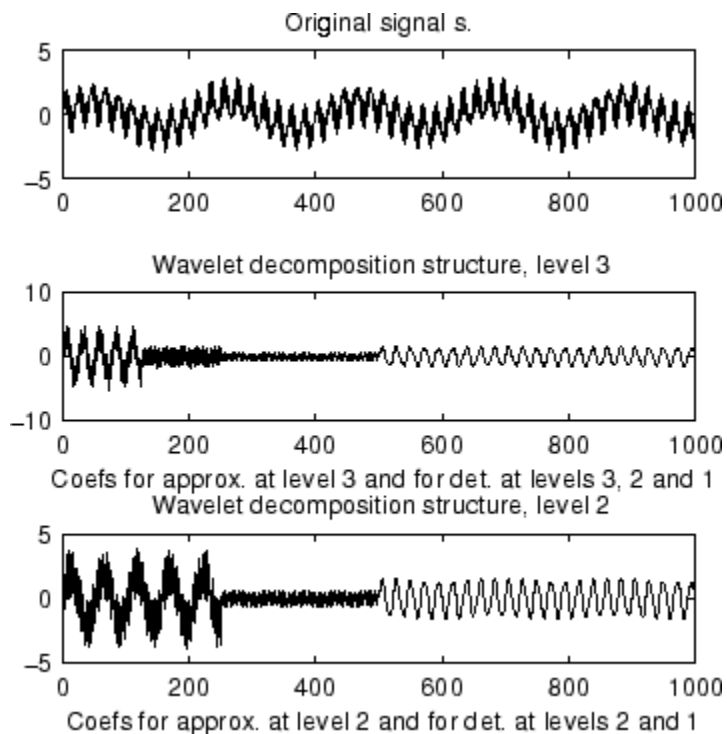
Examples

```
% The current extension mode is zero-padding (see dwtmode).
% Load original one-dimensional signal.
load sumsin; s = sumsin;
```

```
% Perform decomposition at level 3 of s using db1.
[c,l] = wavedec(s,3,'db1');
subplot(311); plot(s);
title('Original signal s. ');
subplot(312); plot(c);
title('Wavelet decomposition structure, level 3')
xlabel(['Coefs for approx. at level 3 ' ...
        'and for det. at levels 3, 2 and 1'])

% One step reconstruction of the wavelet decomposition
% structure at level 3 [c,l], so the new structure [nc,nl]
% is the wavelet decomposition structure at level 2.
[nc,nl] = upwlev(c,l,'db1');
subplot(313); plot(nc);
title('Wavelet decomposition structure, level 2')
xlabel(['Coefs for approx. at level 2 ' ...
        'and for det. at levels 2 and 1'])

% Editing some graphical properties,
% the following figure is generated.
```



See Also

`idwt`

Topics

`upcoef`

`wavedec`

Introduced before R2006a

upwlev2

Single-level reconstruction of 2-D wavelet decomposition

Syntax

```
[NC,NS,cA] = upwlev2(C,S,wname)
[NC,NS,cA] = upwlev2(C,S,Lo_R,Hi_R)
```

Description

`upwlev2` is a two-dimensional wavelet analysis function.

`[NC,NS,cA] = upwlev2(C,S,wname)` performs the single-level reconstruction of wavelet decomposition structure `[C,S]` giving the new one `[NC,NS]`, and extracts the last approximation coefficients matrix `cA`.

`[C,S]` is a decomposition at level `n = size(S,1)-2`, so `[NC,NS]` is the same decomposition at level `n-1` and `cA` is the approximation matrix at level `n`.

`wname` is a character vector or string scalar specifying the wavelet, `C` is the original wavelet decomposition vector, and `S` the corresponding bookkeeping matrix (for detailed storage information, see `wavedec2`).

Instead of giving the wavelet name, you can give the filters.

For `[NC,NS,cA] = upwlev2(C,S,Lo_R,Hi_R)`, `Lo_R` is the reconstruction low-pass filter and `Hi_R` is the reconstruction high-pass filter.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
% Load original image.
load woman;
% X contains the loaded image.
```



```
% Perform decomposition at level 2
% of X using db1.
[c,s] = wavedec2(X,2,'db1');
sc = size(c)

sc =
     1 65536

val_s = s

val_s =
     64  64
     64  64
    128 128
    256 256

% One step reconstruction of wavelet
% decomposition structure [c,s].
[nc,ns] = upwlev2(c,s,'db1');
snc = size(nc)

snc =
     1 65536

val_ns = ns

val_ns =
    128  128
    128  128
    256  256
```

See Also

[idwt2](#) | [upcoef2](#) | [wavedec2](#)

Introduced before R2006a

wave2lp

Laurent polynomials associated with wavelet

Syntax

```
[Hs,Gs,Ha,Ga] = wave2lp(W)
```

Description

`[Hs,Gs,Ha,Ga] = wave2lp(W)` returns the four Laurent polynomials associated with the wavelet W (see `liftwave`).

The pairs (Hs, Gs) and (Ha, Ga) are the synthesis and the analysis pair respectively.

The H-polynomials (G-polynomials) are low-pass (high-pass) polynomials.

For an orthogonal wavelet, $Hs = Ha$ and $Gs = Ga$.

Examples

```
% Get Laurent polynomials associated to the "lazy" wavelet.
```

```
[Hs,Gs,Ha,Ga] = wave2lp('lazy')
```

```
Hs(z) = 1
```

```
Gs(z) = z^(-1)
```

```
Ha(z) = 1
```

```
Ga(z) = z^(-1)
```

```
% Get Laurent polynomials associated to the db1 wavelet.
```

```
[Hs,Gs,Ha,Ga] = wave2lp('db1')
```

```
Hs(z) = + 0.7071 + 0.7071*z^(-1)
```

```

Gs(z) = - 0.7071 + 0.7071*z^(-1)
Ha(z) = + 0.7071 + 0.7071*z^(-1)
Ga(z) = - 0.7071 + 0.7071*z^(-1)

% Get Laurent polynomials associated to the bior1.3 wavelet.
[Hs,Gs,Ha,Ga] = wave2lp('bior1.3')

Hs(z) = + 0.7071 + 0.7071*z^(-1)

Gs(z) = ...
+ 0.08839*z^(+2) + 0.08839*z^(+1) - 0.7071 + 0.7071*z^(-1) -
0.08839*z^(-2) ...
- 0.08839*z^(-3)

Ha(z) = ...
- 0.08839*z^(+2) + 0.08839*z^(+1) + 0.7071 + 0.7071*z^(-1) +
0.08839*z^(-2) ...
- 0.08839*z^(-3)

Ga(z) = - 0.7071 + 0.7071*z^(-1)

```

See Also

laurpoly

Introduced before R2006a

wavedec

1-D wavelet decomposition

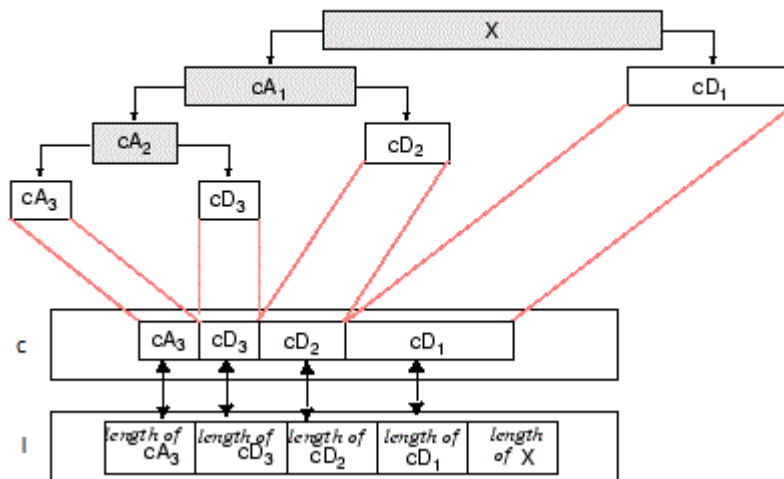
wavedec performs a multilevel one-dimensional wavelet analysis using either a specific wavelet or a specific pair of wavelet decomposition filters.

Syntax

`[c, l] = wavedec(x, n, wname)`
`[c, l] = wavedec(x, n, LoD, HiD)`

Description

`[c, l] = wavedec(x, n, wname)` returns the wavelet decomposition of the signal `x` at level `n` using the wavelet `wname`. The output decomposition structure consists of the wavelet decomposition vector `c` and the bookkeeping vector `l`, which contains the number of coefficients by level. The structure is organized as in this level-3 decomposition diagram.



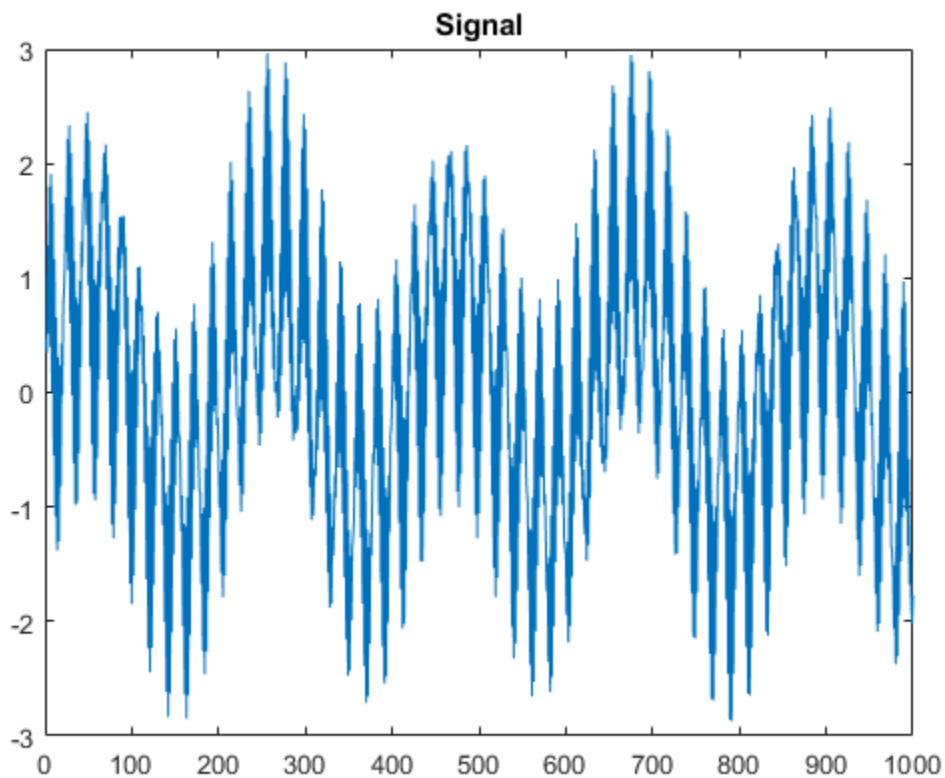
`[c,l] = wavedec(x,n,LoD,HiD)` returns the wavelet decomposition using the specified lowpass, and highpass wavelet decomposition filters `LoD` and `HiD`.

Examples

Multilevel One-Dimensional Wavelet Analysis

Load and plot a one-dimensional signal.

```
load sumsin
plot(sumsin)
title('Signal')
```

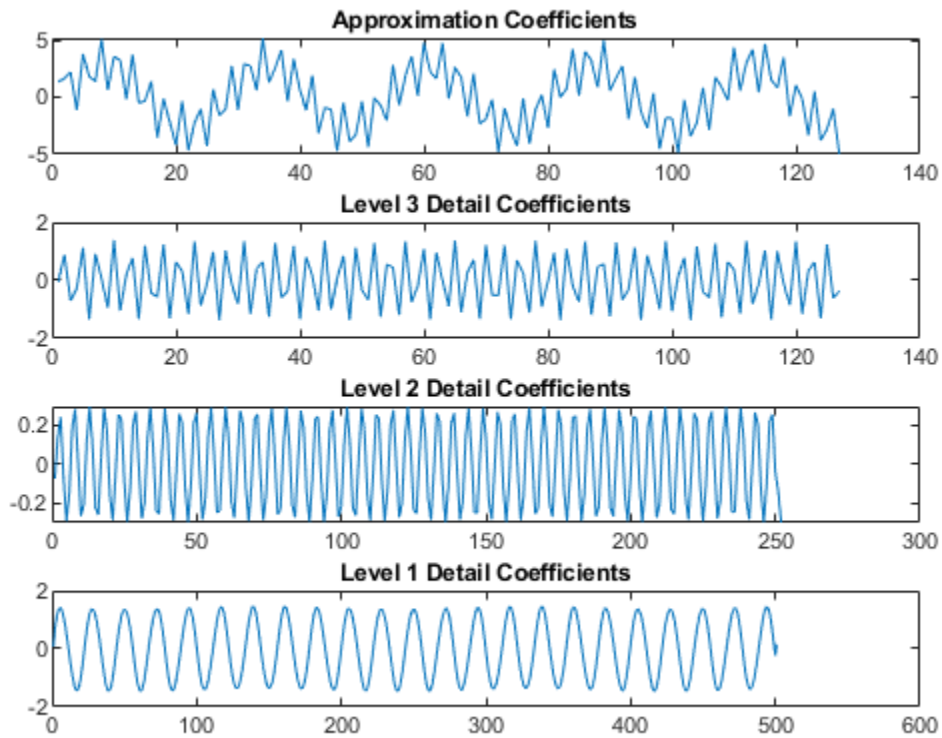


Perform a 3-level wavelet decomposition of the signal using the order 2 Daubechies wavelet. Extract the coarse scale approximation coefficients and the detail coefficients from the decomposition.

```
[c,l] = wavedec(sumsin,3,'db2');  
approx = appcoef(c,l,'db2');  
[cd1,cd2,cd3] = detcoef(c,l,[1 2 3]);
```

Plot the coefficients.

```
subplot(4,1,1)  
plot(approx)  
title('Approximation Coefficients')  
subplot(4,1,2)  
plot(cd3)  
title('Level 3 Detail Coefficients')  
subplot(4,1,3)  
plot(cd2)  
title('Level 2 Detail Coefficients')  
subplot(4,1,4)  
plot(cd1)  
title('Level 1 Detail Coefficients')
```



Input Arguments

x — Input signal

real-valued vector

Input signal, specified as a real-valued vector.

Data Types: double

n — Level of decomposition

positive integer

Level of decomposition, specified as a positive integer. `wavedec` does not enforce a maximum level restriction. Use `wmaxlev` to ensure that the wavelet coefficients are free from boundary effects. If boundary effects are not a concern in your application, a good rule is to set `n` less than or equal to `fix(log2(length(x)))`.

Data Types: `double`

wname — Analyzing wavelet

character vector | string scalar

Analyzing wavelet, specified as a character vector or string scalar.

Note `wavedec` supports only Type 1 (orthogonal) or Type 2 (biorthogonal) wavelets. See `wfilters` for a list of orthogonal and biorthogonal wavelets.

LoD — Wavelet decomposition lowpass filter

even-length real-valued vector

Wavelet decomposition lowpass filter, specified as an even-length real-valued vector. `LoD` must be of the same length as `HiD`. See `wfilters` for details.

HiD — Wavelet decomposition highpass filter

even-length real-valued vector

Wavelet decomposition highpass filter, specified as an even-length real-valued vector. `HiD` must be of the same length as `LoD`. See `wfilters` for details.

Output Arguments

c — Wavelet decomposition vector

real-valued vector

Wavelet decomposition vector, returned as a real-valued vector. The bookkeeping vector `l` contains the number of coefficients by level.

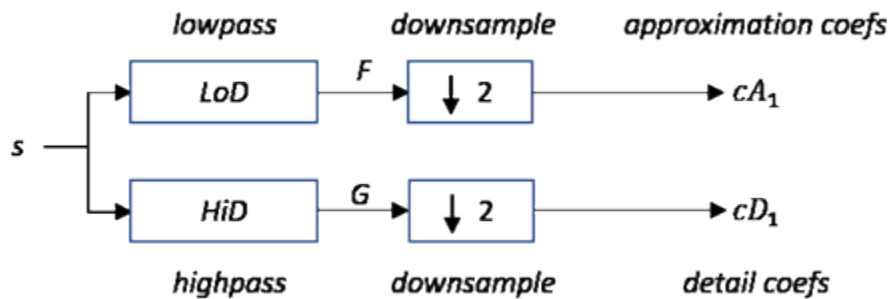
l — Bookkeeping vector

vector of positive integers

Bookkeeping vector, returned as a vector of positive integers. The bookkeeping vector is used to parse the coefficients in the wavelet decomposition vector `c` by level.

Algorithms

Given a signal s of length N , the DWT consists of at most $\log_2 N$ steps. Starting from s , the first step produces two sets of coefficients: approximation coefficients cA_1 and detail coefficients cD_1 . Convolution of s with the lowpass filter LoD and the highpass filter HiD , followed by dyadic decimation (downsampling), results in the approximation and detail coefficients respectively.



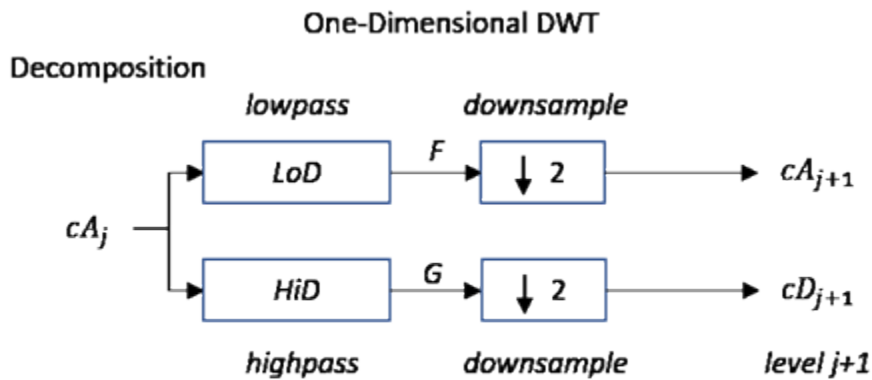
where

- \boxed{X} — Convolve with filter X
- $\boxed{\downarrow 2}$ — Downsample (keep the even-indexed elements)

The length of each filter is equal to $2n$. If $N = \text{length}(s)$, the signals F and G are of length $N + 2n - 1$ and the coefficients cA_1 and cD_1 are of length

$$\text{floor}\left(\frac{N-1}{2}\right) + n.$$

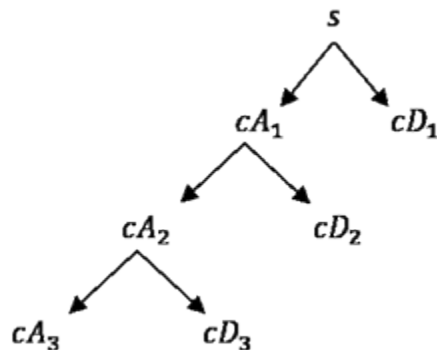
The next step splits the approximation coefficients cA_1 in two parts using the same scheme, replacing s by cA_1 , and producing cA_2 and cD_2 , and so on.



Initialization: $cA_0 = s$

The wavelet decomposition of the signal s analyzed at level j has the following structure: $[cA_j, cD_j, \dots, cD_1]$.

This structure contains, for $j = 3$, the terminal nodes of the following tree:



References

- [1] Daubechies, I. *Ten Lectures on Wavelets*, CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: SIAM Ed, 1992.

[2] Mallat, S. G. "A Theory for Multiresolution Signal Decomposition: The Wavelet Representation," *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 11, Issue 7, July 1989, pp. 674-693.

[3] Meyer, Y. *Wavelets and Operators*. Translated by D. H. Salinger. Cambridge, UK: Cambridge University Press, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.
- The input `wname` must be constant.

See Also

`appcoef` | `detcoef` | `dwt` | `dwtfilterbank` | `waveinfo` | `waverec` | `wfilters` | `wmaxlev`

Introduced before R2006a

wavedec2

2-D wavelet decomposition

Syntax

```
[C,S] = wavedec2(X,N,wname)  
[C,S] = wavedec2(X,N,Lod,Hid)
```

Description

`[C,S] = wavedec2(X,N,wname)` returns the wavelet decomposition of the matrix `X` at level `N` using the wavelet `wname`. The output decomposition structure consists of the wavelet decomposition vector `C` and the bookkeeping matrix `S`, which contains the number of coefficients by level and orientation.

`[C,S] = wavedec2(X,N,Lod,Hid)` returns the wavelet decomposition using the specified lowpass and highpass decomposition filters `LoD` and `HiD`, respectively. See `wfilters` for details.

Examples

Extract and Display Image Decomposition Level

Extract and display images of wavelet decomposition level details. The resulting images are similar to the visualizations in the Example *Analysis At level 2, with haar ---> woman* indexed image accessible in the **Wavelet 2-D** interactive tool. Use `waveletAnalyzer` to launch this tool.

Load an image. Perform a level 2 wavelet decomposition of the image using the `haar` wavelet.

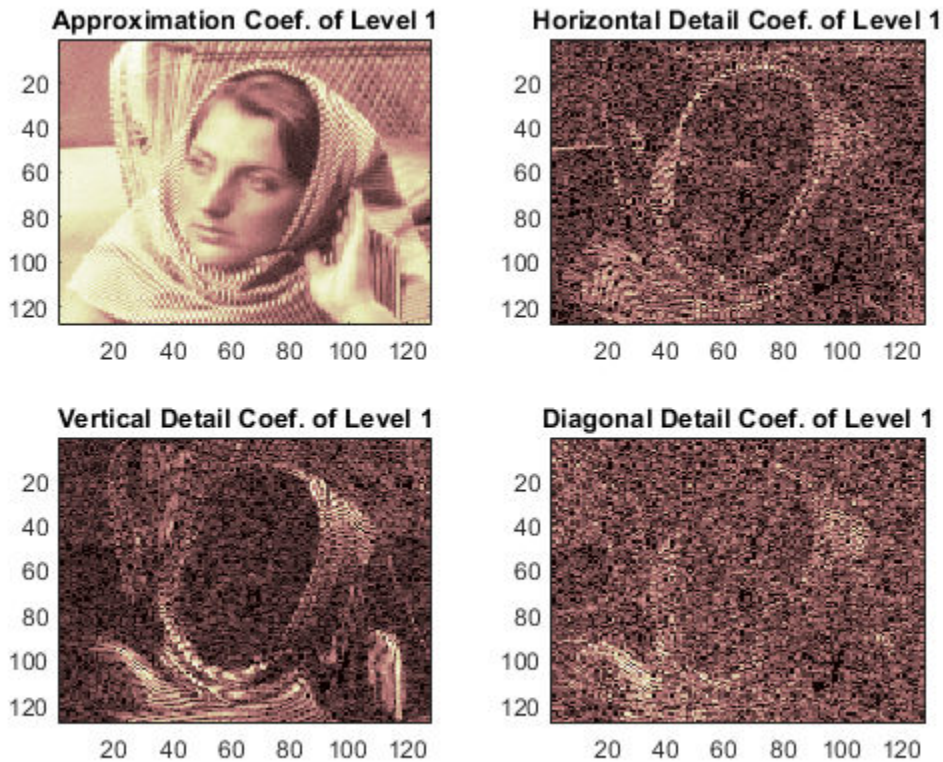
```
load woman  
[c,s]=wavedec2(X,2,'haar');
```

Extract the level 1 approximation and detail coefficients.

```
[H1,V1,D1] = detcoef2('all',c,s,1);  
A1 = appcoef2(c,s,'haar',1);
```

Use `wcodemat` to rescale the coefficients based on their absolute values. Display the rescaled coefficients.

```
V1img = wcodemat(V1,255,'mat',1);  
H1img = wcodemat(H1,255,'mat',1);  
D1img = wcodemat(D1,255,'mat',1);  
A1img = wcodemat(A1,255,'mat',1);  
  
subplot(2,2,1)  
imagesc(A1img)  
colormap pink(255)  
title('Approximation Coef. of Level 1')  
  
subplot(2,2,2)  
imagesc(H1img)  
title('Horizontal Detail Coef. of Level 1')  
  
subplot(2,2,3)  
imagesc(V1img)  
title('Vertical Detail Coef. of Level 1')  
  
subplot(2,2,4)  
imagesc(D1img)  
title('Diagonal Detail Coef. of Level 1')
```



Extract the level 2 approximation and detail coefficients.

```
[H2,V2,D2] = detcoef2('all',c,s,2);
A2 = appcoef2(c,s,'haar',2);
```

Use `wcodemat` to rescale the coefficients based on their absolute values. Display the rescaled coefficients.

```
V2img = wcodemat(V2,255,'mat',1);
H2img = wcodemat(H2,255,'mat',1);
D2img = wcodemat(D2,255,'mat',1);
A2img = wcodemat(A2,255,'mat',1);
```

```
figure
```

```
subplot(2,2,1)
imagesc(A2img)
colormap pink(255)
title('Approximation Coef. of Level 2')

subplot(2,2,2)
imagesc(H2img)
title('Horizontal Detail Coef. of Level 2')

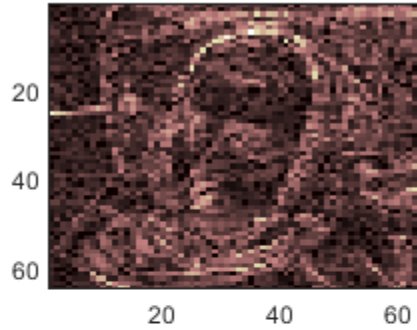
subplot(2,2,3)
imagesc(V2img)
title('Vertical Detail Coef. of Level 2')

subplot(2,2,4)
imagesc(D2img)
title('Diagonal Detail Coef. of Level 2')
```

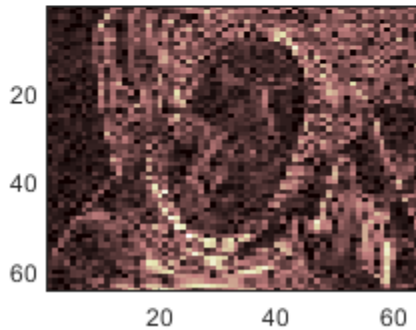
Approximation Coef. of Level 2



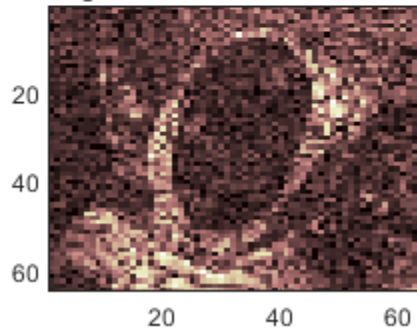
Horizontal Detail Coef. of Level 2



Vertical Detail Coef. of Level 2



Diagonal Detail Coef. of Level 2



2-D Wavelet Decomposition Structure

This example shows the structure of `wavedec2` output matrices.

Load and display an image.

```
load woman
imagesc(X)
colormap(map)
```




Save the current discrete wavelet transform extension mode.

```
origMode = dwtmode('status','nodisplay');
```

Change to periodic boundary handling. The `dwtmode` function displays a message indicating that the DWT extension mode is changing.

```
dwtmode('per');
```

```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!  WARNING: Change DWT Extension Mode  !
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
*****  
** DWT Extension Mode: Periodization **  
*****
```

Perform a level 3 decomposition of the image using the db1 (Haar) wavelet.

```
[c,s] = wavedec2(X,3,'db1');
```

Return the number of elements in the image *X* and coefficient vector *c*. Confirm the number of elements in each are equal.

```
numel(X)
```

```
ans = 65536
```

```
numel(c)
```

```
ans = 65536
```

Display the bookkeeping matrix *s*. The first row displays the dimensions of the coarse scale approximation of the image. The last row displays the dimensions of the original image. The intermediate rows display the dimensions of the detail coefficients at the three levels of the decomposition, proceeding from coarse to fine scale.

```
s
```

```
s = 5x2
```

```
    32    32  
    32    32  
    64    64  
   128   128  
   256   256
```

Reset discrete wavelet transform extension mode to its original mode. The `dwtmode` function displays a message indicating that the DWT extension mode is changing.

```
dwtmode(origMode);
```

```
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!  
! WARNING: Change DWT Extension Mode !  
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
*****
**   DWT Extension Mode: Symmetrization (half-point) **
*****
```

Input Arguments

X — Input data

real-valued matrix | 3-D array of RGB triplets

Input data, specified as a real-valued M -by- N matrix representing an indexed image, or an M -by- N -by-3 array representing a truecolor image. For more information on truecolor images, see “RGB (Truecolor) Images” (MATLAB).

N — Decomposition level

positive integer

Decomposition level, specified as a positive integer. `wavedec2` does not enforce a maximum level restriction. Use `wmaxlev` to determine the maximum decomposition level possible of the matrix X using the wavelet `wname`. The maximum level is the last level for which at least one coefficient is correct.

wname — Analyzing wavelet

character vector | string scalar

Analyzing wavelet, specified as a character vector or string scalar.

Note `wavedec2` supports only Type 1 (orthogonal) or Type 2 (biorthogonal) wavelets. See `wfilters` for a list of orthogonal and biorthogonal wavelets.

Lod, Hid — Wavelet decomposition filters

even-length real-valued vectors

Wavelet decomposition filters associated with an orthogonal or biorthogonal wavelet, specified as even-length real-valued vectors. `Lod` is the lowpass decomposition filter, and `Hid` is the highpass decomposition filter. See `wfilters` for details.

Output Arguments

C — Wavelet decomposition vector

real-valued vector

Wavelet decomposition vector. The vector *C* contains the approximation and detail coefficients organized by level. The bookkeeping matrix *S* is used to parse *C*.

The vector *C* is organized as $A(N), H(N), V(N), D(N), H(N-1), V(N-1), D(N-1), \dots, H(1), V(1), D(1)$, where *A*, *H*, *V*, and *D* are each a row vector. Each vector is the column-wise storage of a matrix.

- *A* contains the approximation coefficients.
- *H* contains the horizontal detail coefficients.
- *V* contains the vertical detail coefficients.
- *D* contains the diagonal detail coefficients.

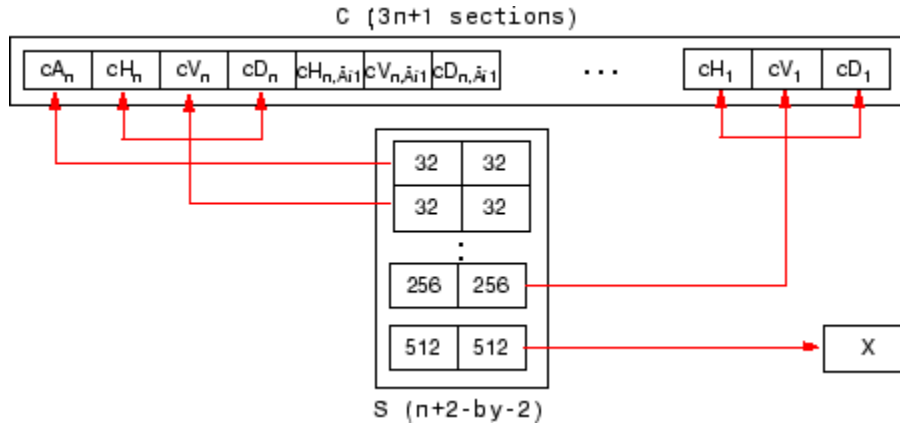
S — Bookkeeping matrix

integer-valued matrix

Bookkeeping matrix. The matrix *S* contains the dimensions of the wavelet coefficients by level and is used to parse the wavelet decomposition vector *C*.

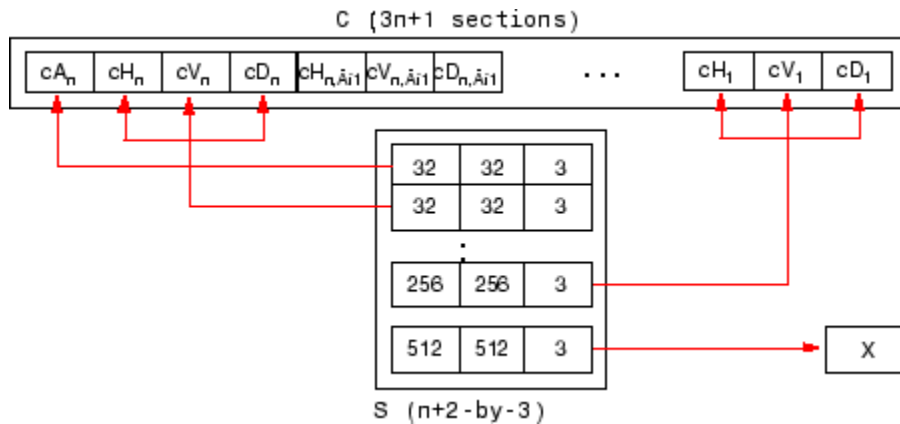
- $S(1, :) = \text{size of approximation coefficients}(N)$.
- $S(i, :) = \text{size of detail coefficients}(N - i + 2)$ for $i = 2, \dots, N+1$ and $S(N+2, :) = \text{size}(X)$.

The following diagram shows the relationship between *C* and *S* in the wavelet decomposition of a 512-by-512 matrix.



When X represents an indexed image, the output arrays cA , cH , cV , and cD are m -by- n matrices. When X represents a truecolor image, it is an m -by- n -by-3 array, where each m -by- n matrix represents a red, green, or blue color plane concatenated along the third dimension. The size of vector C and the size of matrix S depend on the type of analyzed image.

For a truecolor image, the decomposition vector C and the corresponding bookkeeping matrix S can be represented as shown.

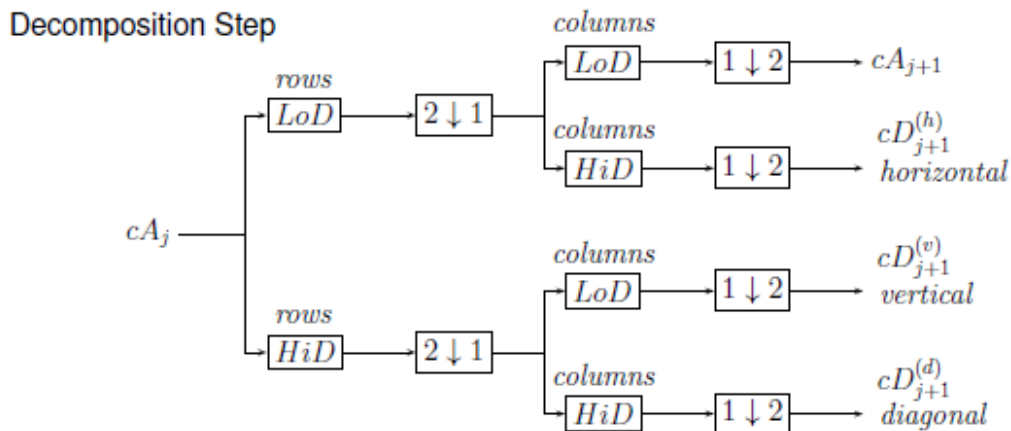


Algorithms

For images, an algorithm similar to the one-dimensional case is possible for two-dimensional wavelets and scaling functions obtained from one-dimensional vectors by tensor product. This kind of two-dimensional DWT leads to a decomposition of approximation coefficients at level j in four components: the approximation at level $j+1$ and the details in three orientations (horizontal, vertical, and diagonal).

The chart describes the basic decomposition step for images:

Two-Dimensional DWT



where

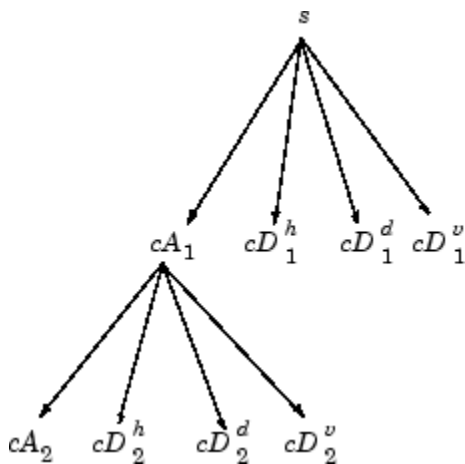
- $\boxed{2 \downarrow 1}$ — Downsample columns: keep the even-indexed columns.
- $\boxed{1 \downarrow 2}$ — Downsample rows: keep the even-indexed rows.
- $\boxed{\overset{\text{rows}}{X}}$ — Convolve with filter X the rows of the entry.

- $\overset{\text{columns}}{\boxed{X}}$ — Convolve with filter X the rows of the entry.

and

Initialization: $cA_0 = s$.

So, for $J = 2$, the two-dimensional wavelet tree has the form



References

[1] Daubechies, I. *Ten Lectures on Wavelets*, CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: SIAM Ed, 1992.

[2] Mallat, S. G. "A Theory for Multiresolution Signal Decomposition: The Wavelet Representation," *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 11, Issue 7, July 1989, pp. 674-693.

[3] Meyer, Y. *Wavelets and Operators*. Translated by D. H. Salinger. Cambridge, UK: Cambridge University Press, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.
- The input '*wname*' must be constant.

See Also

[dwt2](#) | [waveinfo](#) | [waverec2](#) | [wfilters](#) | [wmaxlev](#)

Introduced before R2006a

wavedec3

3-D wavelet decomposition

Syntax

```
WDEC = wavedec3(X,N,wname)
WDEC = wavedec3(X,N,wname,'mode','ExtM')
WDEC = wavedec3(X,N,{LoD,HiD,LoR,HiR})
```

Description

wavedec3 is a three-dimensional wavelet analysis function.

`WDEC = wavedec3(X,N,wname)` returns the wavelet decomposition of the 3-D array `X` at level `N`, using the wavelet specified by the character vector or string scalar `wname` or the particular wavelet filters you specify. It uses the default extension mode `'sym'`. See `dwtmode`. `N` must be a positive integer.

`WDEC = wavedec3(X,N,wname,'mode','ExtM')` uses the specified DWT extension mode.

`WDEC = wavedec3(X,N,{LoD,HiD,LoR,HiR})` uses the decomposition and reconstruction filters you specify in a cell array.

`WDEC` is the output decomposition structure, with the following fields:

<code>sizeINI</code>	Size of the three-dimensional array <code>X</code>
<code>level</code>	Level of the decomposition
<code>mode</code>	Name of the wavelet transform extension mode
<code>filters</code>	Structure with 4 fields, <code>LoD</code> , <code>HiD</code> , <code>LoR</code> , <code>HiR</code> , which contain the filters used for the DWT.

<p>dec</p>	<p>$N \times 1$ cell array containing the coefficients of the decomposition. N is equal to $7 * WDEC.level + 1$.</p> <p>dec{1} contains the lowpass component (approximation) at the level of the decomposition. The approximation is equivalent to the filtering operations 'LLL'.</p> <p>dec{k+2}, ..., dec{k+8} with $k = 0, 7, 14, \dots, 7 * (WDEC.level - 1)$ contain the 3-D wavelet coefficients for the multiresolution starting with the coarsest level when $k=0$.</p> <p>For example, if $WDEC.level=3$, dec{2}, ..., dec{8} contain the wavelet coefficients for level 3 ($k=0$), dec{9}, ..., dec{15} contain the wavelet coefficients for level 2 ($k=7$), and dec{16}, ..., dec{22} contain the wavelet coefficients for level 1 ($k=7 * (WDEC.level - 1)$).</p> <p>At each level, the wavelet coefficients in dec{k+2}, ..., dec{k+8} are in the following order: 'HLL', 'LHL', 'HHL', 'LLH', 'HLH', 'LHH', 'HHH'.</p> <p>The sequence of letters gives the order in which the separable filtering operations are applied from left to right. For example, 'LHH' means that the lowpass (scaling) filter with downsampling is applied to the rows of X, followed by the highpass (wavelet) filter with downsampling applied to the columns of X. Finally, the highpass filter with downsampling is applied to the 3rd dimension of X.</p>
<p>sizes</p>	<p>Successive sizes of the decomposition components</p>

Examples

3-D Wavelet Transform

Find the 3-D DWT of a volume. Construct 8-by-8-by-8 matrix of integers 1 to 64 and make the data 3-D.

```
M = magic(8);
X = repmat(M,[1 1 8]);
```

Obtain the 3-D discrete wavelet transform at level 1 using the Haar wavelet and the default whole-point symmetric extension mode.

```
wd1 = wavedec3(X,1,'db1');
```

3-D Wavelet Transform Using Specified Decomposition and Reconstruction Filters

Specify the decomposition and reconstruction filters as a cell array. Construct 8-by-8-by-8 matrix of integers 1 to 64 and make the data 3-D.

```
M = magic(8);
X = repmat(M,[1 1 8]);
```

Obtain the 3-D discrete wavelet transform down to level 2 using the Daubechies extremal phase wavelet with two vanishing moments. Input the decomposition and reconstruction filters as a cell array. Use the periodic extension mode.

```
[LoD,HiD,LoR,HiR] = wfilters('db2');
wd2 = wavedec3(X,2,{LoD,HiD,LoR,HiR},'mode','per');
```

Coefficient Order in 3-D Wavelet Transform

Compare the output of `wavedec3` and `dwt3` to illustrate the ordering of the 3-D wavelet coefficients described in the `dec` field description.

```
X = reshape(1:512,8,8,8);
dwtOut = dwt3(X,'db1','mode','per');
wdec = wavedec3(X,1,'db1','mode','per');
max(abs((wdec.dec{4}(:)-dwtOut.dec{2,2,1}(:))))

ans = 0

max(abs((wdec.dec{5}(:)-dwtOut.dec{1,1,2}(:))))

ans = 0
```

See Also

dwt3 | dwtmode | waveinfo | waverec3 | wfilters | wmaxlev

Introduced in R2010a

wavefun

Wavelet and scaling functions

Syntax

```
[PHI,PSI,XVAL] = wavefun('wname',ITER)
[PHI1,PSI1,PHI2,PSI2,XVAL] = wavefun('wname',ITER)
[PHI,PSI,XVAL] = wavefun('wname',ITER)
[PSI,XVAL] = wavefun('wname',ITER)
[...] = wavefun(wname,A,B)
[...] = wavefun('wname',max(A,B))
[...] = wavefun('wname',0)
[...] = wavefun('wname',8,0)
[...] = wavefun('wname')
[...] = wavefun('wname',8)
```

Description

The function `wavefun` returns approximations of the wavelet function '*wname*' and the associated scaling function, if it exists. The positive integer `ITER` determines the number of iterations computed; thus, the refinement of the approximations.

For an orthogonal wavelet:

`[PHI,PSI,XVAL] = wavefun('wname',ITER)` returns the scaling and wavelet functions on the points grid `XVAL`.

For a biorthogonal wavelet:

`[PHI1,PSI1,PHI2,PSI2,XVAL] = wavefun('wname',ITER)` returns the scaling and wavelet functions both for decomposition (`PHI1,PSI1`) and for reconstruction (`PHI2,PSI2`).

For a Meyer wavelet:

`[PHI,PSI,XVAL] = wavefun('wname',ITER)`

For a wavelet without scaling function (e.g., Morlet, Mexican Hat, Gaussian derivatives wavelets or complex wavelets):

```
[PSI,XVAL] = wavefun('wname',ITER)
```

[...] = wavefun(wname,A,B), where A and B are positive integers, is equivalent to [...]= wavefun('wname',max(A,B)), and draws plots.

When A is set equal to the special value 0,

- [...] = wavefun('wname',0) is equivalent to
- [...] = wavefun('wname',8,0).
- [...] = wavefun('wname') is equivalent to
- [...] = wavefun('wname',8).

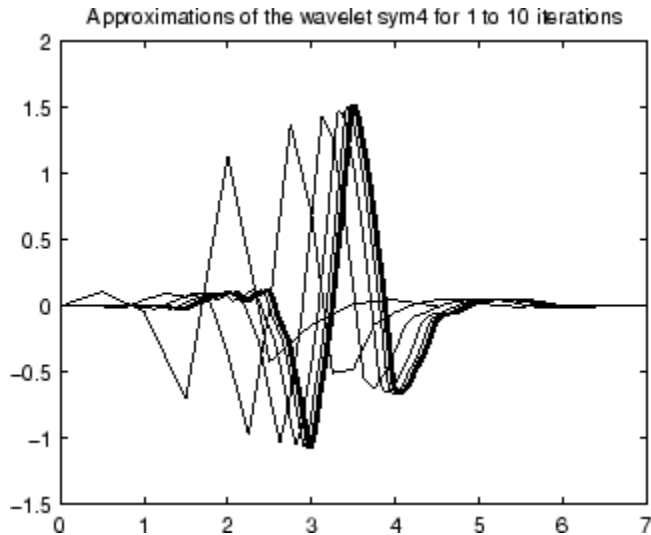
The output arguments are optional.

Examples

On the following graph, 10 piecewise linear approximations of the sym4 wavelet obtained after each iteration of the cascade algorithm are shown.

```
% Set number of iterations and wavelet name.
iter = 10;
wav = 'sym4';

% Compute approximations of the wavelet function using the
% cascade algorithm.
for i = 1:iter
    [phi,psi,xval] = wavefun(wav,i);
    plot(xval,psi);
    hold on
end
title(['Approximations of the wavelet ',wav, ...
' for 1 to ',num2str(iter),' iterations']);
hold off
```



Algorithms

For compactly supported wavelets defined by filters, in general no closed form analytic formula exists.

The algorithm used is the cascade algorithm. It uses the single-level inverse wavelet transform repeatedly.

Let us begin with the scaling function ϕ .

Since ϕ is also equal to $\phi_{0,0}$, this function is characterized by the following coefficients in the orthogonal framework:

- $\langle \phi, \phi_{0,n} \rangle = 1$ only if $n = 0$ and equal to 0 otherwise
- $\langle \phi, \psi_{-j,k} \rangle = 0$ for positive j , and all k .

This expansion can be viewed as a wavelet decomposition structure. Detail coefficients are all zeros and approximation coefficients are all zeros except one equal to 1.

Then we use the reconstruction algorithm to approximate the function ϕ over a dyadic grid, according to the following result:

For any dyadic rational of the form $x = n2^{-j}$ in which the function is continuous and where j is sufficiently large, we have pointwise convergence and

$$\left| \phi(x) - 2^{\frac{j}{2}} \langle \phi, \phi_{-j, n} 2^{j-j} \rangle \right| \leq C \cdot 2^{-j\alpha}$$

where C is a constant, and α is a positive constant depending on the wavelet regularity.

Then using a good approximation of ϕ on dyadic rationals, we can use piecewise constant or piecewise linear interpolations η on dyadic intervals, for which uniform convergence occurs with similar exponential rate:

$$\|\phi - \eta\|_{\infty} \leq C \cdot 2^{-j\alpha}$$

So using a J -step reconstruction scheme, we obtain an approximation that converges exponentially towards ϕ when J goes to infinity.

Approximations are computed over a grid of dyadic rationals covering the support of the function to be approximated.

Since a scaled version of the wavelet function ψ can also be expanded on the $(\phi_{-1, n})_n$, the same scheme can be used, after a single-level reconstruction starting with the appropriate wavelet decomposition structure. Approximation coefficients are all zeros and detail coefficients are all zeros except one equal to 1.

For biorthogonal wavelets, the same ideas can be applied on each of the two multiresolution schemes in duality.

Note This algorithm may diverge if the function to be approximated is not continuous on dyadic rationals.

References

Daubechies, I., *Ten lectures on wavelets*, CBMS, SIAM, 1992, pp. 202–213.

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

See Also

`intwave` | `waveinfo` | `wfilters`

Introduced before R2006a

wavefun2

Wavelet and scaling functions 2-D

Syntax

```
[PHI,PSI,XVAL] = wavefun('wname',ITER)
[S,W1,W2,W3,XYVAL] = wavefun2('wname',ITER,'plot')
[S,W1,W2,W3,XYVAL] = wavefun2(wname,A,B)
[S,W1,W2,W3,XYVAL] = wavefun2('wname',max(A,B))
[S,W1,W2,W3,XYVAL] = wavefun2('wname',0)
[S,W1,W2,W3,XYVAL] = wavefun2('wname',4,0)
[S,W1,W2,W3,XYVAL] = wavefun2('wname')
[S,W1,W2,W3,XYVAL] = wavefun2('wname',4)
```

Description

For an orthogonal wavelet '*wname*', `wavefun2` returns the scaling function and the three wavelet functions resulting from the tensor products of the one-dimensional scaling and wavelet functions.

If `[PHI,PSI,XVAL] = wavefun('wname',ITER)`, the scaling function *S* is the tensor product of *PHI* and *PSI*.

The wavelet functions *W1*, *W2*, and *W3* are the tensor products (*PHI,PSI*), (*PSI,PHI*), and (*PSI,PSI*), respectively.

The two-dimensional variable *XYVAL* is a $2^{\text{ITER}} \times 2^{\text{ITER}}$ points grid obtained from the tensor product (*XVAL,XVAL*).

The positive integer *ITER* determines the number of iterations computed and thus, the refinement of the approximations.

`[S,W1,W2,W3,XYVAL] = wavefun2('wname',ITER,'plot')` computes and also plots the functions.

$[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname, A, B)$, where A and B are positive integers, is equivalent to $[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname, \max(A, B))$. The resulting functions are plotted.

When A is set equal to the special value 0,

- $[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname, 0)$ is equivalent to $[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname, 4, 0)$.
- $[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname)$ is equivalent to $[S, W1, W2, W3, XYVAL] = \text{wavefun2}(wname, 4)$.

The output arguments are optional.

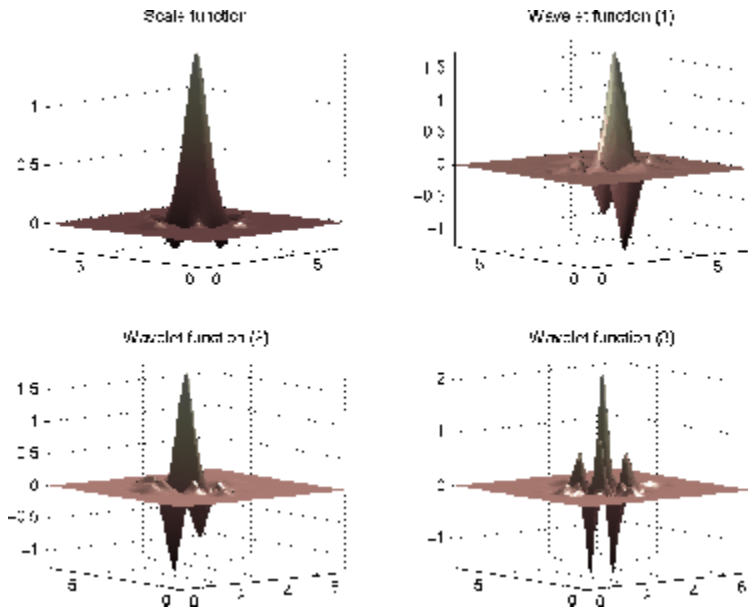
Note The `wavefun2` function can only be used with an orthogonal wavelet.

Examples

On the following graph, a linear approximation of the `sym4` wavelet obtained using the cascade algorithm is shown.

```
% Set number of iterations and wavelet name.
iter = 4;
wav = 'sym4';

% Compute approximations of the wavelet and scale functions using
% the cascade algorithm and plot.
[s,w1,w2,w3,xyval] = wavefun2(wav,iter,0);
```



Algorithms

See `wavefun` for more information.

References

Daubechies, I., *Ten lectures on wavelets*, CBMS, SIAM, 1992, pp. 202–213.

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

See Also

`intwave` | `wavefun` | `waveinfo` | `wfilters`

Introduced before R2006a

waveinfo

Wavelets information

Syntax

`waveinfo('wname')`

Description

`waveinfo` provides information on all wavelets within the toolbox.

`waveinfo('wname')` provides information on the wavelet family whose short name is specified by `'wname'`. Available family short names are listed in the table below.

Wavelet Family Short Name	Wavelet Family Name
'haar'	Haar wavelet
'db'	Daubechies wavelets
'sym'	Symlets
'coif'	Coiflets
'bior'	Biorthogonal wavelets
'fk'	Fejer-Korovkin filters
'rbio'	Reverse biorthogonal wavelets
'meyr'	Meyer wavelet
'dmey'	Discrete approximation of Meyer wavelet
'gaus'	Gaussian wavelets
'mexh'	Mexican hat wavelet (also known as Ricker wavelet)
'morl'	Morlet wavelet
'cgau'	Complex Gaussian wavelets

Wavelet Family Short Name	Wavelet Family Name
'shan'	Shannon wavelets
'fbsp'	Frequency B-Spline wavelets
'cmor'	Complex Morlet wavelets

The family short names can also be user-defined ones (see `wavemngr` for more information).

`waveinfo('wsys')` provides information on wavelet packets.

Examples

```
waveinfo('db')
```

DBINFO Information on Daubechies wavelets.

Daubechies Wavelets

General characteristics: Compactly supported wavelets with extremal phase and highest number of vanishing moments for a given support width. Associated scaling filters are minimum-phase filters.

```
Family           Daubechies
Short name       db
Order N          N strictly positive integer
Examples         db1 or haar, db4, db15
```

```
Orthogonal       yes
Biorthogonal     yes
Compact support  yes
DWT              possible
CWT              possible
```

```
Support width    2N-1
Filters length   2N
Regularity       about 0.2 N for large N
Symmetry         far from
Number of vanishing moments for psi    N
```

Reference: I. Daubechies,
Ten lectures on wavelets CBMS, SIAM, 61, 1994, 194-202.

See Also

wavemngr

Introduced before R2006a

Wavelet Analyzer

Analyze signals and images using wavelets

Description

The **Wavelet Analyzer** app is an interactive tool for using wavelets to visualize and analyze signals and images. Using this app, you can:

- Perform wavelet and wavelet packet analysis
- Denoise and compress signals and images
- Design custom wavelets
- Estimate density and regression
- Perform matching pursuit analysis
- Perform image fusion

Open the Wavelet Analyzer App

- MATLAB Toolstrip: On the **Apps** tab, under **Signal Processing and Communications**, click the app icon.
- MATLAB command prompt: Enter `waveletAnalyzer`

Examples

- “Continuous Wavelet Analysis of Noisy Sinusoid Using the Wavelet Analyzer App”
- “DFT-Based Continuous Wavelet Analysis Using the Wavelet Analyzer App”
- “Interactive 1-D Stationary Wavelet Transform Denoising”
- “Matching Pursuit Using Wavelet Analyzer App”

Definitions

Boundary Conditions

To change the way **Wavelet Analyzer** handles boundary conditions, use the `dwtmode` function.

See Also

Topics

"Continuous Wavelet Analysis of Noisy Sinusoid Using the Wavelet Analyzer App"

"DFT-Based Continuous Wavelet Analysis Using the Wavelet Analyzer App"

"Interactive 1-D Stationary Wavelet Transform Denoising"

"Matching Pursuit Using Wavelet Analyzer App"

"Continuous Wavelet Analysis"

"Discrete Wavelet Analysis"

Introduced before R2006a

waveletfamilies

Wavelet families and family members

Syntax

```
waveletfamilies('f')  
waveletfamilies('n')  
waveletfamilies('a')
```

Description

`waveletfamilies` or `waveletfamilies('f')` displays the names of all available wavelet families.

`waveletfamilies('n')` displays the names of all available wavelets in each family.

`waveletfamilies('a')` displays all available wavelet families with their corresponding properties.

Examples

Wavelet Families

Display the names of all available wavelet families.

```
waveletfamilies
```

```
=====
Haar                haar
Daubechies          db
Symlets             sym
Coiflets            coif
BiorSplines         bior
ReverseBior         rbio
Meyer               meyr
```

```

DMeyer          dmey
Gaussian        gaus
Mexican_hat     mexh
Morlet          morl
Complex Gaussian cgau
Shannon         shan
Frequency B-Spline fbsp
Complex Morlet  cmor
Fejer-Korovkin fk

```

```
=====
```

Display the names of all available wavelets in each family.

```
waveletfamilies('n')
```

```

=====
Haar          haar
=====
Daubechies    db
-----
db1   db2   db3   db4
db5   db6   db7   db8
db9   db10  db**
=====
Symlets       sym
-----
sym2   sym3   sym4   sym5
sym6   sym7   sym8   sym**
=====
Coiflets      coif
-----
coif1   coif2   coif3   coif4
coif5
=====
BiorSplines   bior
-----
bior1.1  bior1.3  bior1.5  bior2.2
bior2.4  bior2.6  bior2.8  bior3.1
bior3.3  bior3.5  bior3.7  bior3.9
bior4.4  bior5.5  bior6.8
=====
ReverseBior    rbio
-----
rbio1.1  rbio1.3  rbio1.5  rbio2.2

```

```
rbio2.4    rbio2.6    rbio2.8    rbio3.1
rbio3.3    rbio3.5    rbio3.7    rbio3.9
rbio4.4    rbio5.5    rbio6.8
=====
Meyer                      meyr
=====
DMeyer                    dmey
=====
Gaussian                  gaus
-----
gaus1    gaus2    gaus3    gaus4
gaus5    gaus6    gaus7    gaus8
=====
Mexican_hat              mexh
=====
Morlet                    morl
=====
Complex Gaussian        cgau
-----
cgau1    cgau2    cgau3    cgau4
cgau5    cgau6    cgau7    cgau8
=====
Shannon                  shan
-----
shan1-1.5    shan1-1    shan1-0.5    shan1-0.1
shan2-3    shan**
=====
Frequency B-Spline      fbsp
-----
fbsp1-1-1.5    fbsp1-1-1    fbsp1-1-0.5    fbsp2-1-1
fbsp2-1-0.5    fbsp2-1-0.1    fbsp**
=====
Complex Morlet          cmor
-----
cmor1-1.5    cmor1-1    cmor1-0.5    cmor1-1
cmor1-0.5    cmor1-0.1    cmor**
=====
Fejer-Korovkin         fk
-----
fk4    fk6    fk8    fk14
fk18    fk22
=====
```

Display all available wavelet families with their corresponding properties.

```
waveletfamilies('a')
```

```
Type of Wavelets
```

```
-----  
type = 1   - orthogonal wavelets           (F.I.R.)  
type = 2   - biorthogonal wavelets        (F.I.R.)  
type = 3   - with scale function  
type = 4   - without scale function  
type = 5   - complex wavelet.  
-----
```

```
-----  
Family Name : Haar
```

```
haar  
1  
no  
no  
dbwavf
```

```
-----  
Family Name : Daubechies
```

```
db  
1  
1 2 3 4 5 6 7 8 9 10 **  
integer  
dbwavf
```

```
-----  
Family Name : Symlets
```

```
sym  
1  
2 3 4 5 6 7 8 **  
integer  
symwavf
```

```
-----  
Family Name : Coiflets
```

```
coif  
1  
1 2 3 4 5  
integer  
coifwavf
```

```
-----  
Family Name : BiorSplines  
bior  
2  
1.1 1.3 1.5 2.2 2.4 2.6 2.8 3.1 3.3 3.5 3.7 3.9 4.4 5.5 6.8  
real  
biorwavf
```

```
-----  
Family Name : ReverseBior  
rbio  
2  
1.1 1.3 1.5 2.2 2.4 2.6 2.8 3.1 3.3 3.5 3.7 3.9 4.4 5.5 6.8  
real  
rbiowavf
```

```
-----  
Family Name : Meyer  
meyr  
3  
no  
no  
meyer  
-8 8
```

```
-----  
Family Name : DMeyer  
dmey  
1  
no  
no  
dmey.mat
```

```
-----  
Family Name : Gaussian  
gaus  
4  
1 2 3 4 5 6 7 8  
integer  
gauswavf  
-5 5
```

```
-----  
Family Name : Mexican_hat  
mexh  
4
```

```
no
no
mexihat
-8 8
-----
Family Name : Morlet
morl
4
no
no
morlet
-8 8
-----
Family Name : Complex Gaussian
cgau
5
1 2 3 4 5 6 7 8
integer
cgauwavf
-5 5
-----
Family Name : Shannon
shan
5
1-1.5 1-1 1-0.5 1-0.1 2-3 **
string
shanwavf
-20 20
-----
Family Name : Frequency B-Spline
fbsp
5
1-1-1.5 1-1-1 1-1-0.5 2-1-1 2-1-0.5 2-1-0.1 **
string
fbspwavf
-20 20
-----
Family Name : Complex Morlet
cmor
5
1-1.5 1-1 1-0.5 1-1 1-0.5 1-0.1 **
string
cmorwavf
-8 8
```

```
-----  
Family Name : Fejer-Korovkin  
fk  
1  
4 6 8 14 18 22  
integer  
fejerkorovkin  
-----
```

See Also

wavemngr

Introduced in R2008a

wavelets

CWT filter bank time-domain wavelets

Syntax

```
psi = wavelets(fb)  
[psi,t] = wavelets(fb)
```

Description

`psi = wavelets(fb)` returns the time-domain wavelets, `psi`, for the continuous wavelet transform (CWT) filter bank, `fb`. The time-domain wavelets are centered at the origin.

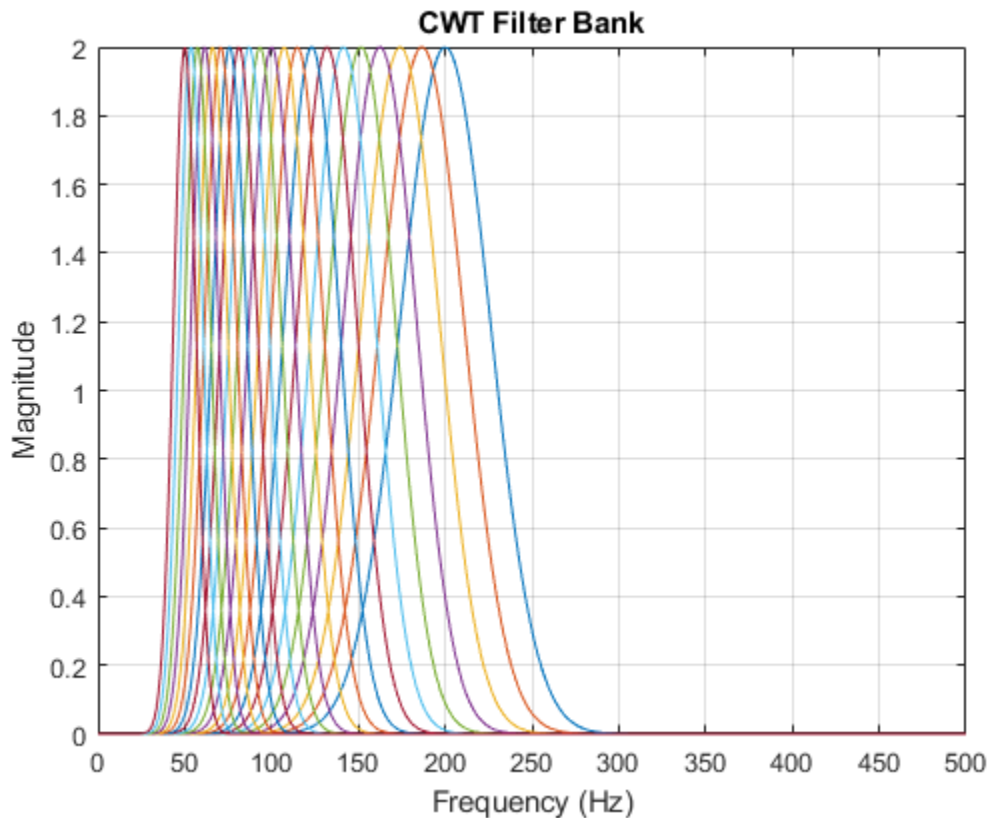
`[psi,t] = wavelets(fb)` returns the sampling instants, `t`, for the wavelets.

Examples

Filter Bank Time Domain Wavelets

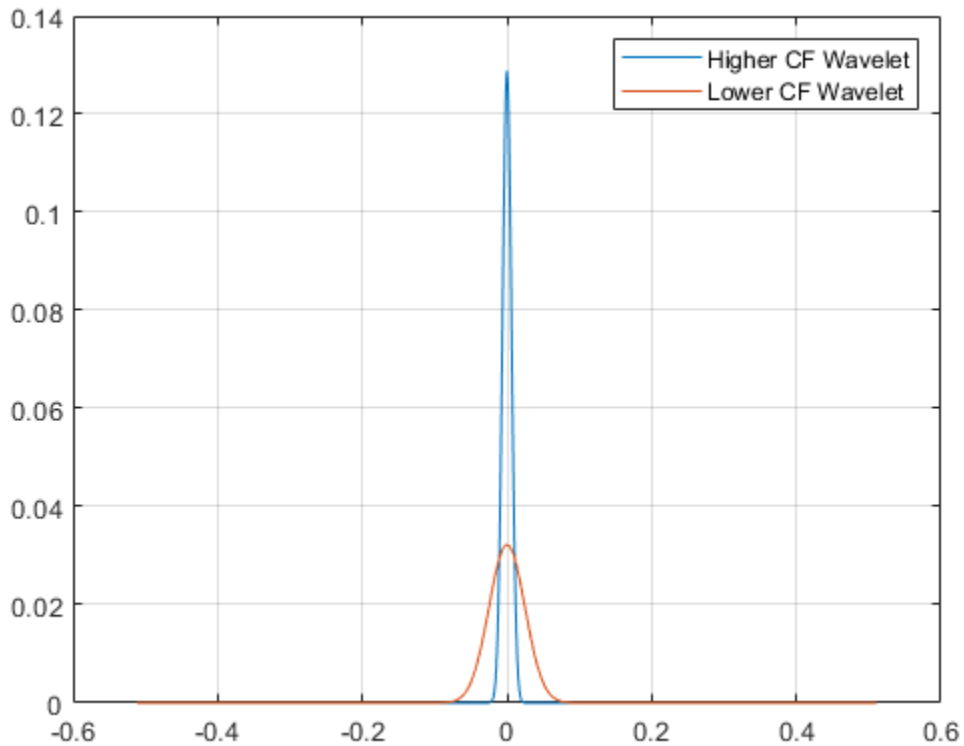
Create a continuous wavelet transform filter bank. Set the sampling frequency to 1000 Hz and the frequency limits to range from 50 Hz to 200 Hz. Plot the frequency response.

```
fb = cwtfilterbank('SamplingFrequency',1000,'FrequencyLimits',[50 200]);  
freqz(fb)
```



Obtain the filter bank time domain wavelets. Plot the magnitudes of the first and last wavelets contained in the output. The first wavelet corresponds to the wavelet filter with center frequency equal to 200 Hz, and the last wavelet corresponds to the wavelet filter with center frequency equal to 50 Hz.

```
[psi,t] = wavelets(fb);  
figure  
plot(t,abs(psi(1,:)))  
hold on  
plot(t,abs(psi(end,:)))  
legend('Higher CF Wavelet','Lower CF Wavelet')  
grid on
```



Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

Output Arguments

psi — Time-domain wavelets

complex-valued matrix

Time-domain wavelets, returned as a N_s -by- N complex-valued matrix, where N_s is the number of wavelet bandpass frequencies (equal to the number of scales) and N is the filter bank `SignalLength`. The wavelets are ordered in `psi` from the highest-frequency passband filter to the lowest-frequency passband filter.

t — Sampling instants

vector

Sampling instants of the time-domain wavelets, returned as a real-valued vector of length N where N is the filter bank `SignalLength`. `t` is the same data type as the `SamplingPeriod`.

See Also

`cwtfilterbank` | `waveletsupport`

Introduced in R2018a

wavelets

DWT filter bank time-domain wavelets

Syntax

```
psi = wavelets(fb)  
[psi,t] = wavelets(fb)
```

Description

`psi = wavelets(fb)` returns the time-domain and centered wavelets corresponding to the wavelet passband filters in `fb`. `psi` is an L-by-N matrix where L is the filter bank Level and N is the filter bank SignalLength.

`[psi,t] = wavelets(fb)` returns the sampling instants, `t`. Sampling instants lie in the interval $-(N*DT)/2 \leq t < (N*DT)/2$, where N is signal length and DT is the filter bank sampling period (reciprocal of the filter bank sampling frequency).

Examples

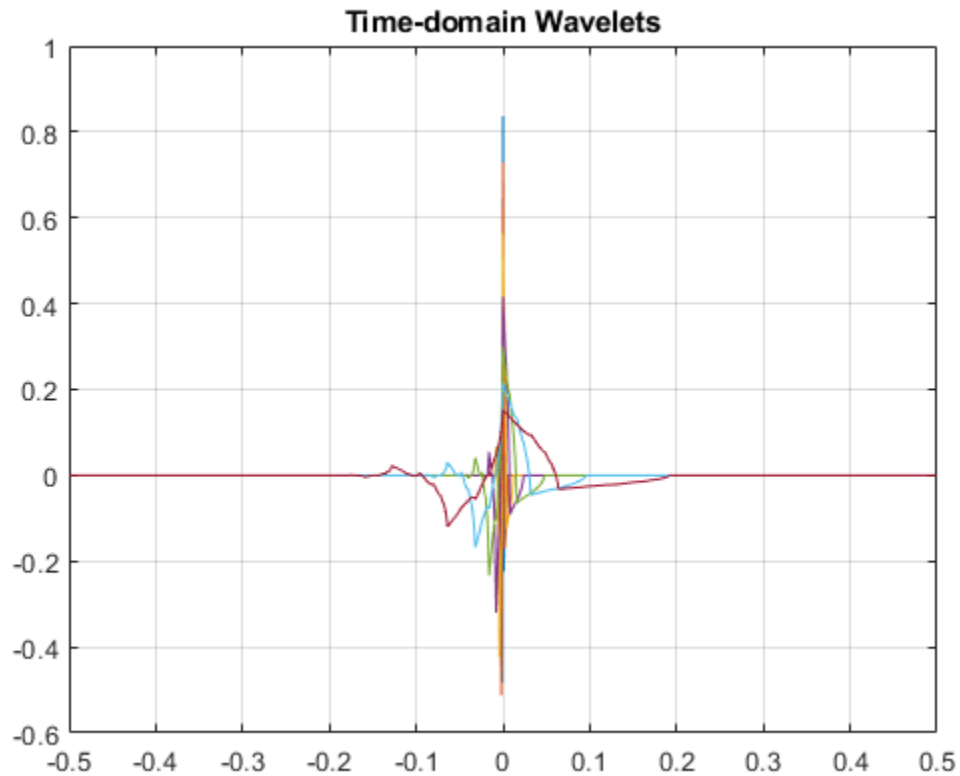
DWT Filter Bank Wavelets

Create a seven level DWT filter bank with a signal length of 1000 samples, using the Daubechies db2 wavelet and a sampling frequency of 1kHz.

```
wv = "db2";  
len = 1000;  
lev = 7;  
Fs = 1e3;  
fb = dwtfilterbank('Wavelet',wv,'SignalLength',len,'Level',lev,'SamplingFrequency',Fs)
```

Obtain and plot the time-domain and centered wavelets corresponding to the wavelet bandpass filters.

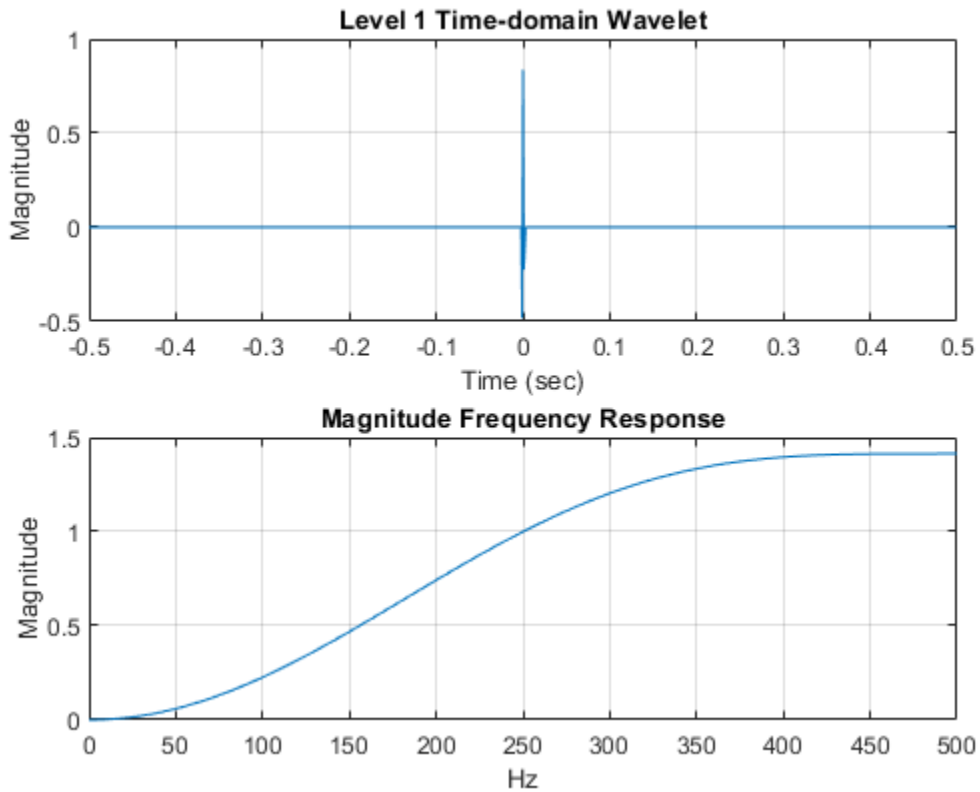
```
[psi,t] = wavelets(fb);  
plot(t,psi')  
grid on  
title('Time-domain Wavelets')
```



Plot the finest scale time-domain wavelet and the one-sided magnitude frequency response of the corresponding wavelet bandpass filter.

```
sc = 1;  
[psidft,f] = freqz(fb);  
subplot(2,1,1)  
plot(t,psi(sc,:))  
grid on  
xlabel('Time (sec)')
```

```
ylabel('Magnitude')
title(['Level ', num2str(sc), ' Time-domain Wavelet'])
subplot(2,1,2)
plot(f(len/2:end),abs(psidft(sc,len/2:end)))
grid on
xlabel('Hz')
ylabel('Magnitude')
title('Magnitude Frequency Response')
```



Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

Output Arguments

psi — Time-centered wavelets

real-valued matrix

Time-centered wavelets corresponding to the wavelet passband filters, returned as an L-by-N matrix, where L is the filter bank `Level` and N is the `SignalLength`. The wavelets are ordered in `psi` from the finest scale resolution to the coarsest scale resolution.

t — Sampling instants

real-valued vector

Sampling instants, returned as a real-valued vector `t` of length N, where N is the filter bank `SignalLength`. Sampling instants lie in the interval $-(N*DT)/2 \leq t < (N*DT)/2$, where DT is the filter bank sampling period (reciprocal of the filter bank sampling frequency).

See Also

`dwtfilterbank` | `scalingfunctions`

Introduced in R2018a

waveletScattering

Wavelet time scattering

Description

Use the `waveletScattering` object to create a framework for a wavelet time scattering decomposition using the Gabor (analytic Morlet) wavelet. The framework uses wavelets and a lowpass scaling function to generate low-variance representations of real-valued time series data. Wavelet time scattering yields representations insensitive to translations in the input signal without sacrificing class discriminability. You can use the framework to extract features that can be used in many learning algorithms, such as k-nearest neighbors (kNN), support vector machines (SVM), and boosted decision trees. You can also use the features in your own classifiers. You can specify the duration of translation invariance and the number of wavelet filters per octave.

Creation

Syntax

```
sf = waveletScattering  
sf = waveletScattering(Name,Value)
```

Description

`sf = waveletScattering` creates a framework for a wavelet time scattering decomposition with two filter banks. The first filter bank has a quality factor of eight wavelets per octave. The second filter bank has a quality factor of one wavelet per octave. By default, `waveletScattering` assumes a signal input length of 1024 samples. The scale invariance length is 512 samples. By default, `waveletScattering` uses periodic boundary conditions.

`sf = waveletScattering(Name,Value)` creates a framework for wavelet scattering, `sf`, with properties specified by one or more `Name, Value` pair arguments. Properties can

be specified in any order as `Name1, Value1, . . . , NameN, ValueN`. Enclose each property name in quotes.

Note With the exception of `OversamplingFactor`, after creation you cannot change a property value of an existing scattering framework. For example, if you have a framework `sf` with a `SignalLength` of 2000, you must create a second framework `sf2` for a signal with 2001 samples. You cannot assign a different `SignalLength` to `sf`.

Properties

SignalLength — Signal length

1024 (default) | positive integer ≥ 16

Signal length, specified as a positive integer ≥ 16 .

Data Types: double

SamplingFrequency — Sampling frequency

1 (default) | positive scalar

Sampling frequency in hertz, specified as a positive scalar. If unspecified, frequencies are in cycles/sample and the Nyquist frequency is $\frac{1}{2}$.

Data Types: double

InvarianceScale — Scattering transform invariance scale

one-half of `SignalLength` (default) | positive scalar

Scattering transform invariance scale, specified as a positive scalar. `InvarianceScale` specifies the translation invariance of the scattering transform. If you do not specify `SamplingFrequency`, `InvarianceScale` is measured in samples. If you specify `SamplingFrequency`, `InvarianceScale` is measured in seconds. By default, `InvarianceScale` is one-half the `SignalLength` in samples.

`InvarianceScale` cannot exceed `SignalLength` in samples.

Example: `sf =`

```
waveletScattering('SignalLength',1000,'SamplingFrequency',200,'InvarianceScale',5)
```

 has the largest possible `InvarianceScale`.

Data Types: double

QualityFactors — Scattering filter bank Q factors

[8 1] (default) | positive integer | vector of positive integers

Scattering filter bank Q factors, specified as a positive integer or a vector of positive integers. A filter bank Q factor is the number of wavelet filters per octave. Quality factors cannot exceed 32 and must be greater than or equal to 1.

If `QualityFactors` is specified as a vector, the elements of `QualityFactors` must be strictly decreasing.

Example: `sf = waveletScattering('QualityFactors',[8 2 1])` creates a wavelet scattering framework with three filter banks.

Data Types: double

Boundary — Signal extension method

'periodic' (default) | 'reflection'

Signal extension method to apply at the boundary:

- 'periodic' — Extend signal periodically to length $2^{\text{ceil}(\log_2(N))}$, where N is the signal length.
- 'reflection' — Extend signal by reflection to length $2^{\text{ceil}(\log_2(2N))}$, where N is the signal length.

The signal is extended to match the length of the wavelet filters. The length is always power of two.

The signal extension method is for internal operations. Results are downsampled back onto the scale of the original signal before being returned.

Precision — Precision of scattering decomposition

'double' (default) | 'single'

Precision of scattering decomposition:

- 'double' — Double precision
- 'single' — Single precision

Note

- All calculations involving the wavelet scattering framework are carried out in Precision. Wavelet scattering functions such as `featureMatrix` and `filterbank` return outputs such as filters in Precision.
 - The precision of the output of the `scatteringTransform` function does not exceed the precision of `sf`.
-

OversamplingFactor — Oversampling factor

0 (default) | nonnegative integer | Inf

Oversampling factor, specified as a nonnegative integer or `Inf`. The factor specifies how much the scattering coefficients are oversampled with respect to the critically downsampled values. The factor is on a \log_2 scale. By default, `OversamplingFactor` is set to 0, which corresponds to critically downsampling the coefficients. You can use `numCoefficients` to determine the number of coefficients obtained for a scattering framework. To obtain a fully undecimated scattering transform, set `OversamplingFactor` to `Inf`.

Setting `OversamplingFactor` to a value that would result in more coefficients than samples is equivalent to setting `OversamplingFactor` to `Inf`. Increasing the `OversamplingFactor` significantly increases the computational complexity and memory requirements of the scattering transform.

Example: If `sf = waveletScattering('OversamplingFactor',2)`, the scattering transform returns 2^2 times as many coefficients for each scattering path with respect to the critically sampled number.

Object Functions

<code>scatteringTransform</code>	Wavelet 1-D scattering transform
<code>featureMatrix</code>	Scattering feature matrix
<code>log</code>	Natural logarithm of scattering transform
<code>filterbank</code>	Wavelet time scattering filter banks
<code>littlewoodPaleySum</code>	Littlewood-Paley sum
<code>scattergram</code>	Visualize scattering or scalogram coefficients
<code>centerFrequencies</code>	Wavelet scattering bandpass center frequencies
<code>numorders</code>	Number of scattering orders
<code>numfilterbanks</code>	Number of scattering filter banks
<code>numCoefficients</code>	Number of wavelet scattering coefficients

Examples

Wavelet Time Scattering with Default Values

Create a scattering framework with default values.

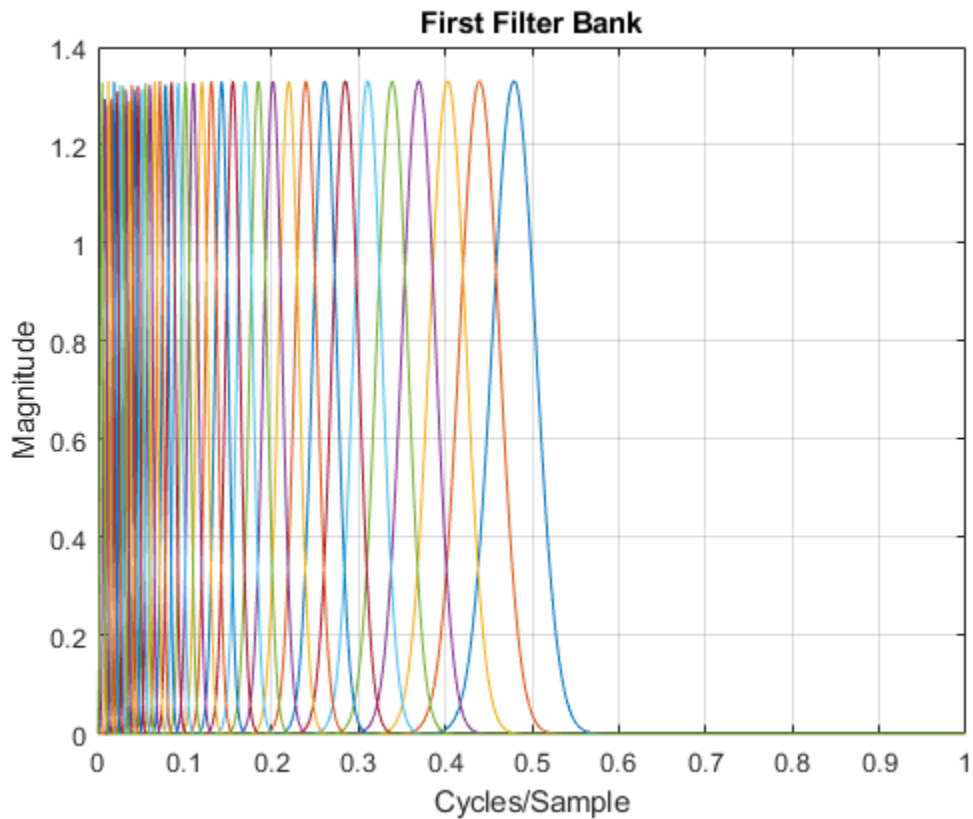
```
sf = waveletScattering

sf =
  waveletScattering with properties:

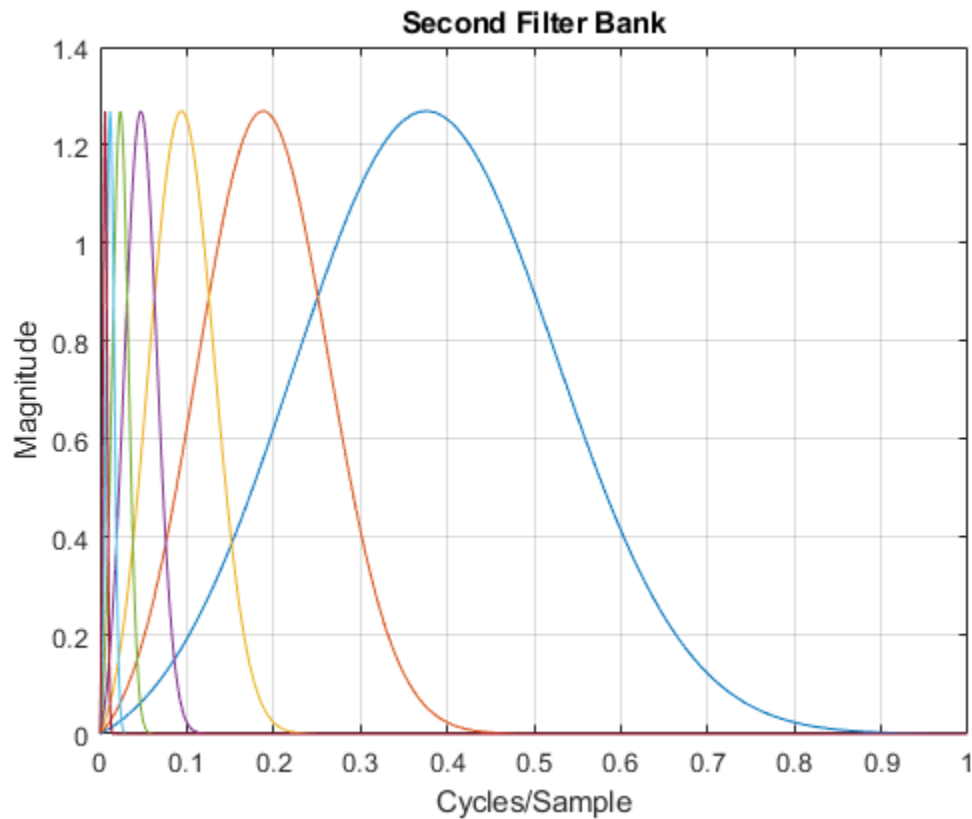
    SignalLength: 1024
    InvarianceScale: 512
    QualityFactors: [8 1]
    Boundary: 'periodic'
    SamplingFrequency: 1
    Precision: 'double'
    OversamplingFactor: 0
```

Plot the wavelet filters used in the first and second filter banks.

```
[filters,f] = filterbank(sf);
plot(f,filters{2}.psift)
title('First Filter Bank')
xlabel('Cycles/Sample')
ylabel('Magnitude')
grid on
```

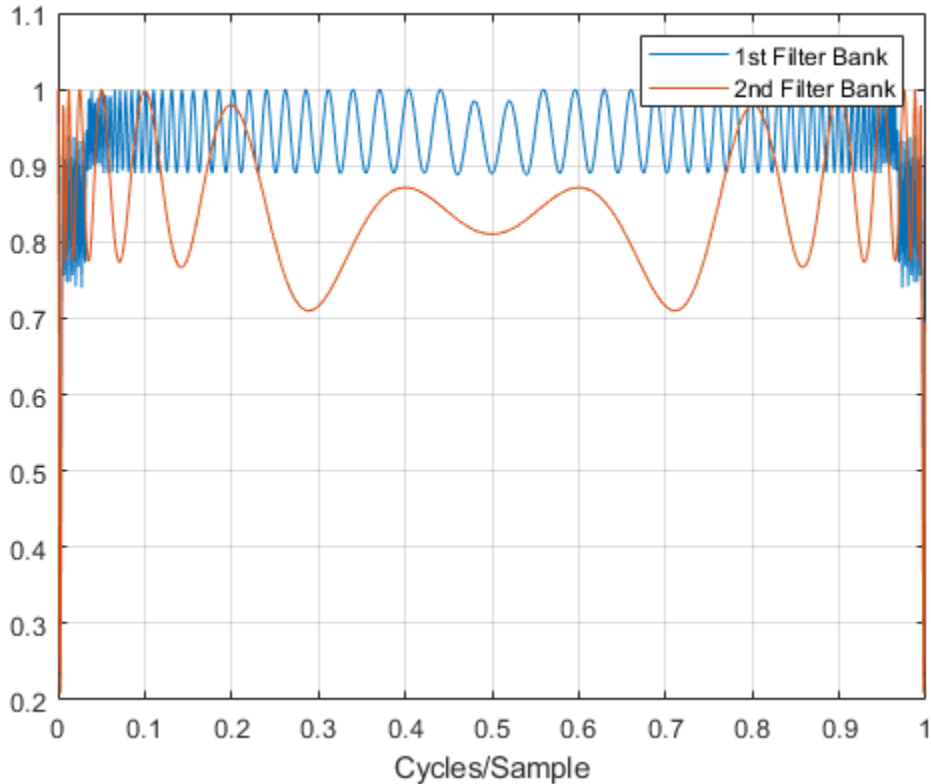


```
figure
plot(f, filters{3}.psift)
title('Second Filter Bank')
xlabel('Cycles/Sample')
ylabel('Magnititude')
grid on
```



Plot the Littlewood-Paley sums of the filter banks.

```
[lpsum,f] = littlewoodPaleySum(sf);  
figure  
plot(f,lpsum)  
legend('1st Filter Bank','2nd Filter Bank')  
xlabel('Cycles/Sample')  
grid on
```



Apply Scattering Decomposition Framework to Data

This example shows how to create and apply a scattering framework with three filter banks to data.

Load in a data set. Create a framework with three filter banks that can be applied to the data.

```
load handel
disp(['Data Sampling Frequency: ', num2str(Fs), ' Hz'])
```

```
Data Sampling Frequency: 8192 Hz
```

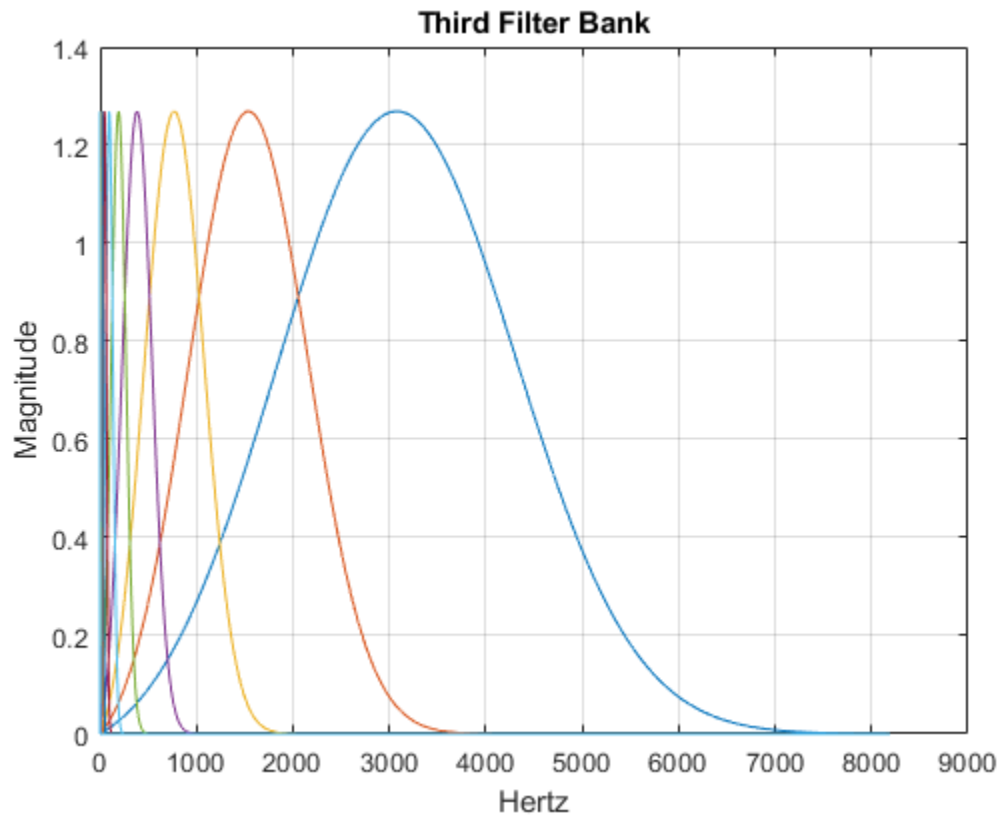


```
sf = waveletScattering('SignalLength', numel(y), ...  
    'SamplingFrequency', Fs, ...  
    'QualityFactors', [4 2 1])
```

```
sf =  
    waveletScattering with properties:  
  
        SignalLength: 73113  
        InvarianceScale: 4.4625  
        QualityFactors: [4 2 1]  
        Boundary: "periodic"  
        SamplingFrequency: 8192  
        Precision: "double"  
        OversamplingFactor: 0
```

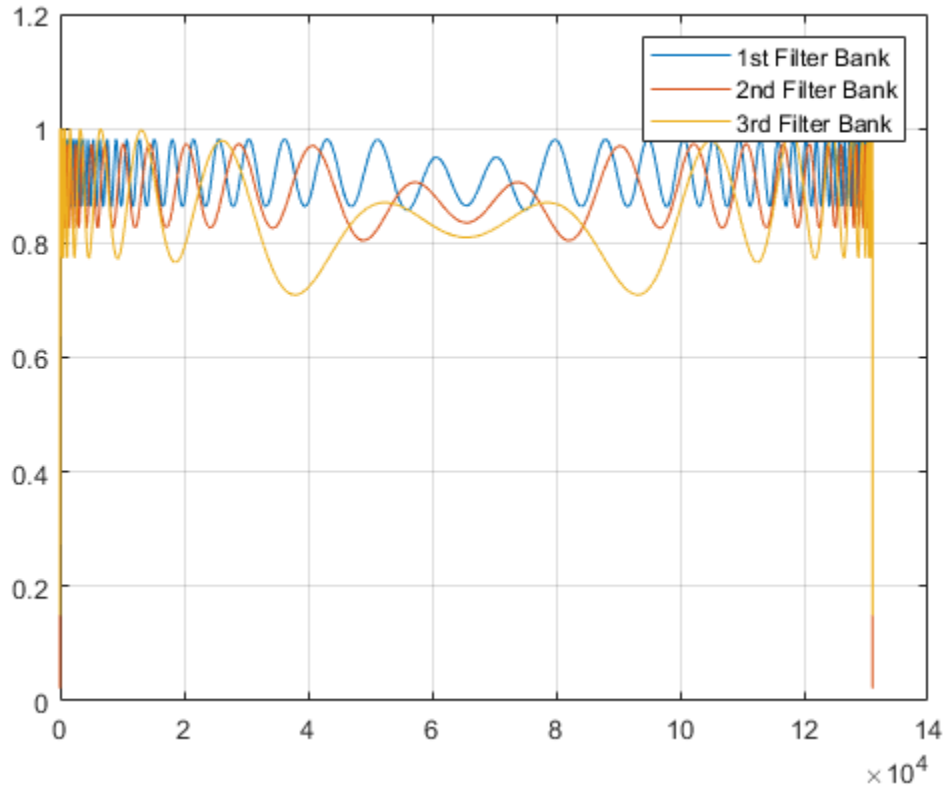
Inspect the framework. Plot the wavelet filters used in the third filter bank.

```
[filters,f] = filterbank(sf);  
plot(f, filters{4}.psift)  
grid on  
title('Third Filter Bank')  
xlabel('Hertz')  
ylabel('Magnitude')
```



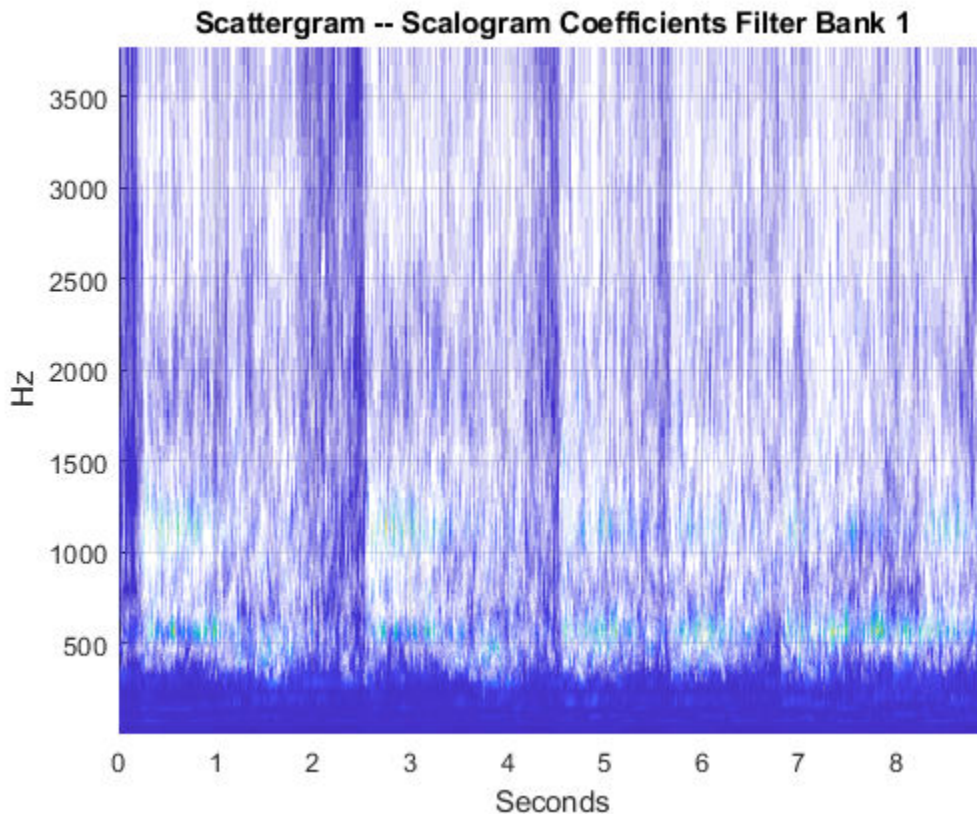
Plot the Littlewood-Paley sums of the three filter banks.

```
lpsum = littlewoodPaleySum(sf);  
figure  
plot(lpsum)  
grid on  
legend('1st Filter Bank', '2nd Filter Bank', '3rd Filter Bank')
```



Calculate the wavelet 1-D scattering transform of the data for `sf`. Visualize the scattergram of the scalogram coefficients for the first filter bank.

```
[S,U] = scatteringTransform(sf,y);  
figure  
scattergram(sf,U, 'FilterBank',1)
```



Compatibility Considerations

waveletScattering property Decimate will be removed

Not recommended starting in R2019a

The waveletScattering property Decimate will be removed in a future release. Use the property OversamplingFactor instead.

Functionality	What Happens When You Use This Functionality?	Use This Instead	Compatibility Considerations
Decimate	Still runs	Use <code>OversamplingFactor</code> or	<ul style="list-style-type: none"> • Replace all instances of 'Decimate', true with 'OversamplingFactor', 0. • Replace all instances of 'Decimate', false with 'OversamplingFactor', Inf.

References

- [1] Andén, J., and S. Mallat. "Deep Scattering Spectrum." *IEEE Transactions on Signal Processing*. Vol. 62, Number 16, 2014, pp. 4114–4128.
- [2] Mallat, S. "Group Invariant Scattering." *Communications in Pure and Applied Mathematics*. Vol. 65, Number 10, 2012, pp. 1331–1398.

See Also

Functions

cwt | cwtfilterbank

Topics

"Wavelet Scattering"

"Music Genre Classification Using Wavelet Time Scattering"

"Wavelet Time Scattering for ECG Signal Classification"

"Wavelet Time Scattering Classification of Phonocardiogram Data"

Introduced in R2018b

waveletScattering2

Wavelet image scattering

Description

Use the `waveletScattering2` object to create a framework for a wavelet image scattering decomposition using complex-valued 2-D Morlet wavelets.

Creation

Syntax

```
sf = waveletScattering2  
sf = waveletScattering2(Name,Value)
```

Description

`sf = waveletScattering2` creates a framework for a wavelet image scattering decomposition with two complex-valued 2-D Morlet filter banks and isotropic scale invariance. Both filter banks have quality factors of one wavelet per octave. There are six rotations linearly spaced between 0 and π radians for each wavelet filter. By default, `waveletScattering2` assumes an image input size of 128-by-128. The scale invariance is 64.

`sf = waveletScattering2(Name,Value)` creates a framework for wavelet image scattering with properties specified by one or more `Name,Value` pair arguments. Properties can be specified in any order as `Name1,Value1,...,NameN,ValueN`. Enclose each property name in single quotes (' ') or double quotes (" ").

Note With the exceptions of `OptimizePath` and `OversamplingFactor`, you cannot change a property value of an existing scattering framework. For example, if you create a framework `sf` with `ImageSize` set to `[256 256]`, you cannot assign a different `ImageSize` to `sf`.

Properties

ImageSize — Image size

[128 128] (default) | two-element integer-valued vector

Image size for wavelet image scattering framework, specified as a two-element integer-valued vector [*numrows numcolumns*]. Images must be at least 10-by-10.

If your input is an RGB image, you do not have to specify the third dimension. `waveletScattering2` only supports color images where the size of the third dimension is 3.

Example: `sf = waveletScattering2('ImageSize',[100 200])` creates a framework for 100-by-200 images and 100-by-200-by-3 color images.

InvarianceScale — Scattering transform invariance scale

64 (default) | positive scalar

Scattering transform invariance scale, specified as a positive scalar. `InvarianceScale` specifies the spatial support in the row and column dimensions of the scaling filter. `InvarianceScale` cannot exceed the minimum size of the row and column dimensions of the image.

By default, `InvarianceScale` is one-half the minimum of the row and column sizes of the image rounded to the nearest integer.

Example: `sf = waveletScattering2('ImageSize',[101 200])` creates a framework with `InvarianceScale` equal to 51.

NumRotations — Number of rotations per wavelet

[6 6] (default) | integer-valued vector

Number of rotations per wavelet per filter bank in the scattering framework, specified as an integer-valued vector. Specify one integer less than or equal to 12 for each filter bank in the scattering framework.

For each wavelet in each filter bank, there are `NumRotations` linearly spaced angles between 0 and π radians. The wavelet is rotated in a clockwise direction. The length of the vector specified in `NumRotations` must equal the length of the vector specified in `QualityFactors`.

Example: `sf = waveletScattering2('NumRotations',[7 5])` creates a framework with seven rotations per wavelet in the first filter bank and five rotations per wavelet in the second filter bank.

Note The 2-D wavelet scattering framework is constructed by rotating the 2-D Morlet wavelets in a clockwise direction. The opposite convention is used in the Image Processing Toolbox™. Creating a Gabor filter bank to apply to an image involves rotating the Gabor filter in a counter-clockwise direction. See “Slant Parameter” on page 1-452, and `gabor` in the Image Processing Toolbox.

QualityFactors — Scattering filter bank quality factors

[1 1] (default) | integer-valued vector

Scattering filter bank quality factors, specified as an integer-valued vector. The quality factor is the number of wavelet filters per octave. The number of wavelet filter banks in the scattering framework is equal to the number of elements in `QualityFactors`. Valid quality factors are integers less than or equal to 4. If `QualityFactors` is specified as a vector, the elements of `QualityFactors` must be nonincreasing.

The length of the vector specified in `QualityFactors` must equal the length of the vector specified in `NumRotations`.

Example: `sf = waveletScattering2('QualityFactors',[2 1])`

Precision — Precision of scattering coefficients and filters

'single' (default) | 'double'

Precision of scattering coefficients and filters:

- 'single' — Single precision
- 'double' — Double precision

Note

- All calculations involving the wavelet scattering framework are carried out in `Precision`.
 - The precision of the output of the `scatteringTransform` function does not exceed the precision of the `waveletScattering2` object.
-

OversamplingFactor – Oversampling factor

0 (default) | nonnegative integer | Inf

Oversampling factor, specified as a nonnegative integer or `Inf`. The factor specifies how much the image scattering coefficients are oversampled with respect to the critically downsampled values. The oversampling factor is on a \log_2 scale. For example, if `sf = waveletScattering2('OversamplingFactor', 1)`, the scattering transform returns 2^1 -by- 2^1 -by- P as many coefficients for each scattering path with respect to the critically sampled number. You can use `coefficientSize` to determine the number of coefficients obtained for a scattering framework. By default, `OversamplingFactor` is set to 0, which corresponds to critically downsampling the coefficients.

If you specify an oversampling factor that would result in an output image size larger than the input, the output size is truncated to the size of the input image. You can also specify the `OversamplingFactor` as `Inf`, which provides a fully undecimated scattering transform where each scattering path contains coefficient matrices equal in size to the input image.

Due to the computational complexity of the scattering transform, the recommended setting for the `OversamplingFactor` property is 0, 1, or 2. Values of 1 and 2 indicate a 2^1 -by- 2^1 -by- P and a 2^2 -by- 2^2 -by- P increase in the number of scattering coefficients per path, respectively.

Example: `sf.OversamplingFactor = 1` sets the `OversamplingFactor` property of an existing framework to 1.

OptimizePath – Optimize scattering transform logical

true (default) | false

Optimize scattering transform logical, which determines whether the scattering transform reduces the number of scattering paths to compute based on a bandwidth consideration.

When `OptimizePath` is set to `true`, a scattering path is computed only if the bandwidth of the parent node overlaps significantly with the bandwidth of the child node.

'Significant' in this context is defined as follows: for a quality factor of 1, 1/2 the 3 dB bandwidth of the child node is subtracted from the child node's wavelet center frequency. If that value is less than the 3 dB bandwidth of the parent, the scattering path is computed. For quality factors greater than 1, significant overlap is defined to be an overlap between the center frequency of the child minus the child's 3 dB bandwidth. If that overlaps with the 3 dB bandwidth of the parent, the scattering path is computed.

You can use `paths` to determine which and how many scattering paths are computed. `OptimizePath` generally results in computational savings in the second and subsequent filter banks only when the quality factors are equal in each filter bank.

Example: `sf.OptimizePath = false` sets the `OptimizePath` property of an existing framework to `false`.

Object Functions

<code>scatteringTransform</code>	Wavelet 2-D scattering transform
<code>featureMatrix</code>	Image scattering feature matrix
<code>log</code>	Natural logarithm of 2-D scattering transform
<code>filterbank</code>	Wavelet and scaling filters
<code>littlewoodPaleySum</code>	Littlewood-Paley sum
<code>coefficientSize</code>	Size of image scattering coefficients
<code>numorders</code>	Number of scattering orders
<code>numfilterbanks</code>	Number of scattering filter banks
<code>paths</code>	Scattering paths

Examples

Wavelet Image Scattering with Default Values

Create a wavelet image scattering framework with default settings. The default image size is 128-by-128, and the default invariance scale is 64.

```
sf = waveletScattering2
```

```
sf =  
    waveletScattering2 with properties:
```

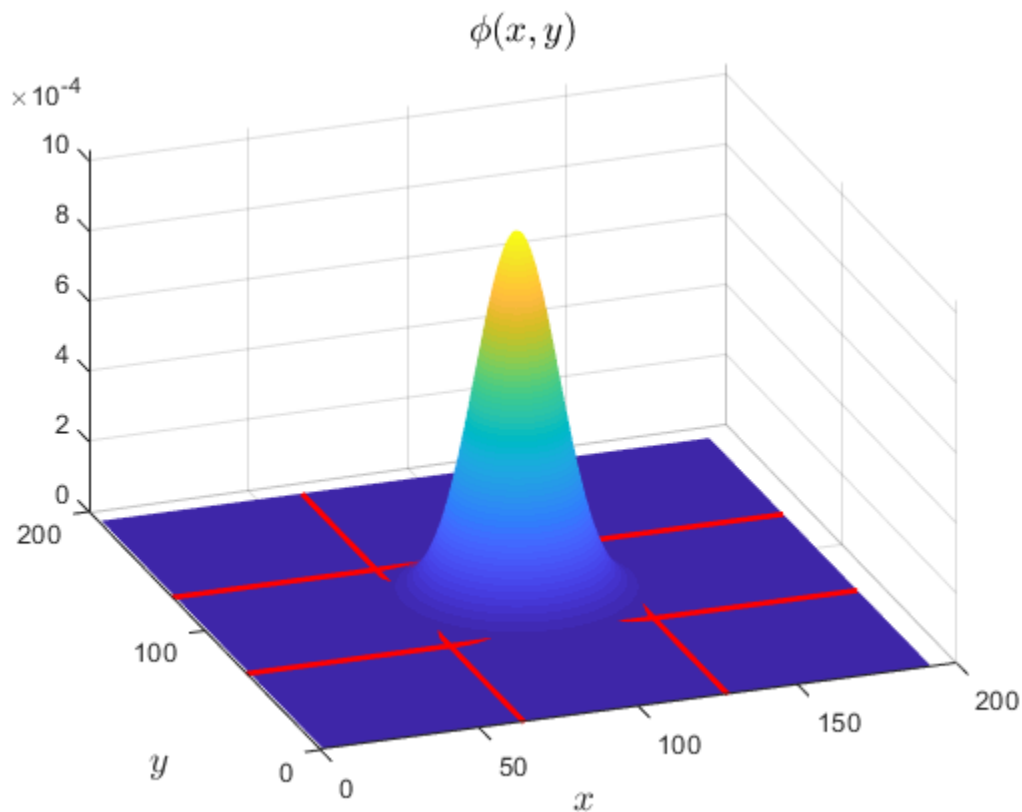
```
        ImageSize: [128 128]  
    InvarianceScale: 64  
        NumRotations: [6 6]  
        QualityFactors: [1 1]  
        Precision: 'single'  
    OversamplingFactor: 0  
        OptimizePath: 1
```

Use the `filterbank` function to obtain the Fourier transform of the scaling function, the wavelet filters, and the center spatial frequencies of the wavelet filters.

```
[phif,psif,f] = filterbank(sf);
```

The invariance scale gives the width in the x - and y -directions of the 2-D Gaussian scaling function. To confirm the scaling function has the expected spatial width, first take the inverse Fourier transform of `phif`. Use the helper function `helperPlotPhiSurface` to plot the scaling function with the extent of the invariance scale in both x and y designated. The source code for `helperPlotPhiSurface` is provided in the appendix at the end of this example.

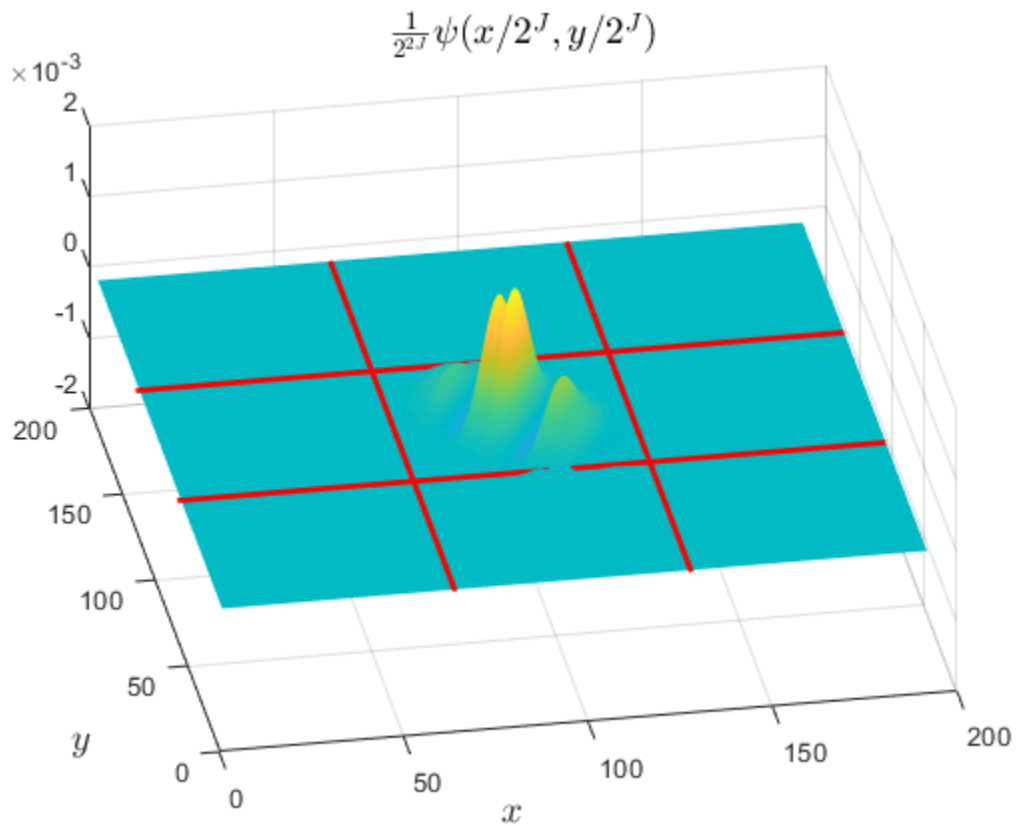
```
phi = ifftshift(ifft2(phif));  
figure  
helperPlotPhiSurface(sf,phi)
```



The scaling function is larger than 128-by-128 because it has been padded to avoid edge effects.

Extract the Fourier transform of the coarsest scale wavelet in the second filter bank and take its inverse Fourier transform. Use `helperPlotPsiSurface` to plot the real and imaginary parts of the wavelet and confirm the spatial extent of the coarsest scale wavelet does not exceed the invariance scale. Similar to the scaling function, the wavelet has been padded to avoid edge effects. The source code for `helperPlotPsiSurface` is provided in the appendix at the end of this example.

```
psiF = psif{2}(:, :, end);
psiL = ifftshift(ifft2(psiF));
figure
helperPlotPsiSurface(sf, psiL)
```



Appendix

The following helper functions are used in this example.

helperPlotPhiSurface

```
function helperPlotPhiSurface(scatFrame,data)
halfscale = scatFrame.InvarianceScale/2;
surf(data)
shading interp
view(-20,35)
Ysize = size(data,1);
Xsize = size(data,2);
Ycenter = Ysize/2;
```

```
Xcenter = Xsize/2;
hold on
plot([Xcenter-halfscale Xcenter-halfscale],[0 Ysize], 'r', 'LineWidth',2);
plot([Xcenter+halfscale Xcenter+halfscale],[0 Ysize], 'r', 'LineWidth',2);
plot([0 Xsize],[Ycenter-halfscale Ycenter-halfscale], 'r', 'LineWidth',2);
plot([0 Xsize],[Ycenter+halfscale Ycenter+halfscale], 'r', 'LineWidth',2);
title('$\phi(x,y)$', 'FontSize',14, 'Interpreter', 'Latex');
xlabel('$x$', 'FontSize',14, 'Interpreter', 'Latex')
ylabel('$y$', 'FontSize',14, 'Interpreter', 'Latex')
end
```

helperPlotPsiSurface

```
function helperPlotPsiSurface(scatFrame,data)
halfscale = scatFrame.InvarianceScale/2;
Ysize = size(data,1);
Xsize = size(data,2);
Ycenter = Ysize/2;
Xcenter = Xsize/2;
surf(real(data))
shading interp
view(-5,13)
hold on
surf(imag(data))
shading interp
plot([Xcenter-halfscale Xcenter-halfscale],[0 Ysize], 'r', 'LineWidth',2);
plot([Xcenter+halfscale Xcenter+halfscale],[0 Ysize], 'r', 'LineWidth',2);
plot([0 Xsize],[Ycenter-halfscale Ycenter-halfscale], 'r', 'LineWidth',2);
plot([0 Xsize],[Ycenter+halfscale Ycenter+halfscale], 'r', 'LineWidth',2);
title('$\frac{1}{2^J}\psi(x/2^J,y/2^J)$', 'FontSize',14, ...
      'Interpreter', 'Latex');
xlabel('$x$', 'FontSize',14, 'Interpreter', 'Latex')
ylabel('$y$', 'FontSize',14, 'Interpreter', 'Latex')
view(-10,51)
end
```

References

- [1] Bruna, A., and S. Mallat. "Invariant Scattering Convolution Networks." *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 35, Number 8, 2013, pp. 1872-1886.
- [2] Sifre, L., and S. Mallat. "Rigid-Motion Scattering for Texture Classification". arXiv preprint. 2014, pp. 1-19. <https://arxiv.org/abs/1403.1687>.

- [3] Sifre, L., and S. Mallat. "Rotation, scaling and deformation invariant scattering for texture discrimination." *2013 IEEE Conference on Computer Vision and Pattern Recognition*. 2013, pp 1233-1240.

See Also

Objects

waveletScattering

Functions

cwtft2 | dddtree2 | wavedec2

Topics

"Wavelet Scattering"

"Texture Classification with Wavelet Image Scattering"

"Digit Classification with Wavelet Scattering"

Introduced in R2019a

Wavelet Signal Denoiser

Visualize and denoise time series data

Description

The **Wavelet Signal Denoiser** app is an interactive tool for visualizing and denoising real-valued 1-D signals and comparing results. With the app, you can:


- Access all the signals in the MATLAB workspace.
- Easily adjust default parameters and apply different denoising techniques.
- Visualize and compare results.
- Export denoised signals to your workspace.
- Recreate the denoised signal in your workspace by generating a MATLAB script.

The **Wavelet Signal Denoiser** app provides a way to work with multiple versions of denoised data simultaneously.

A typical workflow for denoising a signal and comparing results using the app is:

- 1 Start the app and load a 1-D signal from the MATLAB workspace. The app provides an initial denoised version of your data using default parameters.
- 2 Adjust the denoising parameters and produce multiple versions of the denoised signal.
- 3 Compare results and export the desired denoised signal to your workspace.
- 4 To apply the same denoising parameters to other signals in your workspace, generate a MATLAB script and modify it as you see fit.

Open the Wavelet Signal Denoiser App

- MATLAB Toolstrip: On the **Apps** tab, under **Signal Processing and Communications**, click **Wavelet Signal Denoiser** .
- MATLAB command prompt: Enter `waveletSignalDenoiser`.

Examples

Denoise Signal Using Default Settings

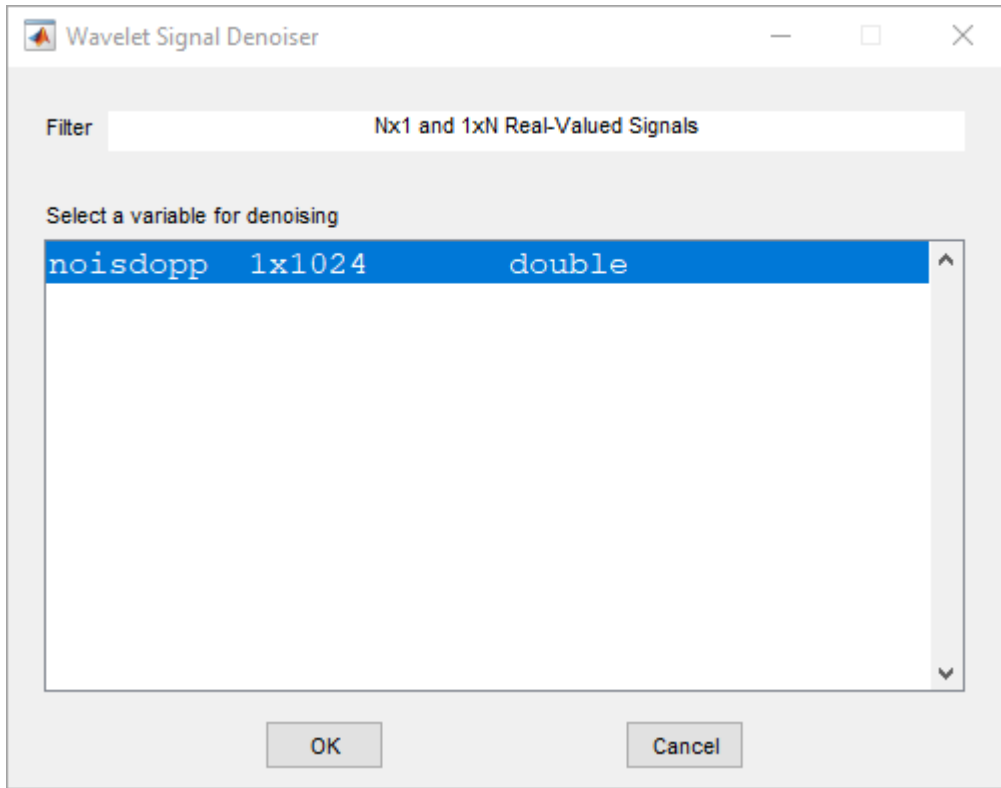
This example shows how to denoise a 1-D signal using the app default settings.

Load the noisy Doppler signal.

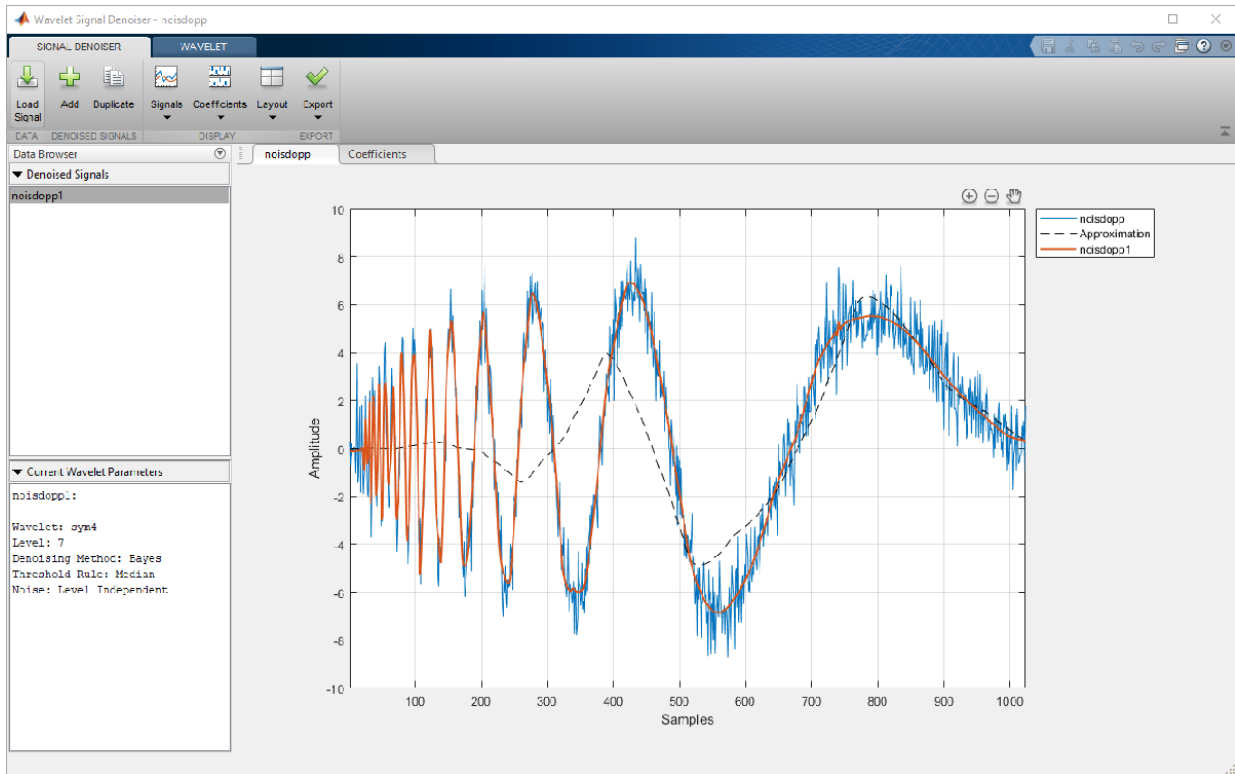
```
load noisdopp
```

Start the **Wavelet Signal Denoiser** app by choosing it from the **Apps** tab on the MATLAB® Toolstrip. You can also start the app by typing `waveletSignalDenoiser` at the MATLAB command prompt.

Load the noisy Doppler signal from the workspace into the app by clicking **Load Signal** in the toolstrip. From the list of workspace variables that can be loaded into the app, select `noisdopp` and click **OK**.



The app displays the original signal, `noisdopp`, the denoised signal, `noisdopp1`, and the coarse scale approximation, `Approximation`.



To toggle what plots are visible, you can:

- Click **Signals** ▼ in the toolstrip and use the drop-down menu to toggle the visibility of the original and approximation plots.
- Click individual signals in the plot legend.

Parameters

Wavelet — Wavelet family

sym (default) | bior | coif | db | fk

Wavelet family used to denoise the signal, specified as one of the following:

- `sym` — Symlets
- `bior` — Biorthogonal spline wavelets
- `coif` — Coiflets
- `db` — Daubechies wavelets
- `fk` — Fejér-Korovkin wavelets

Method — Denoising method

`Bayes (default)` | `BlockJS` | `FDR` | `Minimax` | `SURE` | `UniversalThreshold`

Denoising method to apply, specified as one of the following:

- `Bayes` — Empirical Bayes
- `BlockJS` — Block James-Stein
- `FDR` — False Discovery Rate
- `Minimax` — Minimax Estimation
- `SURE` — Stein's Unbiased Risk Estimate
- `UniversalThreshold` — Universal Threshold $\sqrt{2\ln(\text{length}(x))}$.

Rule — Thresholding rule

`Median (default)` | `Mean` | `Soft` | `Hard`

Thresholding rule to use. The possible rules you can specify for different denoising methods are as follows:

- Block James-Stein — James-Stein
- Empirical Bayes — Median, Mean, Soft, Hard
- False Discovery Rate — Hard
- Minimax Estimation — Soft, Hard
- Stein's Unbiased Risk Estimate — Soft, Hard
- Universal Threshold — Soft, Hard

Tips

To denoise more than one signal simultaneously, you can run multiple instances of the **Wavelet Signal Denoiser** app.

See Also

Functions

wdenoise

Topics

“Denoise a Signal with the Wavelet Signal Denoiser”

Introduced in R2017b

waveletsupport

CWT filter bank time supports

Syntax

```
spsi = waveletsupport(fb)  
spsi = waveletsupport(fb,thresh)
```

Description

`spsi = waveletsupport(fb)` returns the wavelet time supports defined as the time interval in which all of the wavelet's energy occurs. The default tolerance is 99.99% of the energy. The time supports are returned in the MATLAB table `spsi`. The wavelets are normalized to have unit energy.

`spsi = waveletsupport(fb,thresh)` specifies the threshold for the integrated energy. The time support of the wavelet is defined as the first instant the integrated energy exceeds `thresh` and the last instant the integrated energy is less than `1-thresh`. If unspecified, `thresh` defaults to $1e-4$.

Examples

Wavelet Filter Bank Time-Supports

Create a continuous wavelet transform filter bank. Set the sampling frequency to 1000 Hz and the frequency limits to range from 100 Hz to 200 Hz. Obtain the time-supports of the wavelets in the filter bank.

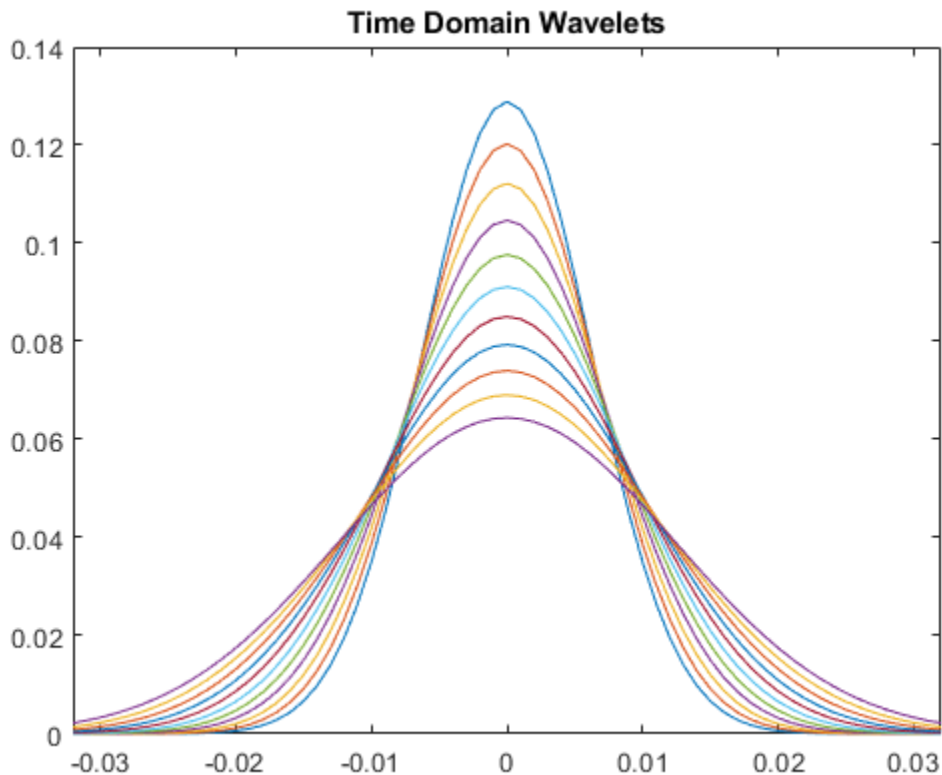
```
fb = cwtfilterbank('SamplingFrequency',1000,'FrequencyLimits',[100 200]);  
spsi = waveletsupport(fb)
```

```
spsi=11x5 table  
      CF      IsAnalytic      TimeSupport      Begin      End  
      _____      _____      _____      _____      _____
```

200	"Analytic"	0.032	-0.016	0.016
186.61	"Analytic"	0.034	-0.017	0.017
174.11	"Analytic"	0.038	-0.019	0.019
162.45	"Analytic"	0.04	-0.02	0.02
151.57	"Analytic"	0.042	-0.021	0.021
141.42	"Analytic"	0.046	-0.023	0.023
131.95	"Analytic"	0.048	-0.024	0.024
123.11	"Analytic"	0.052	-0.026	0.026
114.87	"Analytic"	0.056	-0.028	0.028
107.18	"Analytic"	0.06	-0.03	0.03
100	"Analytic"	0.064	-0.032	0.032

Obtain the time domain wavelets from the filter bank and plot their magnitudes. Use the table to set the x-limits of the plot equal to the wavelet with largest time support.

```
[psi,t] = wavelets(fb);  
plot(t,abs(psi))  
xlim([spsi.Begin(end) spsi.End(end)])  
title('Time Domain Wavelets')
```



Input Arguments

fb — Continuous wavelet transform filter bank

object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

thresh — Time-support threshold

$1e-4$ (default) | positive real number

Time-support threshold for the wavelet, specified as a positive real number between 0 and 0.05. The time support of the wavelet is defined as the first instant the integrated

energy of the wavelet exceeds `thresh` and the last instant the integrated energy is less than `1 - thresh`.

Data Types: `double`

Output Arguments

spsi — Wavelet time supports

`table`

Wavelet time supports, returned as a `Ns`-by-5 MATLAB table, where `Ns` is the number of wavelet bandpass frequencies (equal to the number of scales). The table has the following five variables:

CF — Wavelet center frequency

positive real number

Wavelet center frequency, returned as a positive real number.

Data Types: `double`

IsAnalytic — Wavelet designation

"Analytic" | "Nonanalytic"

Wavelet designation, returned as a string. Wavelets that do not decay to 5% of their peak value at the Nyquist frequency are not considered analytic. The time support information for those wavelets are returned as NaNs.

Data Types: `string`

TimeSupport — Wavelet time support

positive integer | NaN

Wavelet time support, returned in samples, seconds, or MATLAB durations. The units of `TimeSupport` depend on whether you specify `SamplingFrequency` or `SamplingPeriod`. If you specify a `SamplingFrequency`, the units are in seconds. If you specify a `SamplingPeriod`, the units are the same as the `SamplingPeriod`. If no `SamplingFrequency` or `SamplingPeriod` is specified, the units are in samples.

Data Types: `double`

Begin — Beginning of the wavelet time support

integer

Beginning of the wavelet time support, returned as an integer. `Begin` is defined as the first instant the wavelet integrated energy exceeds the default threshold, $1e-4$. `Begin` has the same units as `TimeSupport`.

Data Types: `double`

End — End of the wavelet time support

`integer`

End of the wavelet time support, returned as an integer. `End` is defined as the last instant the wavelet integrated energy is less than $1-1e-4$. `End` has the same units as `TimeSupport`.

Data Types: `double`

Data Types: `table`

See Also

`cwtfilterbank` | `wavelets`

Introduced in R2018a

waveletsupport

DWT filter bank time supports

Syntax

```
spsi = waveletsupport(fb)
spsi = waveletsupport(fb,thresh)
[spsi,sphi] = waveletsupport(fb)
[spsi,sphi,tlow,thigh] = waveletsupport(fb)
```

Description

`spsi = waveletsupport(fb)` returns the wavelet time supports defined as the time interval in which all the wavelet energy occurs ($> 99.99\%$ of the energy for the default threshold). `spsi` is a real-valued L -by-1 vector where L is the number of levels in the DWT filter bank `fb`.

`spsi = waveletsupport(fb,thresh)` specifies the threshold for the integrated energy. `thresh` is a positive real number in the interval $0 < \text{thresh} \leq 0.05$. If unspecified, `thresh` defaults to 10^{-6} . The percent energy contained in the time support is $(1-2*\text{thresh})*100$. The time support of the wavelet is defined as the first instant the integrated energy exceeds `thresh` and the last instant it is less than $1-\text{thresh}$. The wavelets are normalized to have unit energy for the computation.

`[spsi,sphi] = waveletsupport(fb)` returns the scaling function time supports at all levels. `sphi` is a real-valued L -by-1 vector where L is the number of levels in the DWT filter bank `fb`.

`[spsi,sphi,tlow,thigh] = waveletsupport(fb)` returns the instants the integrated energy in the wavelets and scaling functions exceed `thresh` in `tlow` and the last instant the integrated energy is less than $1-\text{thresh}$ in `thigh`. `tlow` and `thigh` are L -by-2 matrices where L is the number of levels in the DWT filter bank `fb`. The first columns of `tlow` and `thigh` contain the times for the wavelets. The second columns of `tlow` and `thigh` contain the times for the scaling functions. The difference of the corresponding columns in `thigh` and `tlow` plus one sampling period equals the values in `spsi` and `sphi` respectively.

Examples

DWT Filter Bank Wavelet Time Supports

Find the time supports for a Haar wavelet filter bank.

```
fb = dwtfilterbank('Wavelet', 'haar', 'Level', 8);  
Spsi = waveletsupport(fb)
```

```
Spsi = 8×1
```

```
     2  
     4  
     8  
    16  
    32  
    64  
   128  
   256
```

Input Arguments

fb — Discrete wavelet transform filter bank

object

Discrete wavelet transform (DWT) filter bank, specified as a `dwtfilterbank` object.

thresh — Threshold for the integrated energy

10^{-6} (default) | positive scalar

Threshold for the integrated energy, specified as a positive scalar in the interval $0 < \text{thresh} \leq 0.05$.

Output Arguments

spsi — Wavelet time supports

real-valued column vector

Wavelet time supports, returned as a real-valued column vector of length L , where L is the number of levels in the DWT filter bank.

sphi — Scaling function time supports

real-valued column vector

Scaling function time supports, returned as a real-valued column vector of length L , where L is the number of levels in the DWT filter bank.

tlow — First instants

real-valued matrix

First instants the integrated energy in the wavelet and scaling functions exceed `thresh`, returned as real-valued L -by-2 matrix where L is the number of levels in the DWT filter bank. The first column of `tlow` contains the times for the wavelets. The second column of `tlow` contains the times for the scaling functions.

thigh — Last instants

real-valued matrix

Last instants the integrated energy in the wavelet and scaling functions is less than `1-thresh`, returned as real-valued L -by-2 matrix where L is the number of levels in the DWT filter bank. The first column of `thigh` contains the times for the wavelets. The second column of `thigh` contains the times for the scaling functions.

See Also

`dwtfilterbank` | `scalingfunctions` | `wavelets`

Introduced in R2018a

wavemngr

Wavelet manager

Syntax

```
wavemngr('add', FN, FSN, WT, NUMS, FILE)
wavemngr('add', FN, FSN, WT, NUMS, FILE, B)
wavemngr('add', FN, FSN, WT, {NUMS, TYPNUMS}, FILE)
wavemngr('add', FN, FSN, WT, {NUMS, TYPNUMS}, FILE, B)
```

Description

wavemngr is a type of wavelets manager. It allows you to add, delete, restore, or read wavelets.

wavemngr('add', FN, FSN, WT, NUMS, FILE) or wavemngr('add', FN, FSN, WT, NUMS, FILE, B) or wavemngr('add', FN, FSN, WT, {NUMS, TYPNUMS}, FILE) or wavemngr('add', FN, FSN, WT, {NUMS, TYPNUMS}, FILE, B), add a new wavelet family to the toolbox.

FN = Family Name (character vector or string scalar)

FSN = Family Short Name (character vector or string scalar of length equal or less than four characters)

WT defines the wavelet type:

- WT = 1, for orthogonal wavelets
- WT = 2, for biorthogonal wavelets
- WT = 3, for wavelet with scaling function
- WT = 4, for wavelet without scaling function
- WT = 5, for complex wavelet without scaling function

If the family contains a single wavelet, NUMS = ''. Note that for this case you specify an empty character vector or string scalar.

Examples:

mexh	' '
morl	' '

If the wavelet is member of a finite family of wavelets, NUMS is a character vector or string scalar containing a space-separated list of items representing wavelet parameters.

Example:

bior	NUMS = '1.1 1.3 ... 4.4 5.5 6.8'
------	----------------------------------

If the wavelet is part of an infinite family of wavelets, NUMS is a character vector or string scalar containing a space-separated list of items representing wavelet parameters, terminated by the special sequence **.

Examples:

db	NUMS = '1 2 3 4 5 6 7 8 9 10 **'
shan	NUMS = '1-1.5 1-1 1-0.5 1-0.1 2-3 **'

In these last two cases, TYPNUMS specifies the wavelet parameter input format: 'integer' or 'real' or 'charactervector'; the default value is 'integer'.

Examples:

db	TYPNUMS = 'integer'
bior	TYPNUMS = 'real'
shan	TYPNUMS = 'charactervector'

FILE = MAT-file or code file name (character vector or string scalar). See usage in the "Examples" section.

B = [lb ub] specifies lower and upper bounds of effective support for wavelets of type = 3, 4, or 5.

wavemngr('del',N), deletes a wavelet or a wavelet family. N is the Family Short Name or the Wavelet Name (in the family). N is a character vector or string scalar.

wavemngr('restore') or wavemngr('restore',IN2) restores previous or initial wavelets. If nargin = 1, the previous wavelets.asc ASCII-file is restored; otherwise the initial wavelets.asc ASCII-file is restored. Here IN2 is a dummy argument.

OUT1 = wavemngr('read') returns all wavelet family names.

OUT1 = wavemngr('read',IN2) returns all wavelet names, IN2 is a dummy argument.

OUT1 = wavemngr('read_asc') reads wavelets.asc ASCII-file and returns all wavelets information.

Examples

Wavelet Names and Family Names

List the wavelet families available by default.

```
wavemngr('read')
```

```
ans = 18x35 char array
'===== '
'Haar          ->->haar      '
'Daubechies    ->->db        '
'Symlets       ->->sym       '
'Coiflets      ->->coif      '
'BiorSplines   ->->bior      '
'ReverseBior   ->->rbio      '
'Meyer         ->->meyr      '
'DMeyer        ->->dmey      '
'Gaussian      ->->gaus      '
'Mexican_hat   ->->mexh      '
'Morlet        ->->morl      '
'Complex Gaussian ->->cgau    '
'Shannon       ->->shan      '
'Frequency B-Spline->->fbsp    '
'Complex Morlet ->->cmor      '
'Fejer-Korovkin ->->fk       '
'===== '
```

List all wavelets.


```
wavemngr('read',1)
```

```
ans = 71x44 char array
```

```

=====
'Haar          ->->haar
=====
'Daubechies   ->->db
-----
'db1->db2->db3->db4->
'db5->db6->db7->db8->
'db9->db10->db**->
=====
'Symlets      ->->sym
-----
'sym2->sym3->sym4->sym5->
'sym6->sym7->sym8->sym**->
=====
'Coiflets     ->->coif
-----
'coif1->coif2->coif3->coif4->
'coif5->
=====
'BiorSplines  ->->bior
-----
'bior1.1->bior1.3->bior1.5->bior2.2->
'bior2.4->bior2.6->bior2.8->bior3.1->
'bior3.3->bior3.5->bior3.7->bior3.9->
'bior4.4->bior5.5->bior6.8->
=====
'ReverseBior  ->->rbio
-----
'rbio1.1->rbio1.3->rbio1.5->rbio2.2->
'rbio2.4->rbio2.6->rbio2.8->rbio3.1->
'rbio3.3->rbio3.5->rbio3.7->rbio3.9->
'rbio4.4->rbio5.5->rbio6.8->
=====
'Meyer        ->->meyr
=====
'DMeyer       ->->dmey
=====
'Gaussian     ->->gaus
-----
'gaus1->gaus2->gaus3->gaus4->
'gaus5->gaus6->gaus7->gaus8->
=====

```

```
'Mexican_hat      ->->mexh
'=====
'Morlet           ->->morl
'=====
'Complex Gaussian ->->cgau
'-----
'cgau1->cgau2->cgau3->cgau4->
'cgau5->cgau6->cgau7->cgau8->
'=====
'Shannon          ->->shan
'-----
'shan1-1.5->shan1-1->shan1-0.5->shan1-0.1->
'shan2-3->shan**->
'=====
'Frequency B-Spline->->fbsp
'-----
'fbsp1-1-1.5->fbsp1-1-1->fbsp1-1-0.5->fbsp2-1-1->
'fbsp2-1-0.5->fbsp2-1-0.1->fbsp**->
'=====
'Complex Morlet   ->->cmor
'-----
'cmor1-1.5->cmor1-1->cmor1-0.5->cmor1-1->
'cmor1-0.5->cmor1-0.1->cmor**->
'=====
'Fejer-Korovkin  ->->fk
'-----
'fk4->fk6->fk8->fk14->
'fk18->fk22->
'=====
```

Add Wavelet Families

In the following example, new compactly supported orthogonal wavelets are added to the toolbox. These wavelets, which are a slight generalization of the Daubechies wavelets, are based on the use of Bernstein polynomials and are due to Kateb and Lemarié.

Add new family of orthogonal wavelets. You must define:

- Family Name: Lemarie
- Family Short Name: lem

- Type of wavelet: 1 (orth)
- Wavelets numbers: 1 2 3 4 5
- File driver: lemwavf

The function `lemwavf.m` must be as follows:

```
function w = lemwavf(wname)
```

where the input argument `wname` is a character vector or string scalar of the form `sh.name + number` (for example, `'lem1'` or `'lem2'`) and `w` the corresponding scaling filter. The addition is obtained using

```
wavemngr('add','Lemarie','lem',1,'1 2 3 4 5','lemwavf');
```

The ASCII file `'wavelets.asc'` is saved as `'wavelets.prv'`, then it is modified and the MAT-file `'wavelets.inf'` is generated.

Note that `wavemngr` works on the current folder. If you add a new wavelet family, it is available in this folder only.

List the available wavelet families.

```
wavemngr('read')
```

```
% ans =
=====
Haar                haar
Daubechies          db
Symlets             sym
Coiflets            coif
BiorSplines         bior
ReverseBior         rbio
Meyer               meyr
DMeyer              dmey
Gaussian            gaus
Mexican_hat         mexh
Morlet              morl
Complex Gaussian    cgau
Shannon             shan
Frequency B-Spline fbsp
Complex Morlet      cmor
Lemarie             lem
=====
```

Remove the added family. Regenerate the list of wavelet families.

```
wavemngr('del','Lemarie');  
wavemngr('read')
```

```
ans =  
=====
```

Haar	haar
Daubechies	db
Symlets	sym
Coiflets	coif
BiorSplines	bior
ReverseBior	rbior
Meyer	meyr
DMeyer	dmey
Gaussian	gaus
Mexican_hat	mexh
Morlet	morl
Complex Gaussian	cgau
Shannon	shan
Frequency B-Spline	fbsp
Complex Morlet	cmor

```
=====
```

Restore the previous ASCII file 'wavelets.prv', then build the MAT-file 'wavelets.inf'. List the restored wavelets.

```
wavemngr('restore');  
wavemngr('read',1)
```

```
ans =  
=====
```

Haar			haar
------	--	--	------

```
=====
```

Daubechies			db
------------	--	--	----

```
-----
```

db1	db2	db3	db4
db5	db6	db7	db8
db9	db10	db**	

```
=====
```

Symlets			sym
---------	--	--	-----

```
-----
```

sym2	sym3	sym4	sym5
sym6	sym7	sym8	sym**

```
=====
```

```

Coiflets                                coif
-----
coif1      coif2      coif3      coif4
coif5
=====
BiorSplines                                bior
-----
bior1.1    bior1.3    bior1.5    bior2.2
bior2.4    bior2.6    bior2.8    bior3.1
bior3.3    bior3.5    bior3.7    bior3.9
bior4.4    bior5.5    bior6.8
=====
ReverseBior                                rbio
-----
rbio1.1    rbio1.3    rbio1.5    rbio2.2
rbio2.4    rbio2.6    rbio2.8    rbio3.1
rbio3.3    rbio3.5    rbio3.7    rbio3.9
rbio4.4    rbio5.5    rbio6.8
=====
Meyer                                meyr
=====
DMeyer                                dmey
=====
Gaussian                                gaus
-----
gaus1      gaus2      gaus3      gaus4
gaus5      gaus6      gaus7      gaus8
gaus**
=====
Mexican_hat                                mexh
=====
Morlet                                morl
=====
Complex Gaussian                                cgau
-----
cgau1      cgau2      cgau3      cgau4
cgau5      cgau**
=====
Shannon                                shan
-----
shan1-1.5  shan1-1    shan1-0.5  shan1-0.1
shan2-3    shan**
=====
Frequency B-Spline                                fbsp

```

```

-----
fbstp1-1-1.5      fbstp1-1-1      fbstp1-1-0.5      fbstp2-1-1
fbstp2-1-0.5      fbstp2-1-0.1      fbstp**
=====
Complex Morlet          cmor
-----
cmor1-1.5      cmor1-1      cmor1-0.5      cmor1-1
cmor1-0.5      cmor1-0.1      cmor**
=====
Lemarie      lem
-----
lem1      lem2      lem3      lem4      lem5
=====

```

Restore the initial wavelets. Restore the initial ASCII file 'wavelets.ini' and the initial MAT-file 'wavelets.bin'. Regenerate the list of wavelet families.

```
wavemngr('restore',0);
wavemngr('read')
```

```
ans =
=====
Haar          haar
Daubechies    db
Symlets       sym
Coiflets      coif
BiorSplines   bior
ReverseBior   rbio
Meyer         meyr
DMeyer        dmey
Gaussian      gaus
Mexican_hat   mexh
Morlet        morl
Complex Gaussian cgau
Shannon       shan
Frequency B-Spline fbsp
Complex Morlet cmor
=====

```

All command line capabilities are available for new families of wavelets. Create a new family. Compute the four associated filters and the scale and wavelet functions.

```
wavemngr('add','Lemarie','lem',1,'1 2 3','lemwavf');
[Lo_D,Hi_D,Lo_R,Hi_R] = wfilters('lem3');
[phi,psi,xval] = wavefun('lem3');
```

Add a new family of orthogonal wavelets in a form specialized for the GUI mode. The file `lemwavf` allows you to compute the filter for any order. If you want to get a popup of the form '1 2 3 **' associated with the family, then wavelets are appended for GUI mode using:

```
wavemngr('restore',0);
wavemngr('add','Lemarie','lem',1,'1 2 3 **','lemwavf');
```

After this sequence, all GUI capabilities are available for the new wavelets. Note that the last command allows a shortcut in the order definition only if the possible orders are integers.

Adding Biorthogonal Wavelet Filters

This example shows how to take analysis and synthesis filters associated with a biorthogonal wavelet and make them compatible with the Wavelet Toolbox™. The Wavelet Toolbox requires that analysis and synthesis lowpass and highpass filters have equal even length. This example uses the nearly-orthogonal biorthogonal wavelets based on the Laplacian pyramid scheme of Burt and Adelson (Table 8.4 on page 283 in [1]). The example also demonstrates how to examine properties of the biorthogonal wavelets.

Define the analysis and synthesis filter coefficients of the biorthogonal wavelet.

```
Hd = [-1 5 12 5 -1]/20*sqrt(2);
Gd = [3 -15 -73 170 -73 -15 3]/280*sqrt(2);
Hr = [-3 -15 73 170 73 -15 -3]/280*sqrt(2);
Gr = [-1 -5 12 -5 -1]/20*sqrt(2);
```

`Hd` and `Gd` are the lowpass and highpass analysis filters, respectively. `Hr` and `Gr` are the lowpass and highpass synthesis filters. They are all finite impulse response (FIR) filters. Confirm the lowpass filter coefficients sum to $\sqrt{2}$ and the highpass filter coefficients sum to 0.

```
sum(Hd)/sqrt(2)
```

```
ans = 1.0000
```

```
sum(Hr)/sqrt(2)
```

```
ans = 1.0000
```

```
sum(Gd)
```

```
ans = -1.0061e-16
```

```
sum(Gr)
```

```
ans = -9.7145e-17
```

The z -transform of an FIR filter h is a Laurent polynomial $h(z)$ given by $h(z) = \sum_{k=k_b}^{k_e} h_k z^{-k}$.

The degree $|h|$ of a Laurent polynomial is defined as $|h| = k_e - k_b$. Therefore the length of the filter h is $1 + |h|$. Examine the Laurent expansion of the scaling and wavelet filters.

```
PHd = laurpoly(Hd, 'dmin', -2)
```

```
PHd(z) = - 0.07071*z^(+2) + 0.3536*z^(+1) + 0.8485 + 0.3536*z^(-1) - 0.07071*z^(-2)
```

```
PHr = laurpoly(Hr, 'dmin', -3)
```

```
PHr(z) = ...
- 0.01515*z^(+3) - 0.07576*z^(+2) + 0.3687*z^(+1) + 0.8586 + 0.3687*z^(-1) ...
- 0.07576*z^(-2) - 0.01515*z^(-3)
```

```
PGd = laurpoly(Gd, 'dmin', -3)
```

```
PGd(z) = ...
+ 0.01515*z^(+3) - 0.07576*z^(+2) - 0.3687*z^(+1) + 0.8586 - 0.3687*z^(-1) ...
- 0.07576*z^(-2) + 0.01515*z^(-3)
```

```
PGr = laurpoly(Gr, 'dmin', -2)
```

```
PGr(z) = - 0.07071*z^(+2) - 0.3536*z^(+1) + 0.8485 - 0.3536*z^(-1) - 0.07071*z^(-2)
```

Since the filters are associated with biorthogonal wavelet, confirm $PHd(z) PHr(z) + PG(z) PGr(z) = 2$.

```
PHd*PHr + PGd*PGr
```

```
ans(z) = 2
```


The Wavelet Toolbox requires that filters associated with the wavelet have even equal length. To use the Laplacian wavelet filters in the Toolbox, you must include the missing powers of the Laurent series as zeros.

The degrees of PHd and PHr are 4 and 6, respectively. The minimum even length filter that can accommodate the four filters has length 8, which corresponds to a Laurent polynomial of degree 7. The strategy is to prepend and append 0s as evenly as possible so that all filters are of length 8. Therefore prepend 0 to all the filters. Then append two 0s to Hd and Gr.

```
Hd = [0 Hd 0 0];
Gd = [0 Gd];
Hr = [0 Hr];
Gr = [0 Gr 0 0];
```

You can examine properties of the biorthogonal wavelets by creating DWT filter banks. Create two custom DWT filter banks using the filters, one for analysis, and the other for synthesis. Confirm the filter banks are biorthogonal.

```
fb = dwtfilterbank('Wavelet','Custom',...
    'CustomScalingFilter',[Hd' Hr'],...
    'CustomWaveletFilter',[Gd' Gr']);

fb2 = dwtfilterbank('Wavelet','Custom',...
    'CustomScalingFilter',[Hd' Hr'],...
    'CustomWaveletFilter',[Gd' Gr'],...
    'FilterType','Synthesis');

fprintf('fb: isOrthogonal = %d\tisBiorthogonal = %d\n',...
    isOrthogonal(fb),isBiorthogonal(fb));

fb: isOrthogonal = 0    isBiorthogonal = 1

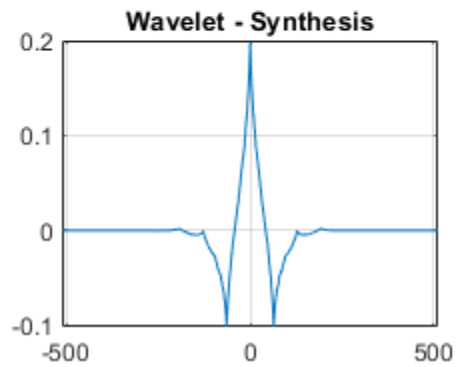
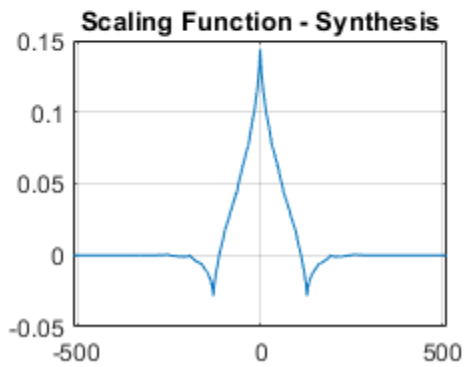
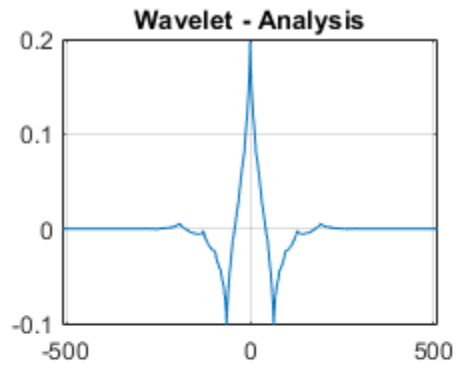
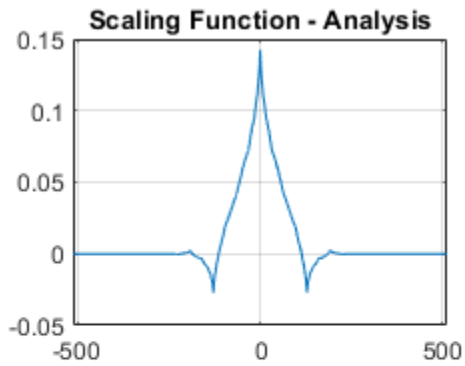
fprintf('fb2: isOrthogonal = %d\tisBiorthogonal = %d\n',...
    isOrthogonal(fb2),isBiorthogonal(fb2));

fb2: isOrthogonal = 0    isBiorthogonal = 1

Plot the scaling and wavelet functions associated with the filter banks at the coarsest
scale.

[phi,t] = scalingfunctions(fb);
[psi,~] = wavelets(fb);
```

```
[phi2,~] = scalingfunctions(fb2);  
[psi2,~] = wavelets(fb2);  
subplot(2,2,1)  
plot(t,phi(end,:))  
grid on  
title('Scaling Function - Analysis')  
subplot(2,2,2)  
plot(t,psi(end,:))  
grid on  
title('Wavelet - Analysis')  
subplot(2,2,3)  
plot(t,phi2(end,:))  
grid on  
title('Scaling Function - Synthesis')  
subplot(2,2,4)  
plot(t,psi2(end,:))  
grid on  
title('Wavelet - Synthesis')
```



Compute the filter bank framebounds.

```
[analysisLowerBound,analysisUpperBound] = framebounds(fb)
```

```
analysisLowerBound = 0.9505
```

```
analysisUpperBound = 1.0211
```

```
[synthesisLowerBound,synthesisUpperBound] = framebounds(fb2)
```

```
synthesisLowerBound = 0.9800
```

```
synthesisUpperBound = 1.0528
```

Limitations

wavemngr allows you to add a new wavelet. You must verify that it is truly a wavelet. No check is performed either about this point or about the type of the new wavelet. You can use `dwtfilterbank` to verify if a wavelet is orthogonal or biorthogonal.

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*. CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: Society for Industrial and Applied Mathematics, 1992.

See Also

Topics

“Adding Quadrature Mirror and Biorthogonal Wavelet Filters”

Introduced before R2006a

wavenames

Wavelet names for LWT

Syntax

`W = wavenames(T)`

Description

`W = wavenames(T)` returns a cell array that contains the name of all wavelets of type *T*. The valid values for *T* are

- 'all' — all wavelets
- 'lazy' — “lazy” wavelet
- 'orth' — orthogonal wavelets
- 'bior' — biorthogonal wavelets

`W = wavenames` is equivalent to `W = wavenames('all')`.

Introduced before R2006a

waverec

1-D wavelet reconstruction

Syntax

```
X = waverec(C,L,Lo_R,Hi_R)
X = waverec(C,L,'wname')
X = appcoef(C,L,'wname',0)
```

Description

`waverec` performs a multilevel one-dimensional wavelet reconstruction using either a specific wavelet (`'wname'`, see `wfilters`) or specific reconstruction filters (`Lo_R` and `Hi_R`).

Note `waverec` supports only Type 1 (orthogonal) or Type 2 (biorthogonal) wavelets.

`X = waverec(C,L,'wname')` reconstructs the signal `X` based on the multilevel wavelet decomposition structure `[C,L]` and wavelet `'wname'`. (For information about the decomposition structure, see `wavedec`.)

`X = waverec(C,L,Lo_R,Hi_R)` reconstructs the signal `X` as above, using the reconstruction filters you specify. `Lo_R` is the reconstruction low-pass filter and `Hi_R` is the reconstruction high-pass filter.

Note that `X = waverec(C,L,'wname')` is equivalent to `X = appcoef(C,L,'wname',0)`.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
% Load original one-dimensional signal.
```

```
load leleccum; s = leleccum(1:3920); ls = length(s);

% Perform decomposition of signal at level 3 using db5.
[c,l] = wavedec(s,3,'db5');

% Reconstruct s from the wavelet decomposition structure [c,l].
a0 = waverec(c,l,'db5');

% Check for perfect reconstruction.
err = norm(s-a0)
err =
    3.2079e-09
```

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

appcoef | idwt | wavedec

Introduced before R2006a

waverec2

2-D wavelet reconstruction

Syntax

```
X = waverec2(C,S,wname)
X = waverec2(C,S,Lo_R,Hi_R)
X = waverec2(C,S,wname)
X = appcoef2(C,S,wname,0)
```

Description

`X = waverec2(C,S,wname)` performs a multilevel wavelet reconstruction of the matrix `X` based on the wavelet decomposition structure `[C,S]`. For detailed storage information, see `wavedec2`. `wname` is a character vector or string scalar specifying the wavelet. See `wfilters` for more information.

Instead of specifying the wavelet name, you can specify the filters.

- `X = waverec2(C,S,Lo_R,Hi_R)`, `Lo_R` is the reconstruction low-pass filter
- `Hi_R` is the reconstruction high-pass filter.

`waverec2` is the inverse function of `wavedec2` in the sense that the abstract statement `waverec2(wavedec2(X,N,wname),wname)` returns `X`.

`X = waverec2(C,S,wname)` is equivalent to `X = appcoef2(C,S,wname,0)`.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
% Load original image.
load woman;
% X contains the loaded image.
% Perform decomposition at level 2
% of X using sym4.
```



```
[c,s] = wavedec2(X,2,'sym4');  
% Reconstruct X from the wavelet  
% decomposition structure [c,s].  
a0 = waverec2(c,s,'sym4');  
% Check for perfect reconstruction.  
max(max(abs(X-a0)))  
ans =  
    2.5565e-10
```

Tips

If *C* and *S* are obtained from an indexed image analysis or a truecolor image analysis, *X* is an *m*-by-*n* matrix or an *m*-by-*n*-by-3 array, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

`appcoef2` | `idwt2` | `wavedec2`

Introduced before R2006a

waverec3

3-D wavelet reconstruction

Syntax

```
X = waverec3(WDEC)
C = waverec3(WDEC,TYPE,N)
X = waverec3(WDEC,'a',0)
X = waverec3(WDEC,'ca',0)
C = waverec3(WDEC,TYPE)
C = waverec3(WDEC,TYPE,N)
```

Description

`waverec3` performs a multilevel 3-D wavelet reconstruction starting from a multilevel 3-D wavelet decomposition.

`X = waverec3(WDEC)` reconstructs the 3-D array `X` based on the multilevel wavelet decomposition structure `WDEC`. You can also use `waverec3` to extract coefficients from a 3-D wavelet decomposition.

`WDEC` is a structure with the fields shown in the table.

`C = waverec3(WDEC,TYPE,N)` reconstructs the multilevel components at level `N` of a 3-D wavelet decomposition. `N` must be a positive integer less than or equal to the level of the decomposition.

Valid values for `TYPE` are:

- A group of three characters `'xyz'`, one per direction, with `'x'`, `'y'` and `'z'` selected in the set `{'a','d','l','h'}` or in the corresponding uppercase set `{'A','D','L','H'}`, where `'A'` (or `'L'`) is a low-pass filter and `'D'` (or `'H'`) is a high-pass filter.
- The char `'d'` (or `'h'` or `'D'` or `'H'`) gives the sum of all the components different from the low-pass.
- The char `'a'` (or `'l'` or `'A'` or `'L'`) gives the low-pass component (the approximation at level `N`).

For extraction, the valid values for TYPE are the same but prefixed by 'c' or 'C'.

$X = \text{waverec3}(\text{WDEC}, 'a', 0)$ or $X = \text{waverec3}(\text{WDEC}, 'ca', 0)$ is equivalent to $X = \text{waverec3}(\text{WDEC})$. X is a reconstruction of the coefficients in WDEC at level 0.

$C = \text{waverec3}(\text{WDEC}, \text{TYPE})$ is equivalent to $C = \text{waverec3}(\text{WDEC}, \text{TYPE}, N)$ with N equal to the level of the decomposition.

sizeINI	Size of the three-dimensional array X
level	Level of the decomposition
mode	Name of the wavelet transform extension mode
filters	Structure with 4 fields, LoD, HiD, LoR, and HiR, which contain the filters used for DWT

<p>dec</p>	<p>$N \times 1$ cell array containing the coefficients of the decomposition. N is equal to $7 * WDEC.level + 1$.</p> <p>dec{1} contains the lowpass component (approximation) at the level of the decomposition. The approximation is equivalent to the filtering operations 'LLL'.</p> <p>dec{k+2}, ..., dec{k+8} with $k = 0, 7, 14, \dots, 7 * (WDEC.level - 1)$ contain the 3-D wavelet coefficients for the multiresolution starting with the coarsest level when $k=0$.</p> <p>For example, if $WDEC.level=3$, dec{2}, ..., dec{8} contain the wavelet coefficients for level 3 ($k=0$), dec{9}, ..., dec{15} contain the wavelet coefficients for level 2 ($k=7$), and dec{16}, ..., dec{22} contain the wavelet coefficients for level 1 ($k=7 * (WDEC.level - 1)$).</p> <p>At each level, the wavelet coefficients in dec{k+2}, ..., dec{k+8} are in the following order: 'HLL', 'LHL', 'HHL', 'LLH', 'HLH', 'LHH', 'HHH'.</p> <p>The sequence of letters gives the order in which the separable filtering operations are applied from left to right. For example, 'LHH' means that the lowpass (scaling) filter with downsampling is applied to the rows of X, followed by the highpass (wavelet) filter with downsampling applied to the columns of X. Finally, the highpass filter with downsampling is applied to the 3rd dimension of X.</p>
<p>sizes</p>	<p>Successive sizes of the decomposition components</p>

Examples

Perfect Reconstruction with 3-D Discrete Wavelet Transform

Construct a 3-D matrix, obtain the wavelet transform down to level 2 using the 'db2' wavelet, and reconstruct the matrix to verify perfect reconstruction.

Create 3-D matrix.

```
M = magic(8);
X = repmat(M,[1 1 8]);
```

Obtain the 3-D discrete wavelet transform of the matrix and reconstruct the input based on the 3-D approximation and detail coefficients.

```
wd = wavedec3(X,2,'db2');
XR = waverec3(wd);
```

Verify perfect reconstruction using the wavelet decomposition down to level 2.

```
err1 = max(abs(X(:)-XR(:)))
```

```
err1 = 8.6057e-11
```

Verify that the data matrix is the sum of the approximation and the details from levels 2 and 1. Reconstruct the sum of components different from the lowpass component and check that $X = A + D$.

```
A = waverec3(wd,'LLL');
D = waverec3(wd,'d');
err2 = max(abs(X(:)-A(:)-D(:)))
```

```
err2 = 8.6054e-11
```

Compare waverec3 and idwt3

Compare level-1 reconstructions based on the filtering operations 'LLH' using idwt3 and waverec3.

```
M = magic(8);
X = repmat(M,[1 1 8]);
wd = wavedec3(X,2,'db2','mode','per');
dwtOut = dwt3(X,'db2');
Xr = idwt3(dwtOut,'LLH');
Xrec = waverec3(wd,'LLH',1);
norm(Xr(:)-Xrec(:))
```

```
ans = 2.7511e-14
```

See Also

`idwt3` | `wavedec3` | `waveinfo`

Introduced in R2010a

wavsupport

Wavelet support

Syntax

```
[LB,UB] = wavsupport(wname)
```

Description

`[LB,UB] = wavsupport(wname)` returns the lower bound, LB, and upper bound, UB, of the support for the wavelet specified by `wname`. `wname` is any valid wavelet. For real-valued wavelets with and without scaling functions and complex-valued wavelets without scaling functions (wavelets type 3,4, and 5), the bounds indicate the effective support of the wavelet. For orthogonal and biorthogonal wavelets (type 1 and type 2), the lower and upper bounds are $-0.5*(LF-1)$ and $0.5*(LF-1)$, where LF is the length of the wavelet filter.

Examples

Support of Haar Wavelet

Return the lower bound and upper bound of the support for the Haar wavelet.

```
[LB, UB] = wavsupport('haar')
```

```
LB = -0.5000
```

```
UB = 0.5000
```

Compare LB and UB to the lower and upper bounds for orthogonal and biorthogonal wavelets (type 1 and type 2).

```
LowerBound = -0.5*(2-1);
```

```
UpperBound = 0.5*(2-1);
```

Support of Complex-Valued Gaussian Wavelet

Return the lower bound and upper bound of the support for the complex-valued Gaussian wavelet.

```
[LB,UB] = wavsupport( 'cgau3' )
```

```
LB = -5
```

```
UB = 5
```

See Also

wavemngr

Introduced in R2010b

wbmpen

Penalized threshold for wavelet 1-D or 2-D denoising

Syntax

```
THR = wbmpen(C,L,SIGMA,ALPHA)
wbmpen(C,L,SIGMA,ALPHA,ARG)
```

Description

`THR = wbmpen(C,L,SIGMA,ALPHA)` returns global threshold `THR` for denoising. `THR` is obtained by a wavelet coefficients selection rule using a penalization method provided by Birgé-Massart.

`[C,L]` is the wavelet decomposition structure of the signal or image to be denoised.

`SIGMA` is the standard deviation of the zero mean Gaussian white noise in denoising model (see `wnoisest` for more information).

`ALPHA` is a tuning parameter for the penalty term. It must be a real number greater than 1. The sparsity of the wavelet representation of the denoised signal or image grows with `ALPHA`. Typically `ALPHA = 2`.

`THR` minimizes the penalized criterion given by the following:

Let t^* be the minimizer of

$$\text{crit}(t) = -\sum(c(k)^2, k \leq t) + 2 \cdot \text{SIGMA}^2 \cdot t \cdot (\text{ALPHA} + \log(n/t))$$

where $c(k)$ are the wavelet coefficients sorted in decreasing order of their absolute value and n is the number of coefficients; then $\text{THR} = |c(t^*)|$.

`wbmpen(C,L,SIGMA,ALPHA,ARG)` computes the global threshold and, in addition, plots three curves:

- $2 \cdot \text{SIGMA}^2 \cdot t \cdot (\text{ALPHA} + \log(n/t))$

- `sum(c(k)^2,k-<=t)`
- `crit(t)`

Examples

```
% Example 1: Signal denoising.
% Load noisy bumps signal.
load noisbump; x = noisbump;

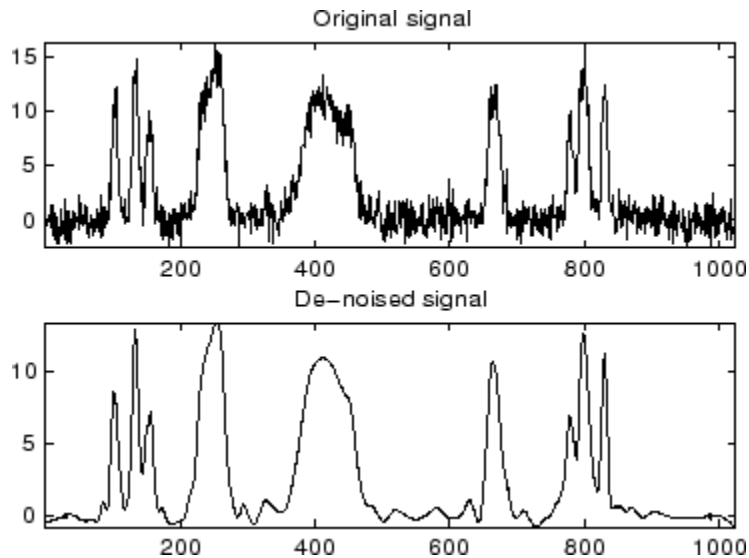
% Perform a wavelet decomposition of the signal
% at level 5 using sym6.
wname = 'sym6'; lev = 5;
[c,l] = wavedec(x,lev,wname);
% Estimate the noise standard deviation from the
% detail coefficients at level 1, using wnoisest.
sigma = wnoisest(c,l,1);

% Use wbmphen for selecting global threshold
% for signal denoising, using the tuning parameter.
alpha = 2;
thr = wbmphen(c,l,sigma,alpha)
thr =

    2.7681

% Use wdencomp for denoising the signal using the above
% threshold with soft thresholding and approximation kept.
keepapp = 1;
xd = wdencomp('gbl',c,l,wname,lev,thr,'s',keepapp);

% Plot original and denoised signals.
figure(1)
subplot(211), plot(x), title('Original signal')
subplot(212), plot(xd), title('De-noised signal')
```



```

% Example 2: Image denoising.
% Load original image.
load noiswom;
nbc = size(map,1);

% Perform a wavelet decomposition of the image
% at level 3 using coif2.
wname = 'coif2'; lev = 3;
[c,s] = wavedec2(X,lev,wname);

% Estimate the noise standard deviation from the
% detail coefficients at level 1.
det1 = detcoef2('compact',c,s,1);
sigma = median(abs(det1))/0.6745;

% Use wbmpen for selecting global threshold
% for image denoising.
alpha = 1.2;
thr = wbmpen(c,l,sigma,alpha)

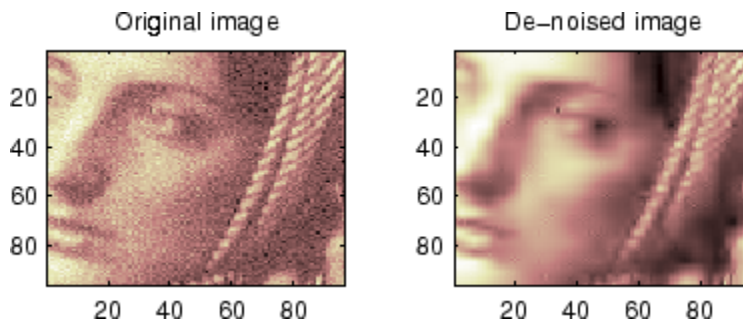
thr =

    36.0621

```

```
% Use wdencmp for denoising the image using the above
% thresholds with soft thresholding and approximation kept.
keepapp = 1;
xd = wdencmp('gbl',c,s,wname,lev,thr,'s',keepapp);

% Plot original and denoised images.
figure(2)
colormap(pink(nbc));
subplot(221), image(wcodemat(X,nbc))
title('Original image')
subplot(222), image(wcodemat(xd,nbc))
title('De-noised image')
```



See Also

wden | wdencmp | wdenoise | wpbmpen | wpcnmp

Introduced before R2006a

wcodemat

Extended pseudocolor matrix scaling

Syntax

```
Y = wcodemat(X)
Y = wcodemat(X,NBCODES)
Y = wcodemat(X,NBCODES,OPT)
Y = wcodemat(X,NBCODES,OPT,ABSOL)
```

Description

wcodemat rescales an input matrix to a specified range for display. If the specified range is the full range of the current colormap, wcodemat is similar in behavior to `imagesc`.

`Y = wcodemat(X)` rescales the matrix `X` to integers in the range `[1,16]`.

`Y = wcodemat(X,NBCODES)` rescales the input `X` as integers in the range `[1,NBCODES]`. The default value of `NBCODES` is 16.

`Y = wcodemat(X,NBCODES,OPT)` rescales the matrix along the dimension specified by `OPT`. `OPT` can be one of: 'column' (or 'c'), 'row' (or 'r'), and 'mat' (or 'm'). 'rows' scales `X` row-wise, 'column' scales `X` column-wise, and 'mat' scales `X` globally. The default value of `OPT` is 'mat'.

`Y = wcodemat(X,NBCODES,OPT,ABSOL)` rescales the input matrix `X` based on the absolute values of the entries in `X` if `ABSOL` is nonzero, or based on the signed values of `X` if `ABSOL` is equal to zero. The default value of `ABSOL` is 1.

Examples

Extended Pseudocolor Matrix Scaling

Scale level-one approximation coefficients globally to the full range of the colormap.

Load an image.

```
load woman;
```

Get the range of the colormap.

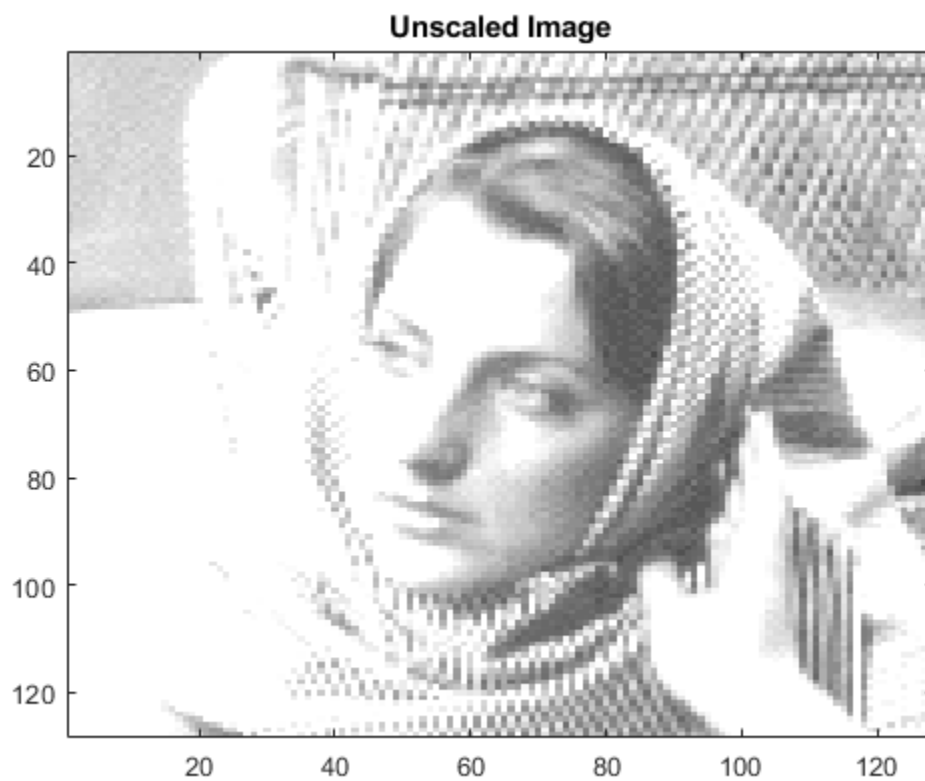
```
NBCOL = size(map,1);
```

Obtain the 2D dwt using the Haar wavelet.

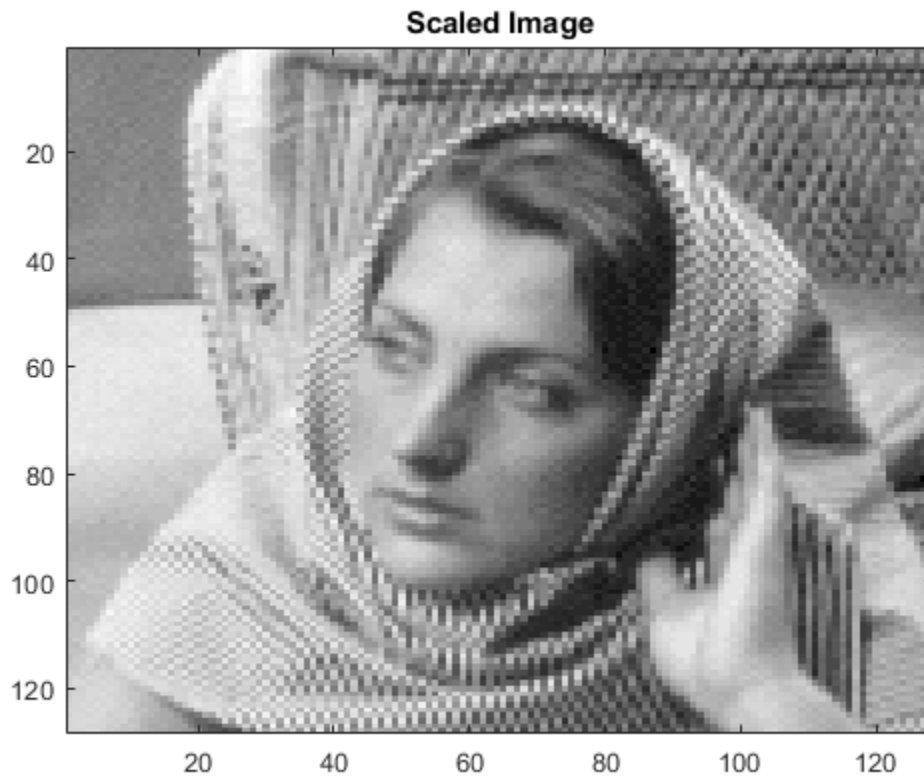
```
[cA1,cH1,cV1,cD1] = dwt2(X,'db1');
```

Display without scaling and with scaling.

```
image(cA1);  
colormap(map);  
title('Unscaled Image');
```



```
figure
image(wcodemat(cA1,NBCOL));
colormap(map);
title('Scaled Image');
```



Introduced before R2006a

wcoher

(Not recommended) Wavelet coherence

Note wcoher is not recommended. Use wcoherence instead.

Syntax

```
WCOH = wcoher(Sig1,Sig2,Scales,wname)
WCOH = wcoher(...,Name,Value)
[WCOH,WCS] = wcoher(...)
[WCOH,WCS,CWT_S1,CWT_S2] = wcoher(...)
[...] = wcoher(...,'plot')
```

Description

`WCOH = wcoher(Sig1,Sig2,Scales,wname)` returns the wavelet coherence for the input signals `Sig1` and `Sig2` using the wavelet specified in `wname` at the scales in `Scales`. The input signals must be real-valued and equal in length.

`WCOH = wcoher(...,Name,Value)` returns the wavelet coherence with additional options specified by one or more `Name,Value` pair arguments.

`[WCOH,WCS] = wcoher(...)` returns the wavelet cross spectrum.

`[WCOH,WCS,CWT_S1,CWT_S2] = wcoher(...)` returns the continuous wavelet transforms of `Sig1` and `Sig2`.

`[...] = wcoher(...,'plot')` displays the modulus and phase of the wavelet cross spectrum.

Input Arguments

Sig1

A real-valued one-dimensional input signal. **Sig1** is a row or column vector.

Sig2

A real-valued one-dimensional input signal. **Sig2** is a row or column vector.

Scales

Scales is a vector of real-valued, positive scales at which to compute the wavelet coherence.

wname

Wavelet used in the wavelet coherence. **wname** is any valid wavelet name.

Name-Value Pair Arguments

asc

Scale factor for arrows in quiver plot. **wcoher** represents the phase using **quiver**. **asc** corresponds to the **scale** input argument in **quiver**.

Default: 1

nas

Number of arrows in scale. Together with the number of scales, **nas** determines the spacing between the y coordinates in the input to **quiver**. The y input to **quiver** is `1:length(Scales)/(nas-1):Scales(end)`

Default: 20

nsw

Length of smoothing window in scale. **nsw** is a positive integer that specifies the length of a moving average filter in scale.

Default: 1

ntw

Length of smoothing window in time. `ntw` is a positive integer that specifies the length of a moving average filter in time.

Default: `min[20,0.05*length(Sig1)]`

plot

Type of plot. `plot` is one of the following:

- 'cwt'

Displays the continuous wavelet transforms of signals 1 and 2.

- 'wcs'

Displays the wavelet cross spectrum.

- 'wcoh'

Displays the phase of the wavelet cross spectrum.

- 'all'

Displays all plots in separate figures.

Output Arguments

WCOH

Wavelet coherence.

WCS

Wavelet cross spectrum.

CWT_S1

Continuous wavelet transform of signal 1.

CWT_S2

Continuous wavelet transform of signal 2.

Examples

Wavelet coherence of sine waves in noise with delay:

```
t = linspace(0,1,2048);
x = sin(16*pi*t)+0.5*randn(1,2048);
y = sin(16*pi*t+pi/4)+0.5*randn(1,2048);
wname = 'cgau3';
scales = 1:512;
ntw = 21; % smoothing parameter
% Display the modulus and phased of the wavelet cross spectrum.
wcoher(x,y,scales,wname,'ntw',ntw,'plot');
```

Sine wave and Doppler signal:

```
t = linspace(0,1,1024);
x = -sin(8*pi*t) + 0.4*randn(1,1024);
x = x/max(abs(x));
y = wnoise('doppler',10);
wname = 'cgau3';
scales = 1:512;
ntw = 21; % smoothing parameter
% Display of the CWT of the two signals.
wcoher(x,y,scales,wname,'ntw',ntw,'plot','cwt');
% Display of the wavelet cross spectrum.
wcoher(x,y,scales,wname,'ntw',ntw,'nsw',1,'plot','wcs');
% Display of the modulus and phased of the wavelet cross spectrum.
wcoher(x,y,scales,wname,'ntw',ntw,'plot');
```

Definitions

Wavelet Cross Spectrum

The wavelet cross spectrum of two time series, x and y is:

$$C_{xy}(a,b) = S(C_x^*(a,b)C_y(a,b))$$

where $C_x(a,b)$ and $C_y(a,b)$ denote the continuous wavelet transforms of x and y at scales a and positions b . The superscript $*$ is the complex conjugate and S is a smoothing operator in time and scale.

For real-valued time series, the wavelet cross spectrum is real-valued if you use a real-valued analyzing wavelet, and complex-valued if you use a complex-valued analyzing wavelet.

Wavelet Coherence

The wavelet coherence of two time series x and y is:

$$\frac{|S(C_x^*(a,b)C_y(a,b))|^2}{S(|C_x(a,b)|^2)S(|C_y(a,b)|^2)}$$

where $C_x(a,b)$ and $C_y(a,b)$ denote the continuous wavelet transforms of x and y at scales a and positions b . The superscript $*$ is the complex conjugate and S is a smoothing operator in time and scale.

For real-valued time series, the wavelet coherence is real-valued if you use a real-valued analyzing wavelet, and complex-valued if you use a complex-valued analyzing wavelet.

References

Grinsted, A, J.C. Moore, and S. Jevrejeva. "Application of the cross wavelet transform and wavelet coherence to geophysical time series. *Nonlinear Processes in Geophysics*. 11, 2004, pp. 561-566.

Torrence. C., and G. Compo. "A Practical Guide to Wavelet Analysis". *Bulletin of the American Meteorological Society*, 79, pp. 61-78.

See Also

cwt | wcoherence

Topics

"Compare Time-Frequency Content in Signals with Wavelet Coherence"

“Continuous Wavelet Analysis of Cusp Signal”
“Continuous and Discrete Wavelet Transforms”

Introduced in R2010b

wcoherence

Wavelet coherence and cross-spectrum

Syntax

```
wcoh = wcoherence(x,y)
[wcoh,wcs] = wcoherence(x,y)
[wcoh,wcs,period] = wcoherence(x,y,ts)
[wcoh,wcs,f] = wcoherence(x,y,fs)
[wcoh,wcs,f,coi] = wcoherence( ___ )
[wcoh,wcs,period,coi] = wcoherence( ___ ,ts)
wcoherence( ___ )
```

Description

`wcoh = wcoherence(x,y)` returns the magnitude-squared wavelet coherence, which is a measure of the correlation between signals `x` and `y` in the time-frequency plane. Wavelet coherence is useful for analyzing nonstationary signals. The inputs `x` and `y` must be equal length, 1-D, real-valued signals. The coherence is computed using the analytic Morlet wavelet.

`[wcoh,wcs] = wcoherence(x,y)` returns the wavelet cross-spectrum of `x` and `y`. You can use the phase of the wavelet cross-spectrum values to identify the relative lag between the input signals.

`[wcoh,wcs,period] = wcoherence(x,y,ts)` uses a duration `ts` with a positive, scalar input, as the sampling interval. The duration can be in years, days, hours, minutes, or seconds. `ts` is used to compute the scale-to period conversion, `period`. The period in an array of durations with the same time unit as specified in `ts`

`[wcoh,wcs,f] = wcoherence(x,y,fs)` uses the positive sampling frequency, `fs`, to compute the scale-to-frequency conversion, `f`. The `fs` input is in Hz.

`[wcoh,wcs,f,coi] = wcoherence(___)` returns the cone of influence, `coi`, for the wavelet coherence in cycles per sample. If you specify the sampling frequency, `fs`, the function returns both `f` and the cone of influence in Hz.


```

rng default;
t = 0:0.001:2;
x = cos(2*pi*10*t).*(t>=0.5 & t<1.1)+ ...
cos(2*pi*50*t).*(t>= 0.2 & t< 1.4)+0.25*randn(size(t));
y = sin(2*pi*10*t).*(t>=0.6 & t<1.2)+...
sin(2*pi*50*t).*(t>= 0.4 & t<1.6)+ 0.35*randn(size(t));
[wcoh,~,period,coi] = wcoherence(x,y,seconds(0.001));

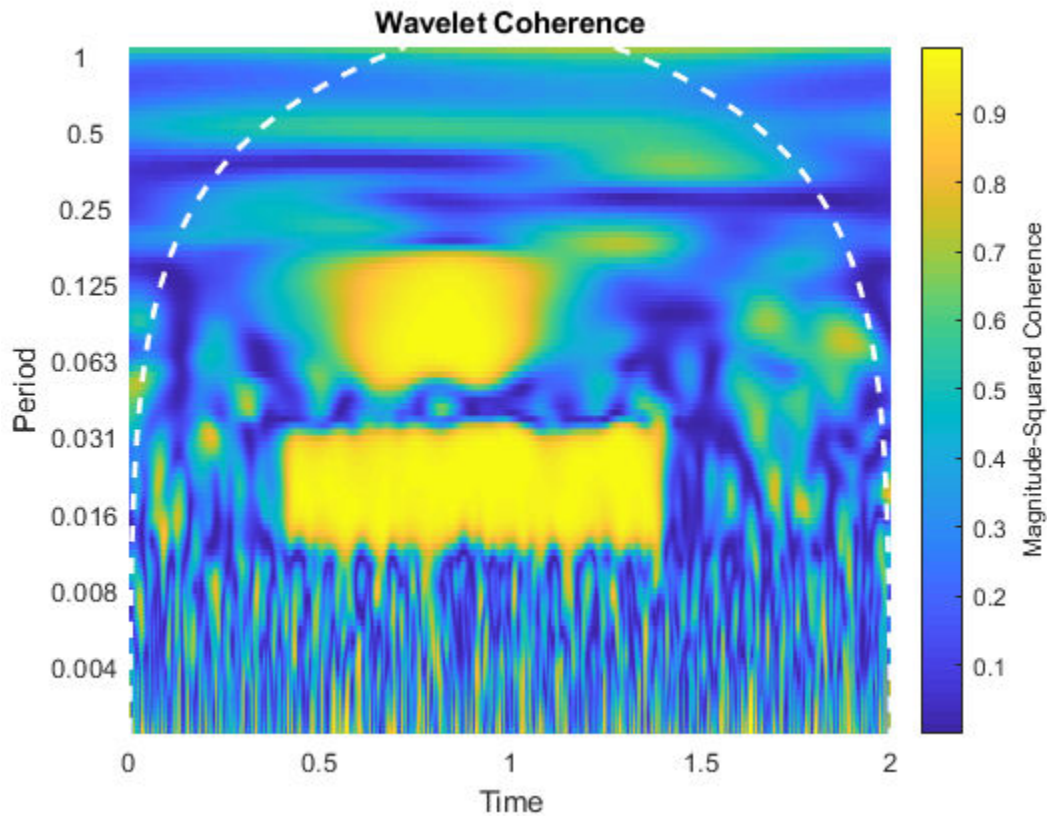
```

Use the `pcolor` command to plot the coherence and cone of influence.

```

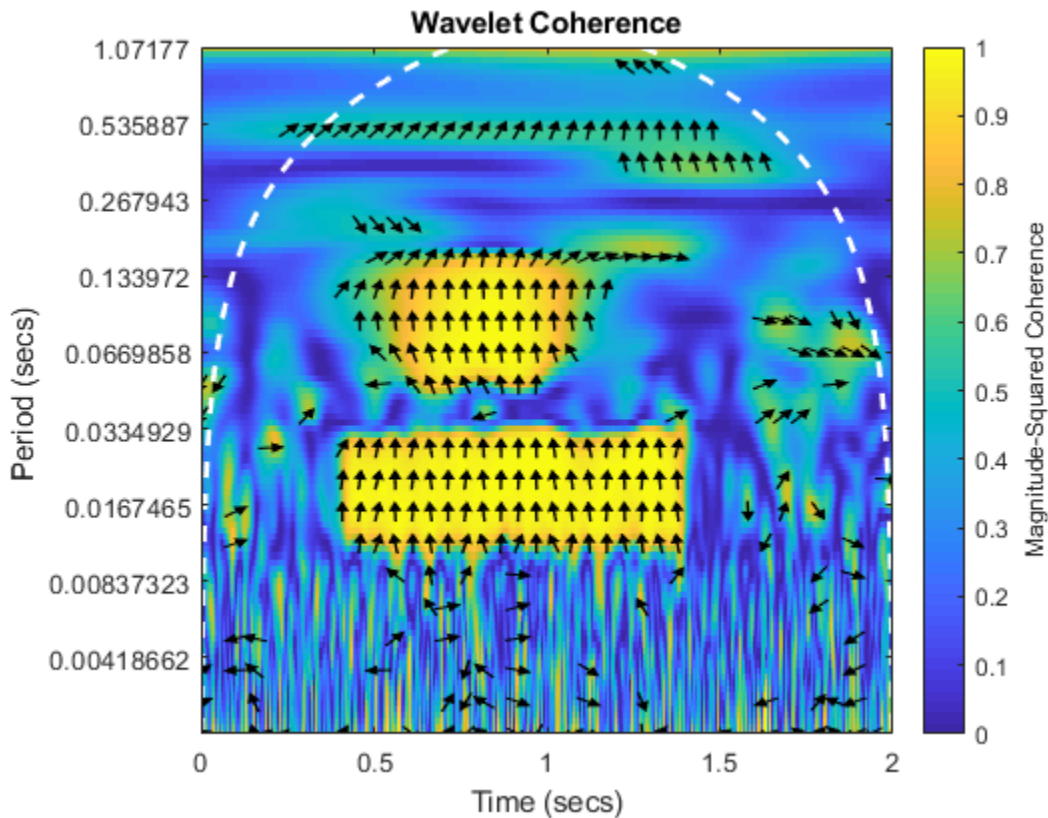
period = seconds(period);
coi = seconds(coi);
h = pcolor(t,log2(period),wcoh);
h.EdgeColor = 'none';
ax = gca;
ytick=round(pow2(ax.YTick),3);
ax.YTickLabel=ytick;
ax.XLabel.String='Time';
ax.YLabel.String='Period';
ax.Title.String = 'Wavelet Coherence';
hcol = colorbar;
hcol.Label.String = 'Magnitude-Squared Coherence';
hold on;
plot(ax,t,log2(coi),'w--','linewidth',2);

```



Use `wcoherence(x,y,seconds(0.001))` without any outputs arguments. This plot includes the phase arrows and the cone of influence.

```
wcoherence(x,y,seconds(0.001));
```



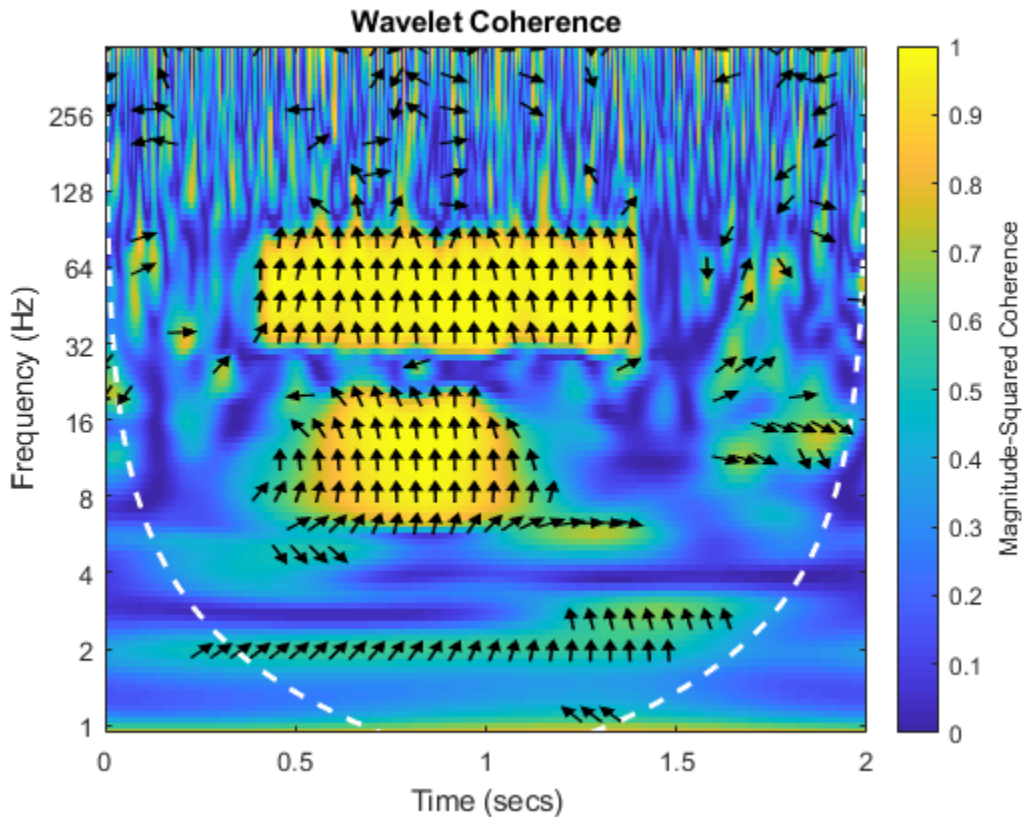
Effect of Sampling Frequency on Wavelet Coherence

Obtain the wavelet coherence for two signals, specifying a sampling frequency of 1000 Hz. Both signals consist of two sine waves (10 Hz and 50 Hz) in white noise. The sine waves have different time supports.

Set the random number generator to its default settings for reproducibility. Then, create the two signals and obtain the coherence.

```
rng default
t = 0:0.001:2;
```

```
x = cos(2*pi*10*t).*(t>=0.5 & t<1.1)+...
    cos(2*pi*50*t).*(t>= 0.2 & t< 1.4)+0.25*randn(size(t));
y = sin(2*pi*10*t).*(t>=0.6 & t<1.2)+...
    sin(2*pi*50*t).*(t>= 0.4 & t<1.6)+ 0.35*randn(size(t));
wcoherence(x,y,1000)
```



This coherence plot is flipped with respect to the coherence plot in the previous example, which specifies a sampling interval instead of a sampling frequency.

Obtain the scale-to-frequency conversion output in `f`.

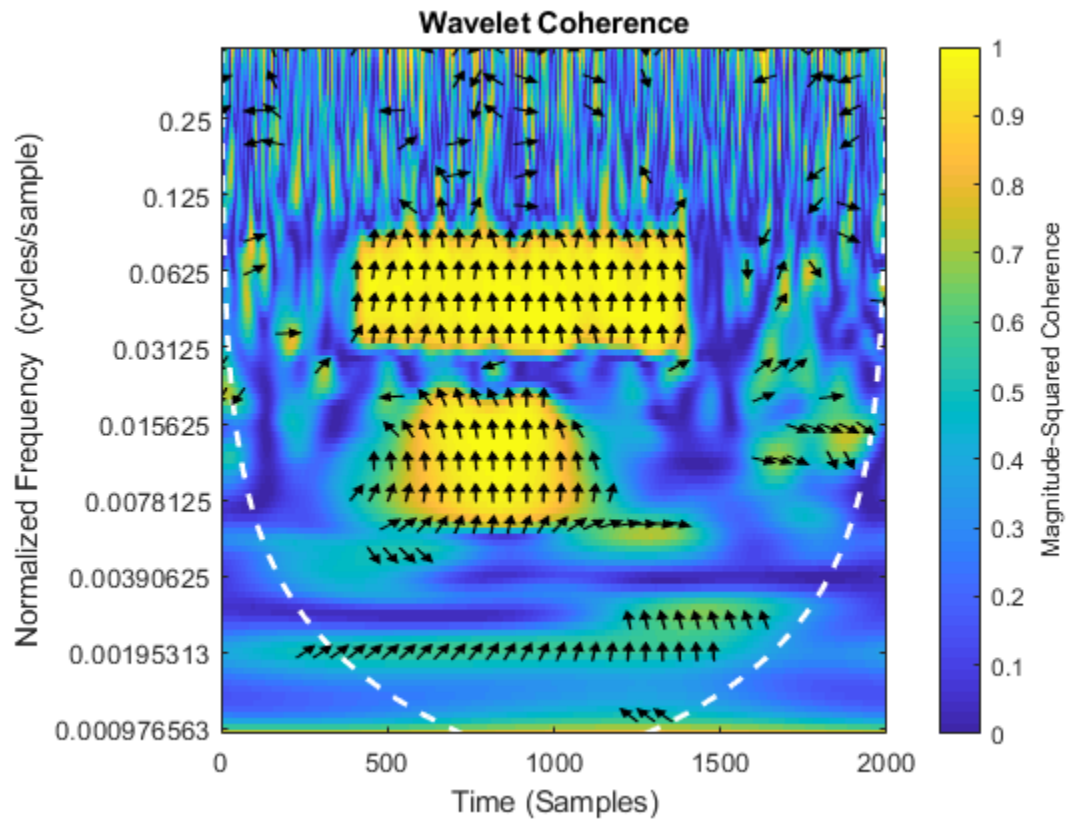
```
[wcoh,wcs,f] = wcoherence(x,y,1000);
```

Effect of Number of Smoothed Scales on Wavelet Coherence

Obtain the wavelet coherence for two signals. Both signals consist of two sine waves (10 Hz and 50 Hz) in white noise. Use the default number of scales to smooth. This value is equivalent to the number of voices per octave. Both values default to 12.

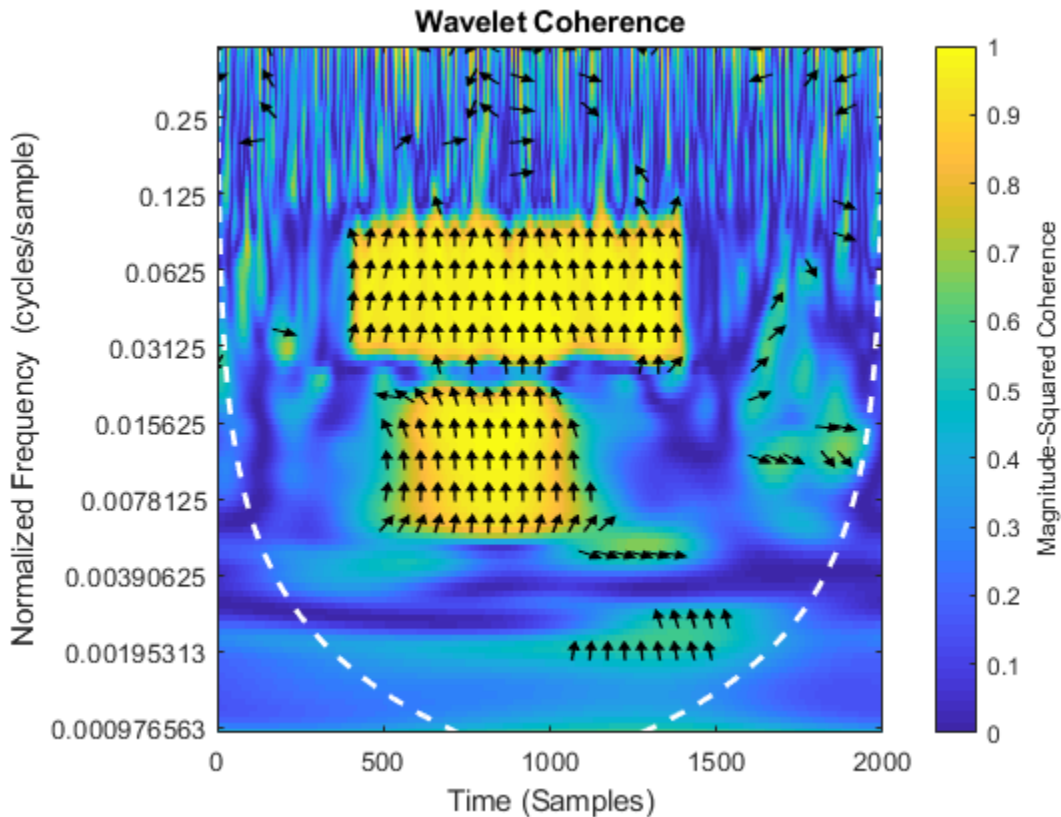
Set the random number generator to its default settings for reproducibility. Then, create the two signals and obtain the coherence.

```
rng default;
t = 0:0.001:2;
x = cos(2*pi*10*t).*(t>=0.5 & t<1.1)+ ...
cos(2*pi*50*t).*(t>= 0.2 & t< 1.4)+0.25*randn(size(t));
y = sin(2*pi*10*t).*(t>=0.6 & t<1.2)+...
sin(2*pi*50*t).*(t>= 0.4 & t<1.6)+ 0.35*randn(size(t));
wcoherence(x,y);
```



Set the number of scales to smooth to 16. The increased smoothing causes reduced low frequency resolution.

```
wcoherence(x,y,'NumScalesToSmooth',16);
```

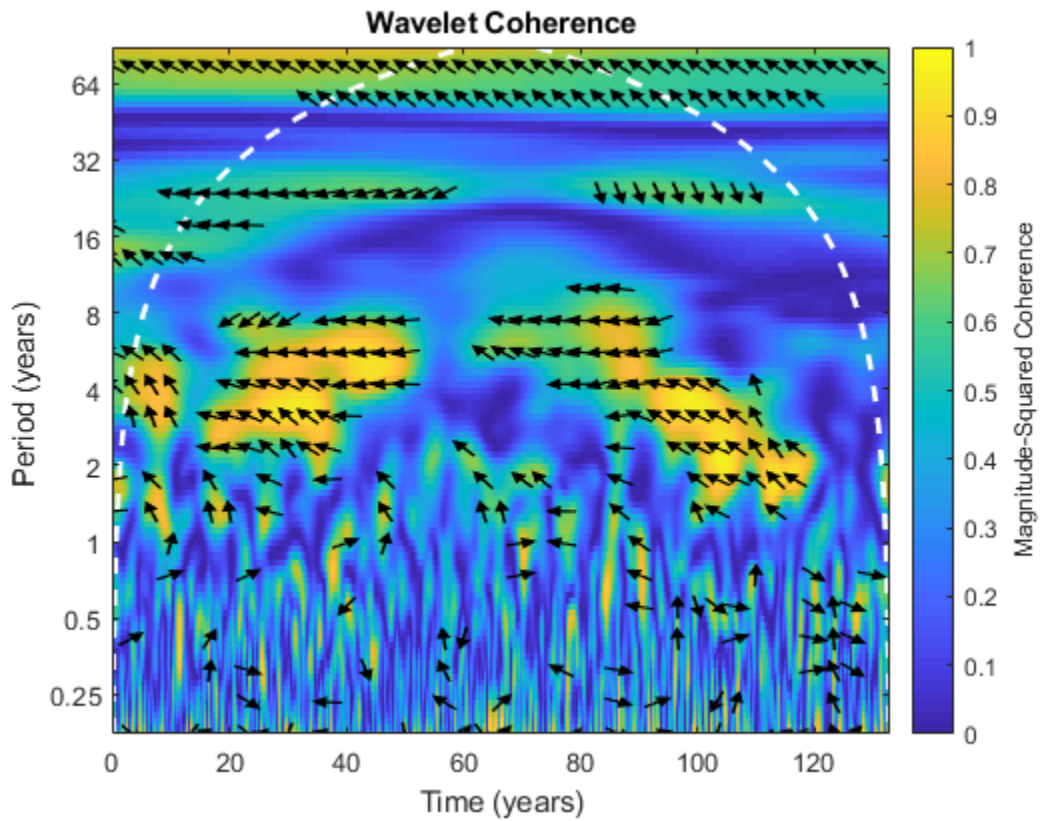


Effect of Phase Display Threshold on Wavelet Coherence of Weather Data

Compare the effects of using different phase display thresholds on the wavelet coherence.

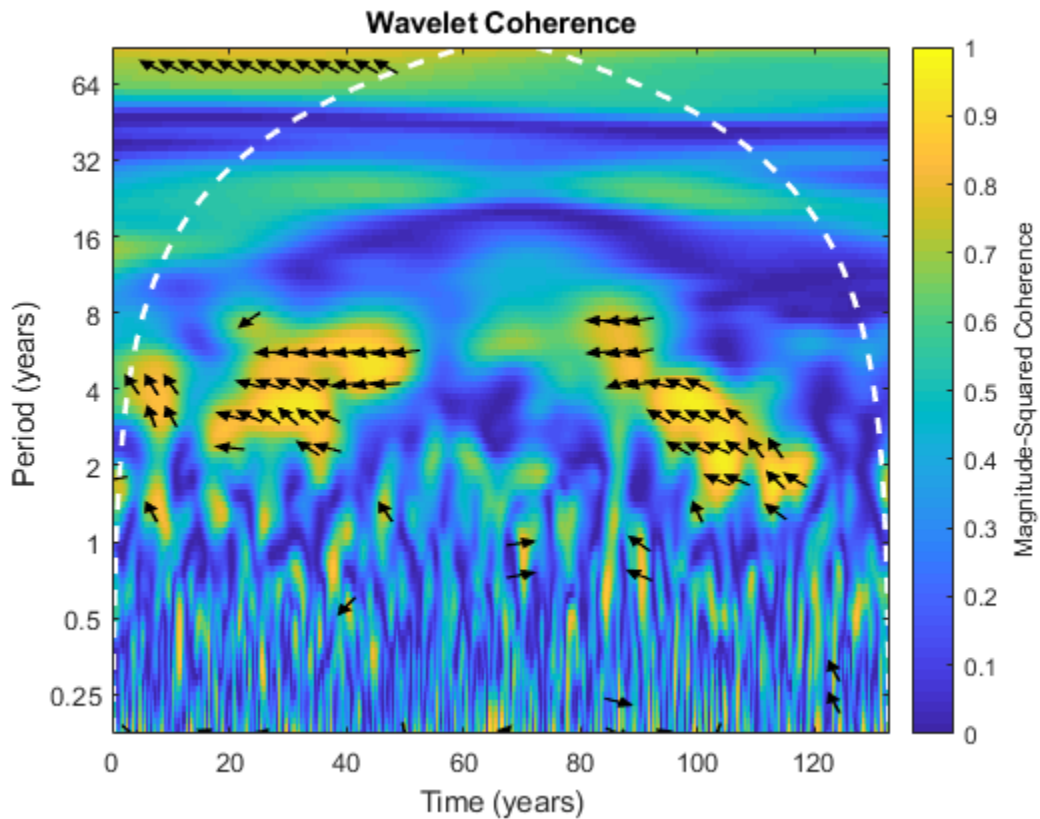
Plot the wavelet coherence between the El Nino time series and the All India Average Rainfall Index. The data are sampled monthly. Specify the sampling interval as 1/12 of a year to display the periods in years. Use the default phase display threshold of 0.5, which shows phase arrows only where the coherence is greater than or equal to 0.5.

```
load ninoairdata;
wcoherence(nino,air,years(1/12));
```



Set the phase display threshold to 0.7. The number of phase arrows decreases.

```
wcoherence(nino,air,years(1/12),'PhaseDisplayThreshold',0.7);
```

Input Arguments

x – Input signal

vector of real values

Input signal, specified as a vector of real values. x must be a 1-D, real-valued signal. The two input signals, x and y , must be the same length and must have at least four samples.

y – Input signal

vector of real values

Input signal, specified as vector of real values. `y` must be a 1-D, real-valued signal. The two input signals, `x` and `y`, must be the same length and must have at least four samples.

ts — Sampling interval

duration with positive scalar input

Sampling interval, also known as the sampling period, specified as a duration with positive scalar input. Valid durations are years, days, hours, seconds, and minutes. You can also use the `duration` function to specify `ts`. You cannot use calendar durations (`caldays`, `calweeks`, `calmonths`, `calquarters`, or `calyears`). You cannot specify both `ts` and `fs`.

fs — Sampling frequency

positive scalar

Sampling frequency, specified as a positive scalar.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

Example: `'PhaseDisplayThreshold', 0.7`; specifies the threshold for displaying phase vectors.

VoicesPerOctave — Number of voices per octave

12 (default) | integer from 10 to 32

Number of voices per octave to use in the wavelet coherence, specified as an integer from 10 to 32.

NumScalesToSmooth — Number of scales to smooth

number of voices per octave (default) | positive integer

Number of scales to smooth in time and scale, specified as a positive integer. This value must be less than one half the total number of scales. The function uses a moving average filter to smooth across scale. If you do not specify the number of scales to smooth, the value of `NumScalesToSmooth` defaults to the number of voices per octave. If your coherence is noisy, you can specify a larger `NumScalesToSmooth` value to smooth the coherence more.

NumOctaves — Number of octaves

positive integer

Number of octaves to use in the wavelet coherence, specified as a positive integer between 1 and $\text{floor}(\log_2(\text{numel}(x))) - 1$. If you do not need to examine lower frequency values, use a smaller NumOctaves value.

PhaseDisplayThreshold — Threshold for displaying phase vectors

0.5 (default) | real scalar between 0 and 1

Threshold for displaying phase vectors, specified as a real scalar between 0 and 1. This function displays phase vectors for regions with coherence greater than or equal to the specified threshold value. Lowering the threshold value displays more phase vectors. If you use wcoherence with any output arguments, the PhaseDisplayThreshold value is ignored.

Output Arguments

wcoh — Wavelet coherence

matrix

Wavelet coherence, returned as a matrix. The coherence is computed using the analytic Morlet wavelet over logarithmic scales, with a default value of 12 voices per octave. The default number of octaves is equal to $\text{floor}(\log_2(\text{numel}(x))) - 1$. If you do not specify a sampling interval, sampling frequency is assumed.

wcs — Wavelet cross spectrum

matrix of complex values

Wavelet cross-spectrum, returned as a matrix of complex values. You can use the phase of the wavelet cross-spectrum values to identify the relative lag between the input signals.

Data Types: double

Complex Number Support: Yes

period — Scale-to-period conversion

array of durations

Scale-to-period conversion, returned as an array of durations. The conversion values are computed from the sampling period specified in `ts`. Each `period` element has the same format as `ts`.

f — Scale-to-frequency conversion

vector

Scale-to-frequency conversion, returned as a vector. The vector contains the peak frequency values for the wavelets used to compute the coherence. If you want to output **f**, but do not specify a sampling frequency input, **fs**, the returned wavelet coherence is in cycles per sample.

coi — Cone of influence

array of doubles or array of durations

Cone of influence for the wavelet coherence, returned as either an array of doubles or array of durations. The cone of influence indicates where edge effects occur in the coherence data. If you specify a sampling frequency, **fs**, the cone of influence is in Hz. If you specify a sampling interval or period, **ts**, the cone of influence is in periods. Due to the edge effects, give less credence to areas of apparent high coherence that are outside or overlap the cone of influence. The cone of influence is indicated by a dashed line.

Definitions

Wavelet Cross Spectrum

The wavelet cross- spectrum is a measure of the distribution of power of two signals.

The wavelet cross spectrum of two time series, *x* and *y*, is:

$$C_{xy}(a, b) = S(C_x^*(a, b)C_y(a, b))$$

$C_x(a, b)$ and $C_y(a, b)$ denote the continuous wavelet transforms of *x* and *y* at scales *a* and positions *b*. The superscript * is the complex conjugate, and *S* is a smoothing operator in time and scale.

For real-valued time series, the wavelet cross-spectrum is real-valued if you use a real-valued analyzing wavelet, and complex-valued if you use a complex-valued analyzing wavelet.

Wavelet Coherence

Wavelet coherence is a measure of the correlation between two signals.

The wavelet coherence of two time series x and y is:

$$\frac{|S(C_x^*(a, b)C_y(a, b))|^2}{S(|C_x(a, b)|^2)S(|C_y(a, b)|^2)}$$

$C_x(a, b)$ and $C_y(a, b)$ denote the continuous wavelet transforms of x and y at scales a and positions b . The superscript $*$ is the complex conjugate and S is a smoothing operator in time and scale.

For real-valued time series, the wavelet coherence is real-valued if you use a real-valued analyzing wavelet, and complex-valued if you use a complex-valued analyzing wavelet.

References

- [1] Grinsted, A. J., C. Moore, and S. Jevrejeva. "Application of the cross wavelet transform and wavelet coherence to geophysical time series." *Nonlinear Processes in Geophysics*. Vol. 11, Issue 5/6, 2004, pp. 561-566.
- [2] Maraun, D., J. Kurths, and M. Holschneider. "Nonstationary Gaussian processes in wavelet domain: Synthesis, estimation and significance testing." *Physical Review E* 75. 2007, pp. 016707-1-016707-14.
- [3] Torrence, C., and P. Webster. "Interdecadal changes in the ENSO-Monsoon System." *Journal of Climate*. Vol. 12, 1999, pp. 2679-2690.

See Also

cwt | days | duration | hours | minutes | seconds | years

Introduced in R2016a

wcompress

True compression of images using wavelets

Syntax

```
wcompress('c',X,SAV_FILENAME,COMP_METHOD)
wcompress(...,'ParName1',ParVal1,'ParName2',ParVal2,...)
[COMPRAT,BPP] = wcompress('c',...)
XC = wcompress('u',SAV_FILENAME)
XC = wcompress('u',SAV_FILENAME,'plot')
XC = wcompress('u',SAV_FILENAME,'step')
```

Description

The `wcompress` command performs either compression or uncompression of grayscale or truecolor images.

More theoretical information on true compression is in “Wavelet Compression for Images” of the Wavelet Toolbox User's Guide.

Compression

`wcompress('c',X,SAV_FILENAME,COMP_METHOD)` compresses the image `X` using the compression method `COMP_METHOD`.

The compressed image is saved in the file `SAV_FILENAME`. You must have write permission in the current working directory or MATLAB will change directory to `tempdir` and write the `.wtc` file in that directory. `X` can be either a 2-D array containing an indexed image or a 3-D array of `uint8` containing a truecolor image. Both the row and column size of the image must be powers of two.

`wcompress('c',FILENAME,...)` loads the image `X` from the file `FILENAME` which is a MATLAB Supported Format (MSF) file: MAT-file or other image files (see `imread`).

`wcompress('c',I,...)` converts the indexed image `X = I{1}` to a truecolor image `Y` using the colormap `map = I{2}` and then compresses `Y`.

Note Data written to .wtc files uses `uint64` precision. In releases previous to R2016b, data was written using `uint32`. If your code is affected adversely by this change, use the `legacy` option to compress and uncompress your data using the previous behavior.

```
wcompress('c',X,SAV_FILENAME,COMP_METHOD,'legacy')
```

The valid compression methods are divided in three categories.

1 Progressive Coefficients Significance Methods (**PCSM**):

MATLAB Name	Compression Method Name
'ezw'	Embedded Zerotree Wavelet
'spiht'	Set Partitioning In Hierarchical Trees
'stw'	Spatial-orientation Tree Wavelet
'wdr'	Wavelet Difference Reduction
'aswdr'	Adaptively Scanned Wavelet Difference Reduction
'spiht_3d'	Set Partitioning In Hierarchical Trees 3D for truecolor images

For more details on these methods, see the references and especially Walker and also Said and Pearlman.

1 Coefficients Thresholding Methods (**CTM-1**):

MATLAB Name	Compression Method Name
'lvl_mmc'	Subband thresholding of coefficients and Huffman encoding

For more details on this method, see the Strang and Nguyen reference.

1 Coefficients Thresholding Methods (**CTM-2**):

MATLAB Name	Compression Method Name
'gbl_mmc_f'	Global thresholding of coefficients and fixed encoding
'gbl_mmc_h'	Global thresholding of coefficients and Huffman encoding

Note The Discrete Wavelet Transform uses the periodized extension mode.

All the compression methods use parameters which have default values. You can change these values using the following syntax:

```
wcompress(..., 'ParName1', ParVal1, 'ParName2', ParVal2, ...)
```

Some of the parameters are related to display or to data transform functionalities. The others are linked to the compression process itself.

Data transform parameters

- 'ParName' = 'wname' or 'WNAME' sets the wavelet name.

ParVal is a character vector or string scalar (see `waveletfamilies`). The default for is `bior4.4`

- 'ParName' = 'level' or 'LEVEL' sets the level of decomposition.

ParVal is an integer such that: $1 \leq \text{level} \leq \text{levmax}$ which is the maximum possible level (see `wmaxlev`).

The default level depends on the method:

- for **PCSM** methods `level` is equal to `levmax`.
- for **CTM** methods `level` is equal to `fix(levmax/2)`

- 'ParName' = 'it' or 'IT' sets Image type Transform.

ParVal must be one of the following:

'n' : no transformation (default), image type (truecolor or grayscale) is automatically detected.

'g' : grayscale transformation type.

'c' : color transformation type (RGB `uint8`).

- 'ParName' = 'cc' or 'CC' sets Color Conversion parameter if X is a truecolor image.

ParVal must be one of the following:

'rgb' or 'none' : No conversion (default).

'yuv' : YUV color space transform.

'klt' : Karhunen-Loeve transform.

'yiq' : YIQ color space transform.

'xyz' : CIEXYZ color space transform.

Parameter for Progressive Coefficients Significance Methods (PCSM)

- 'ParName' = 'maxloop' or 'MAXLOOP' sets the maximum number of steps for the compression algorithm.

ParVal must be a positive integer or Inf (default is 10).

Parameters for Coefficients Thresholding Methods (CTM-1)

Either of the following parameters may be used:

- 'ParName' = 'bpp' or 'BPP' sets the bit-per-pixel ratio.

ParVal must be such that $0 \leq \text{ParVal} \leq 8$ (grayscale) or 24 (truecolor).

- 'ParName' = 'comprat' or 'COMPRAT' sets the compression ratio.

ParVal must be such that $0 \leq \text{ParVal} \leq 100$.

Parameters for Coefficients Thresholding Methods (CTM-2)

Two parameters may be used. The first is related to the threshold and the second is the number of classes for quantization.

The first one may be chosen among the five following parameters:

- 'ParName' = 'threshold' or 'THRESHOLD' sets the threshold value for compression.

ParVal must be a positive (or zero) real number.

- 'ParName' = 'nbcfs' or 'NBCFS' sets the number of preserved coefficients in the wavelet decomposition.

ParVal must be an integer such that: $0 \leq \text{ParVal} \leq$ total number of coefficients of wavelet decomposition.

- 'ParName' = 'percfs' or 'PERCFS' sets the percentage of preserved coefficients in the wavelet decomposition.

ParVal must be a real number such that: $0 \leq \text{ParVal} \leq 100$.

- 'ParName' = 'bpp' or 'BPP' sets the bit-per-pixel ratio.

ParVal must be such that: $0 \leq \text{ParVal} \leq 8$ (grayscale) or 24 (truecolor)

- 'ParName' = 'comprat' or 'COMPRAT' sets the compression ratio.

ParVal must be such that: $0 \leq \text{ParVal} \leq 100$.

The second parameter sets the number of classes for quantization:

- 'ParName' = 'nbclas' or 'NBCLAS' sets the number of classes.

ParVal must be a real number such that: $2 \leq \text{ParVal} \leq 200$.

Display parameter

- 'ParName' = 'plotpar' or 'PLOTPAR' sets the plot parameter.

ParVal must be one of the following:

'plot' or 0: plots only the compressed image.

'step' or 1: displays each step of the encoding process (only for **PCSM** methods).

[COMPRAT, BPP] = wcompress('c', ...) returns the compression ratio COMPRAT and the bit_per_pixel ratio BPP.

Uncompression

XC = wcompress('u', SAV_FILENAME) uncompresses the file SAV_FILENAME, which contains the compressed image, and returns the image XC. Depending on the initial compressed image, XC can be a 2-D array containing either an indexed image or a 3-D array of uint8 containing a truecolor image.

XC = wcompress('u', SAV_FILENAME, 'plot') plots the uncompressed image.

`XC = wcompress('u', SAV_FILENAME, 'step')` shows the step-by-step uncompression, only for **PCSM** methods.

Examples

Image Compression Using Basic Parameters

This example shows how to compress and uncompress the jpeg image `arms.jpg`.

Use the spatial orientation tree wavelet ('stw') compression method and save the compressed image to a file.

```
wcompress('c', 'arms.jpg', 'comp_arms.wtc', 'stw');
```

Load the stored image and display the step-by-step uncompression to produce the uncompressed image.

```
wcompress('u', 'comp_arms.wtc', 'step');
```

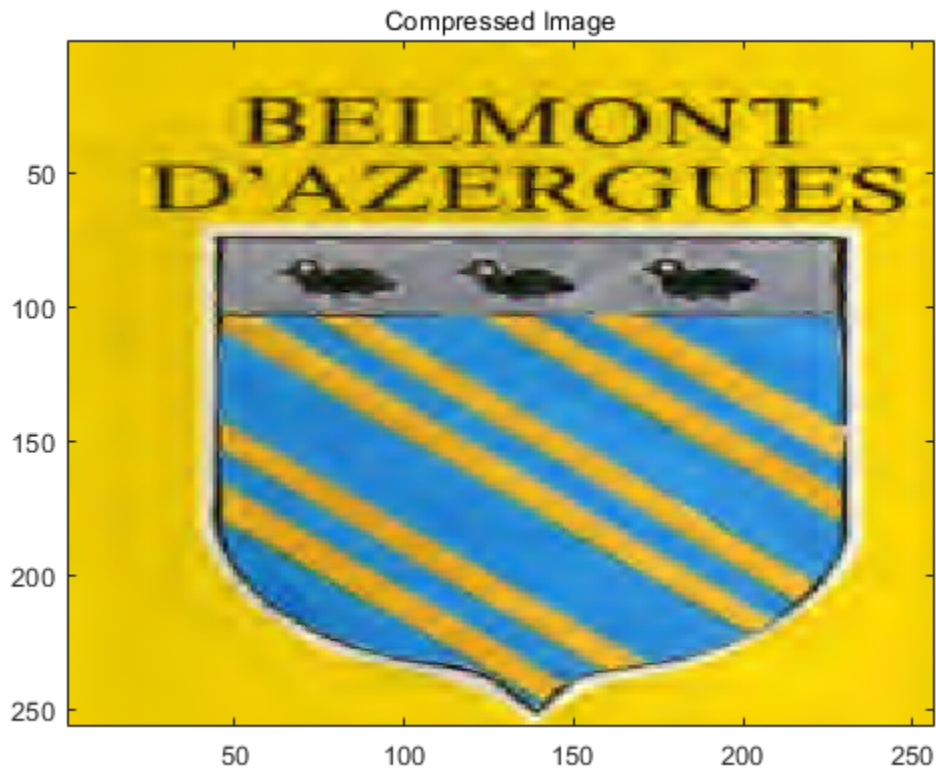


Image Compression and Uncompression Using Advanced Parameters.

This example shows how to compress a jpeg image using the adaptively scanned wavelet difference reduction compression method ('aswdr'). The conversion color ('cc') uses the Karhunen-Loeve transform ('klt'). The maximum number of loops ('maxloop') is set to 11 and the plot type ('plotpar') is set to step through the compression. Show the compression ratio (cratio) and the bit-per-pixel ratio (bpp), which indicate the quality of the compression.

```
[cratio,bpp] = wcompress('c','woodstatue.jpg','woodstatue.wtc', ...  
                        'aswdr','cc','klt','maxloop',11,'plotpar','step');
```

```
cratio  
bpp
```

```
cratio =  
    3.0792
```

```
bpp =  
    0.7390
```



Load the compressed image and step through the uncompression process.

```
wcompress('u', 'woodstatue.wtc', 'step');
```



Compression and Uncompression of a Grayscale Image

This example shows how to compress a grayscale image using the set partitioning in hierarchical trees ('spih') compression method. It also computes the mean square error (MSE) and the peak signal to noise ratio (PSNR) error values. You use these two measures to quantify the error between two images. The PSNR is expressed in decibels.

Load the image and store it in a file.

```
load mask;
[cr,bpp] = wcompress('c',X,'mask.wtc','spiht','maxloop',12)

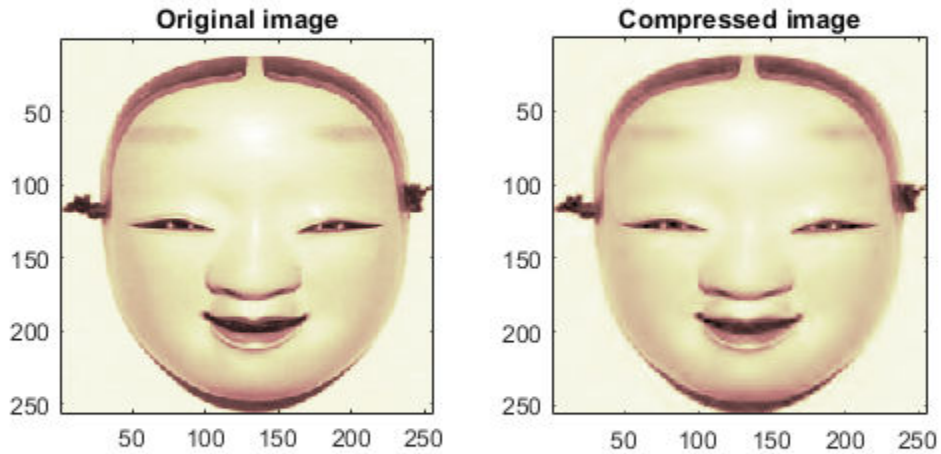
cr = 2.8610
bpp = 0.2289
```

Load the stored image from the file, uncompress it, and delete the file.

```
Xc = wcompress('u','mask.wtc');
delete('mask.wtc')
```

Display the original and compressed images.

```
colormap(pink(255))
subplot(1,2,1); image(X); title('Original image')
axis square
subplot(1,2,2); image(Xc); title('Compressed image')
axis square
```



Compute the MSE and PSNR.

```
D = abs(X-Xc).^2;  
mse = sum(D(:))/numel(X)  
  
mse = 33.6564  
  
psnr = 10*log10(255*255/mse)  
  
psnr = 32.8601
```


Compression and Uncompression of a Truecolor Image

This example shows how to compress a truecolor image using the set partitioning in hierarchical trees - 3D ('spiht_3D') compression method.

Load, compress, and store the image in a file. Plot the original and compressed images. Display the compression ratio ('cratio') and the bits-per-pixel ('bpp'), which indicate the quality of the compression.

```
load mask;
X = imread('wpeppers.jpg');
[cratio,bpp] = wcompress('c',X,'wpeppers.wtc','spiht','maxloop',12)

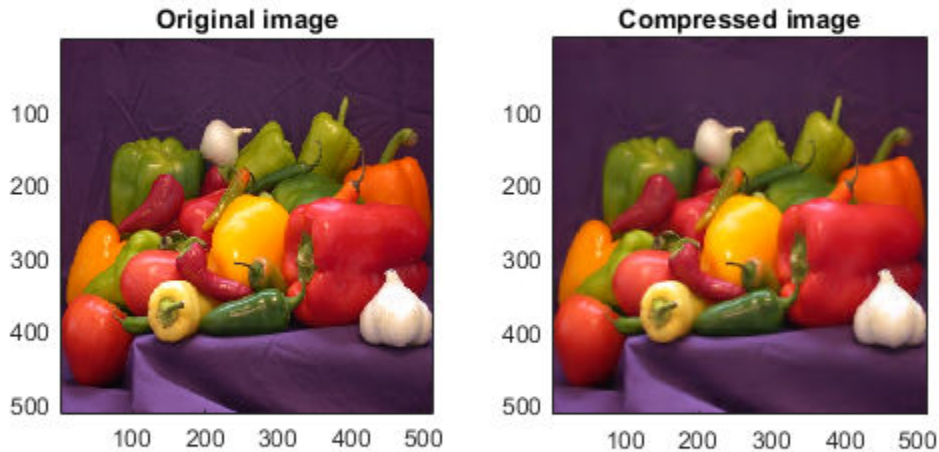
cratio = 1.6549

bpp = 0.3972

Xc = wcompress('u','wpeppers.wtc');
delete('wpeppers.wtc')
```

Display the original and compressed images.

```
subplot(1,2,1)
image(X)
title('Original image')
axis square
subplot(1,2,2)
image(Xc)
title('Compressed image')
axis square
```



Compute the mean square error (MSE) and the peak signal-to-noise ratio (PSNR) error values. You use these two measures to quantify the error between two images. The PSNR is expressed in decibels.

```
D = abs(double(X) - double(Xc)).^2;  
mse = sum(D(:))/numel(X)
```

```
mse = 26.7808
```

```
psnr = 10*log10(255*255/mse)
```

```
psnr = 33.8526
```

References

Christophe, E., C. Mailhes, P. Duhamel (2006), "Adaptation of zerotrees using signed binary digit representations for 3 dimensional image coding," *EURASIP Journal on Image and Video Processing*, 2007, to appear in the special issue on Wavelets in Source Coding, Communications, and Networks, Paper ID 54679.

Misiti, M., Y. Misiti, G. Oppenheim, J.-M. Poggi (2007), *Wavelets and their applications*, ISTE DSP Series.

Said A., W.A. Pearlman (1996), "A new, fast, and efficient image codec based on set partitioning in hierarchical trees," *IEEE Trans. on Circuits and Systems for Video Technology*, Vol. 6, No. 3, pp. 243-250.

Shapiro J.M. (1993), "Embedded image coding using zerotrees of wavelet coefficients", *IEEE Trans. Signal Proc.*, Vol. 41, No. 12, pp. 3445-3462.

Strang, G.; T. Nguyen (1996), *Wavelets and Filter Banks*, Wellesley-Cambridge Press.

Walker J.S. (1999), "Wavelet-Based Image Compression," University of Wisconsin, Eau Claire, Wisconsin, USA, , Sub-chapter of CRC Press book: *Transform and Data Compression. A Primer on Wavelets and Their Scientific Applications*.

See Also

`imread` | `imwrite` | `path` | `tempdir` | `wmaxlev`

Introduced in R2008b

wdcbm

Thresholds for wavelet 1-D using Birgé-Massart strategy

Syntax

```
[THR,NKEEP] = wdcbm(C,L,ALPHA,M)
wdcbm(C,L,ALPHA)
wdcbm(C,L,ALPHA,L(1))
```

Description

[THR,NKEEP] = wdcbm(C,L,ALPHA,M) returns level-dependent thresholds THR and numbers of coefficients to be kept NKEEP, for denoising or compression. THR is obtained using a wavelet coefficients selection rule based on the Birgé-Massart strategy.

[C,L] is the wavelet decomposition structure of the signal to be denoised or compressed, at level $j = \text{length}(L) - 2$. ALPHA and M must be real numbers greater than 1.

THR is a vector of length j ; THR(i) contains the threshold for level i .

NKEEP is a vector of length j ; NKEEP(i) contains the number of coefficients to be kept at level i .

j , M and ALPHA define the strategy:

- At level $j+1$ (and coarser levels), everything is kept.
- For level i from 1 to j , the n_i largest coefficients are kept with $n_i = M / (j+2-i)^{\text{ALPHA}}$.

Typically ALPHA = 1.5 for compression and ALPHA = 3 for denoising.

A default value for M is $M = L(1)$, the number of the coarsest approximation coefficients, since the previous formula leads for $i = j+1$, to $n_{j+1} = M = L(1)$. Recommended values for M are from $L(1)$ to $2*L(1)$.

wdcbm(C,L,ALPHA) is equivalent to wdcbm(C,L,ALPHA,L(1)).

Examples

```

% Load electrical signal and select a part of it.
load leleccum; indx = 2600:3100;
x = leleccum(indx);

% Perform a wavelet decomposition of the signal
% at level 5 using db3.
wname = 'db3'; lev = 5;
[c,l] = wavedec(x,lev,wname);

% Use wdcbm for selecting level dependent thresholds
% for signal compression using the advised parameters.
alpha = 1.5; m = l(1);
[thr,nkeep] = wdcbm(c,l,alpha,m)

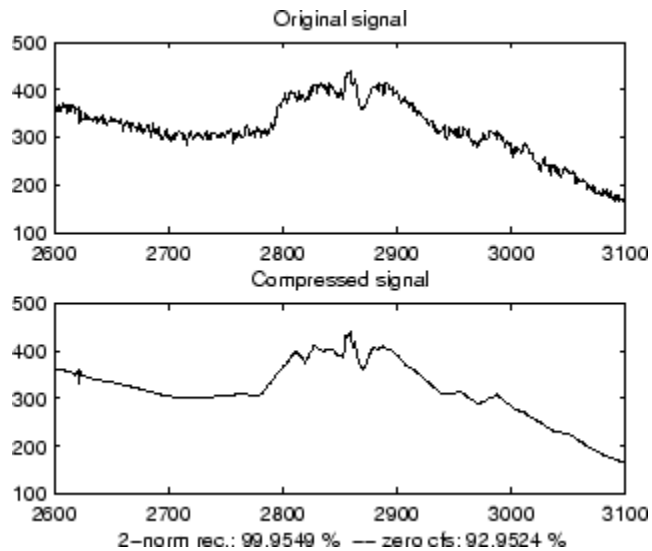
thr =
    19.5569    17.1415    20.2599    42.8959    15.0049

nkeep =
     1     2     3     4     7

% Use wdencomp for compressing the signal using the above
% thresholds with hard thresholding.
[xd,cxd,lxd,perf0,perfl2] = ...
    wdencomp('lvd',c,l,wname,lev,thr,'h');

% Plot original and compressed signals.
subplot(211), plot(indx,x), title('Original signal');
subplot(212), plot(indx,xd), title('Compressed signal');
xlabel1 = ['2-norm rec.: ',num2str(perfl2)];
xlabel2 = [' % -- zero cfs: ',num2str(perf0), ' %'];
xlabel([xlabel1 xlabel2]);

```



References

Birgé, L.; P. Massart (1997), "From model selection to adaptive estimation," in D. Pollard (ed), *Festschrift for L. Le Cam*, Springer, pp. 55-88.

See Also

`wden` | `wdencmp` | `wdenoise` | `wpdencmp`

Introduced before R2006a

wdcbm2

Thresholds for wavelet 2-D using Birgé-Massart strategy

Syntax

```
[THR,NKEEP] = wdcbm2(C,S,ALPHA,M)
wdcbm2(C,S,ALPHA)
wdcbm2(C,S,ALPHA,prod(S(1,:)))
```

Description

[THR,NKEEP] = wdcbm2(C,S,ALPHA,M) returns level-dependent thresholds THR and numbers of coefficients to be kept NKEEP, for de-noising or compression. THR is obtained using a wavelet coefficients selection rule based on the Birgé-Massart strategy.

[C,S] is the wavelet decomposition structure of the image to be de-noised or compressed, at level $j = \text{size}(S,1) - 2$.

ALPHA and M must be real numbers greater than 1.

THR is a matrix 3 by j; THR(:,i) contains the level dependent thresholds in the three orientations: horizontal, diagonal, and vertical, for level i.

NKEEP is a vector of length j; NKEEP(i) contains the number of coefficients to be kept at level i.

j, M and ALPHA define the strategy:

- At level $j+1$ (and coarser levels), everything is kept.
- For level i from 1 to j , the n_i largest coefficients are kept with $n_i = M / (j+2-i)^{\text{ALPHA}}$.

Typically ALPHA = 1.5 for compression and ALPHA = 3 for de-noising.

A default value for M is $M = \text{prod}(S(1,:))$, the length of the coarsest approximation coefficients, since the previous formula leads for $i = j+1$, to $n_{j+1} = M = \text{prod}(S(1,:))$.

Recommended values for M are from $\text{prod}(S(1,:))$ to $6 * \text{prod}(S(1,:))$.

`wdcbm2(C,S,ALPHA)` is equivalent to `wdcbm2(C,S,ALPHA,prod(S(1,:)))`.

Examples

```
% Load original image.
load detfingr;
nbc = size(map,1);

% Perform a wavelet decomposition of the image
% at level 3 using sym4.
wname = 'sym4'; lev = 3;
[c,s] = wavedec2(X,lev,wname);

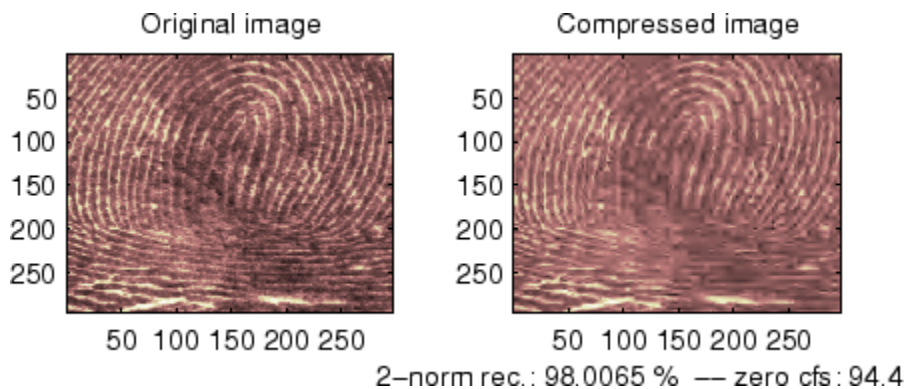
% Use wdcbm2 for selecting level dependent thresholds
% for image compression using the advised parameters.
alpha = 1.5; m = 2.7*prod(s(1,:));
[thr,nkeep] = wdcbm2(c,s,alpha,m)

thr =
    21.4814    46.8354    40.7907
    21.4814    46.8354    40.7907
    21.4814    46.8354    40.7907

nkeep =
         624         961        1765

% Use wdecmp for compressing the image using the above
% thresholds with hard thresholding.
[xd,cxd,sxd,perf0,perfl2] = ...
    wdecmp('lvd',c,s,wname,lev,thr,'h');

% Plot original and compressed images.
colormap(pink(nbc));
subplot(221), image(wcodemat(X,nbc)),
title('Original image')
subplot(222), image(wcodemat(xd,nbc)),
title('Compressed image')
xlab1 = ['2-norm rec.: ',num2str(perfl2)];
xlab2 = ['% -- zero cfs: ',num2str(perf0), '%'];
xlabel([xlab1 xlab2]);
```

References

Birgé, L.; P. Massart (1997). "From model selection to adaptive estimation," in D. Pollard (ed), *Festchrift for L. Le Cam*, Springer, pp. 55-88.

See Also

wdencmp | wpdencmp

Introduced before R2006a

wdecenergy

Multisignal 1-D decomposition energy distribution

Syntax

```
[E,PEC,PECFS] = wdecenergy(DEC)
[E,PEC,PECFS,IDXSORT,LONGS] = wdecenergy(DEC,'sort')
[E,PEC,PECFS] = wdecenergy(DEC,OPTSORT,IDXSIG)
[E,PEC,PECFS,IDXSORT,LONGS] = wdecenergy(DEC,OPTSORT,IDXSIG)
```

Description

`[E,PEC,PECFS] = wdecenergy(DEC)` computes the vector `E` that contains the energy (L2-Norm) of each decomposed signal, the matrix `PEC` that contains the percentage of energy for each wavelet component (approximation and details) of each signal, and the matrix `PECFS` that contains the percentage of energy for each coefficient.

- `E(i)` is the energy (L2-norm) of the `ith` signal.
- `PEC(i,1)` is the percentage of energy for the approximation of level `MAXLEV = DEC.level` of the `ith` signal.
- `PEC(i,j)`, `j=2,...,MAXLEV+1` is the percentage of energy for the detail of level `(MAXLEV+1-j)` of the `ith` signal.
- `PECFS(i,j)`, is the percentage of energy for `jth` coefficients of the `ith` signal.

`[E,PEC,PECFS,IDXSORT,LONGS] = wdecenergy(DEC,'sort')` returns `PECFS` sorted (by row) in ascending order and an index vector `IDXSORT`.

- Replacing 'sort' by 'ascend' returns the same result.
- Replacing 'sort' by 'descend' returns `PECFS` sorted in descending order.

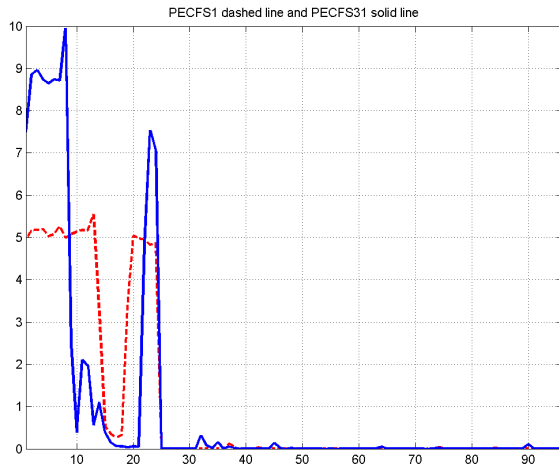
`LONGS` is a vector containing the lengths of each family of coefficients.

`[E,PEC,PECFS] = wdecenergy(DEC,OPTSORT,IDXSIG)` returns the values for the signals whose indices are given by the `IDXSIG` vector.

[E,PEC,PECFS,IDXSORT,LONGS] = wdecenergy(DEC,OPTSORT,IDXSIG) returns the values for the signals whose indices are given by the IDXSIG vector, the index vector IDXSORT, and LONGS, which is a vector containing the lengths of each family of coefficients. Valid values for OPTSORT are 'none', 'sort', 'ascend', 'descend'.

Examples

```
% Load original 1D-multisignal.
load thinker
% Perform a decomposition at level 2 using wavelet db2.
dec = mdwtdec('r',X,2,'db2');
% Compute the energy distribution.
[E,PEC,PECFS] = wdecenergy(dec);
% Display the total energy and the distribution of energy
% for each wavelet component (A2, D2, D1).
E31 = E(31)
perA2D2D1 = PEC(31,:);
% Compare the coefficient energy distribution
% for signal 1 and signal 31.
PECFS_1 = PECFS(1,:);
PECFS_31 = PECFS(31,:);
figure;
plot(PECFS_1,'--r','linewidth',2); hold on
plot(PECFS_31,'b','linewidth',2);
grid; set(gca,'Xlim',[1,size(PECFS,2)])
title('PECFS1 dashed line and PECFS31 solid line')
```



See Also

[mdwtdec](#) | [mdwtrec](#)

Introduced in R2012a

wden

Automatic 1-D denoising

Note wden is no longer recommended. Use `wdenoise` instead.

Syntax

```

XD = wden(X,TPTR,SORH,SCAL,N,wname)
XD = wden(C,L,___)
XD = wden(W,'modwtsqtwo log',SORH,'mln',N,wname)
[XD,CXD] = wden(___)
[XD,CXD,LXD] = wden(___)
[XD,CXD,LXD,THR] = wden(___)
[XD,CXD,THR] = wden(___)

```

Description

`XD = wden(X,TPTR,SORH,SCAL,N,wname)` returns a denoised version `XD` of the signal `X`. The function uses an `N`-level wavelet decomposition of `X` using the specified orthogonal or biorthogonal wavelet `wname` to obtain the wavelet coefficients. The thresholding selection rule `TPTR` is applied to the wavelet decomposition. `SORH` and `SCAL` define how the rule is applied.

`XD = wden(C,L,___)` returns a denoised version `XD` of the signal `X` using the same options as in the previous syntax, but obtained directly from the wavelet decomposition structure `[C,L]` of `X`. `[C,L]` is the output of `wavedec`.

`XD = wden(W,'modwtsqtwo log',SORH,'mln',N,wname)` returns the denoised signal `XD` obtained by operating on the maximal overlap discrete wavelet transform (MODWT) matrix `W`, where `W` is the output of `modwt`. You must use the same orthogonal wavelet in both `modwt` and `wden`.

`[XD,CXD] = wden(___)` returns the denoised wavelet coefficients. For discrete wavelet transform (DWT) denoising, `CXD` is a vector (see `wavedec`). For MODWT

denoising, CXD is a matrix with N+1 rows (see `modwt`). The number of columns of CXD is equal to the length of the input signal X.

`[XD, CXD, LXD] = wden(___)` returns the number of coefficients by level for DWT denoising. See `wavedec` for details. The LXD output is not supported for MODWT denoising. The additional output arguments `[CXD, LXD]` are the wavelet decomposition structure (see `wavedec` for more information) of the denoised signal XD.

`[XD, CXD, LXD, THR] = wden(___)` returns the denoising thresholds by level for DWT denoising.

`[XD, CXD, THR] = wden(___)` returns the denoising thresholds by level for MODWT denoising when you specify the `'modwtsqtwolog'` input argument.

Examples

Automatic 1-D Denoising Using Wavelets

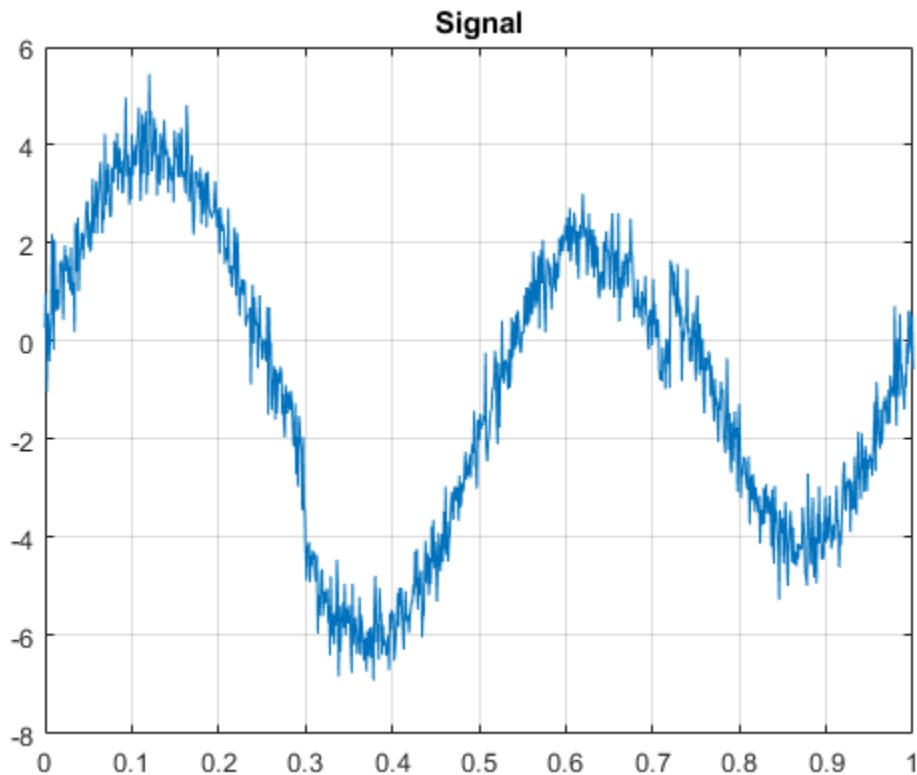
This example shows how to apply three different denoising techniques to a noisy signal. It compares the results with plots and the threshold values produced by each technique.

First, to ensure reproducibility of results, set a seed that will be used to generate the random noise.

```
rng('default')
```

Create a signal consisting of a 2 Hz sine wave with transients at 0.3 and 0.72 seconds. Add randomly generated noise to the signal and plot the result.

```
N = 1000;  
t = linspace(0,1,N);  
x = 4*sin(4*pi*t);  
x = x - sign(t-0.3) - sign(0.72-t);  
sig = x + 0.5*randn(size(t));  
plot(t,sig)  
title('Signal')  
grid on
```

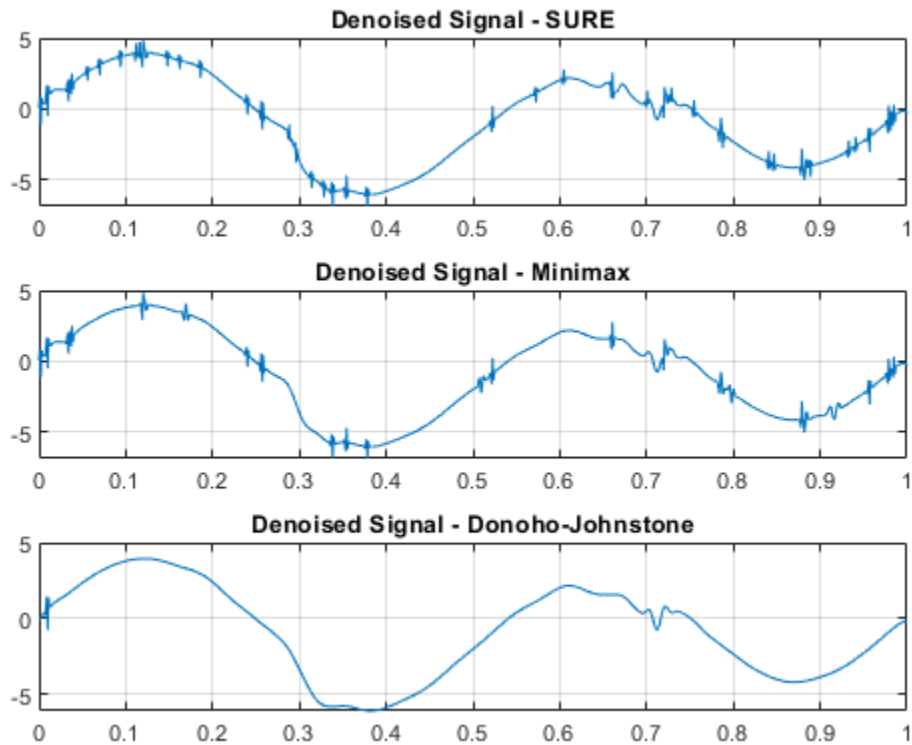


Using the `sym8` wavelet, perform a level 5 wavelet decomposition of the signal and denoise it by applying three different threshold selection rules to the wavelet coefficients: SURE, minimax, and Donoho and Johnstone's universal threshold with level-dependent estimation of the noise. In each case, apply hard thresholding.

```
lev = 5;
wname = 'sym8';
[dnSig1,c1,l1,threshold_SURE] = wden(sig,'rigrsure','h','mln',lev,wname);
[dnSig2,c2,l2,threshold_Minimax] = wden(sig,'minimaxi','h','mln',lev,wname);
[dnSig3,c3,l3,threshold_DJ] = wden(sig,'sqtwolog','h','mln',lev,wname);
```

Plot and compare the three denoised signals.

```
subplot(3,1,1)
plot(t,dnsig1)
title('Denoised Signal - SURE')
grid on
subplot(3,1,2)
plot(t,dnsig2)
title('Denoised Signal - Minimax')
grid on
subplot(3,1,3)
plot(t,dnsig3)
title('Denoised Signal - Donoho-Johnstone')
grid on
```



Compare the thresholds applied at each detail level for the three denoising methods.


```

threshold_SURE
threshold_SURE = 1x5
    0.8079    1.1448    1.3915    0.9816    0.8667

threshold_Minimax
threshold_Minimax = 1x5
    1.0522    1.0881    1.2237    1.3366    1.1985

threshold_DJ
threshold_DJ = 1x5
    1.7644    1.8247    2.0520    2.2413    2.0097

```

Compare DWT and MODWT Denoising of a Sinusoid with Two Jumps

This example denoises a signal using the DWT and MODWT. It compares the results with plots and the threshold values produced by each technique.

First, to ensure reproducibility of results, set a seed that will be used to generate random noise.

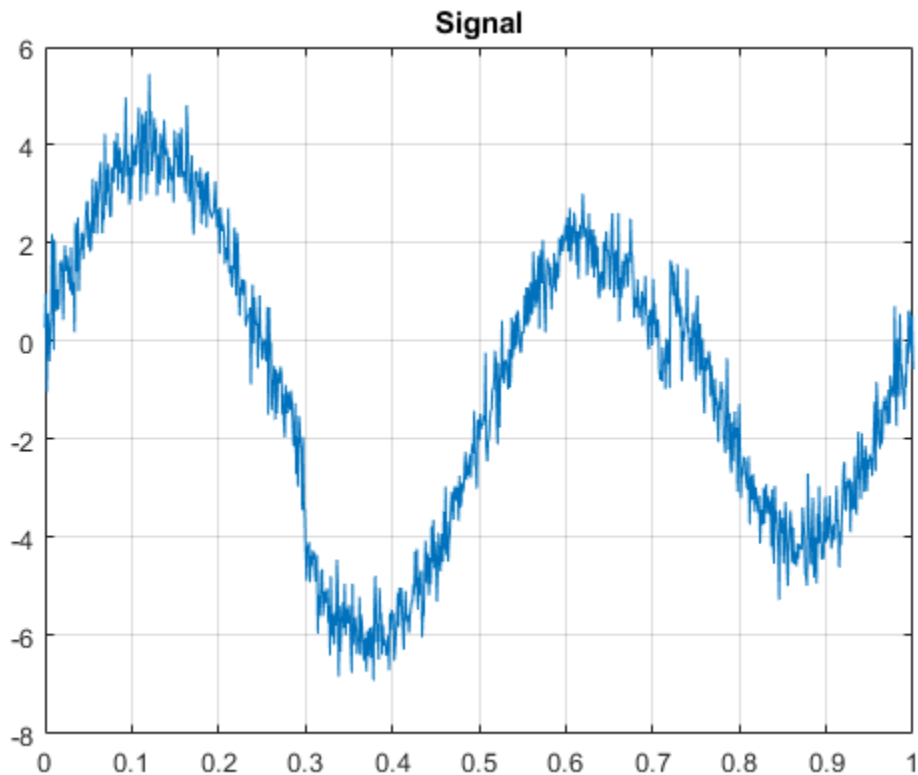
```
rng('default')
```

Create a signal consisting of a 2 Hz sine wave with transients at 0.3 and 0.72 seconds. Add randomly generated noise to the signal and plot the result.

```

N = 1000;
t = linspace(0,1,N);
x = 4*sin(4*pi*t);
x = x - sign(t-0.3) - sign(0.72-t);
sig = x + 0.5*randn(size(t));
plot(t,sig)
title('Signal')
grid on

```



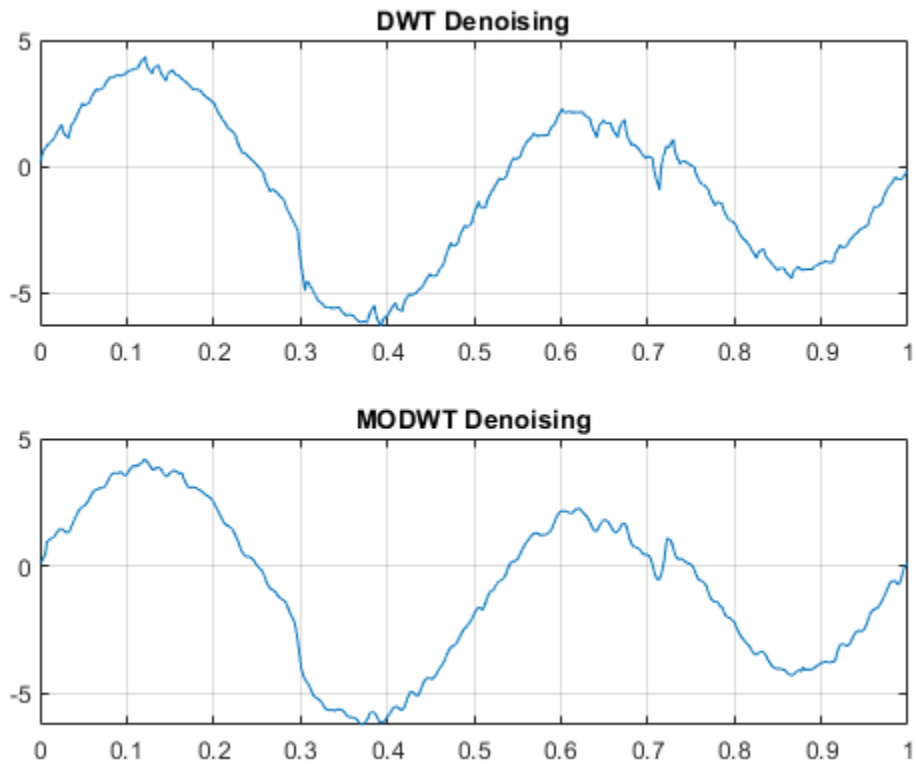
Using the `db2` wavelet, perform a level 3 wavelet decomposition of the signal and denoise it using Donoho and Johnstone's universal threshold with level-dependent estimation of the noise. Obtain denoised versions using DWT and MODWT, both with soft thresholding.

```
wname = 'db2';  
lev = 3;  
[xdDWT,c1,l1,threshold_DWT] = wden(sig,'sqrtwolog','s','mln',lev,wname);  
[xdMODWT,c2,threshold_MODWT] = wden(sig,'modwtsqrtwolog','s','mln',lev,wname);
```

Plot and compare the results.

```
subplot(2,1,1)  
plot(t,xdDWT)  
grid on
```

```
title('DWT Denoising')
subplot(2,1,2)
plot(t,xdMODWT)
grid on
title('MODWT Denoising')
```



Compare the thresholds applied in each case.

```
threshold_DWT
```

```
threshold_DWT = 1×3
```

```
    1.7783    1.6876    2.0434
```

```
threshold_MODWT
threshold_MODWT = 1×3
    1.2760    0.6405    0.3787
```

Compare DWT and MODWT Denoising of a Blocky Signal

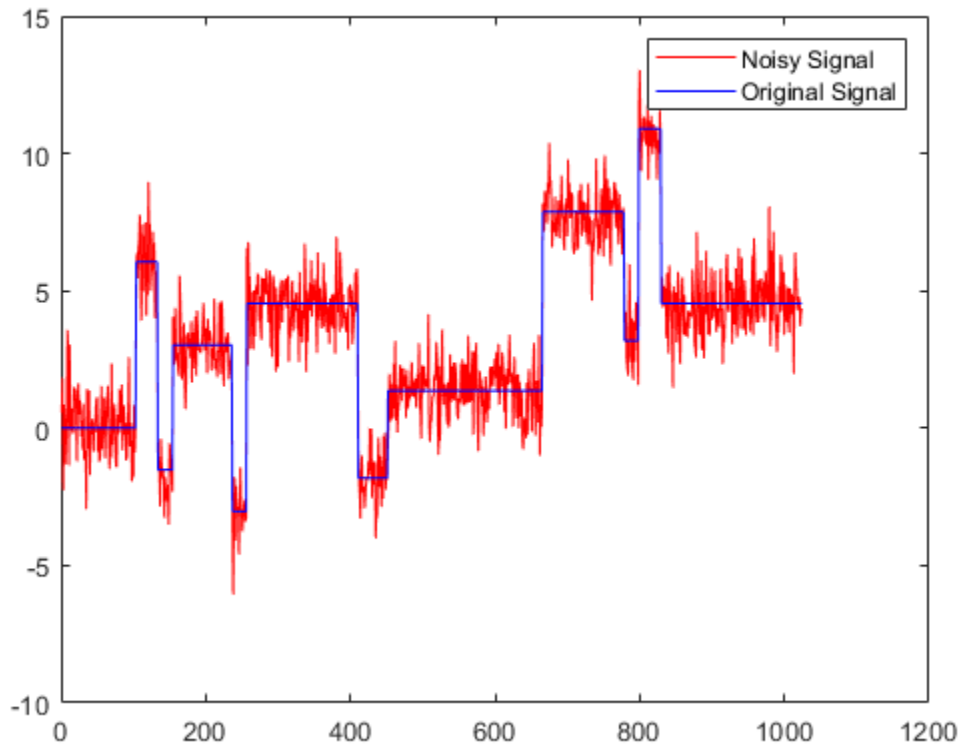
This example denoises a blocky signal using the Haar wavelet with DWT and MODWT denoising. It compares the results with plots and metrics for the original and denoised versions.

First, to ensure reproducibility of results, set a seed that will be used to generate random noise.

```
rng('default')
```

Generate a signal and a noisy version with the square root of the signal-to-noise ratio equal to 3. Plot and compare each.

```
[osig,nsig] = wnoise('blocks',10,3);
plot(nsig,'r')
hold on
plot(osig,'b')
legend('Noisy Signal','Original Signal')
```

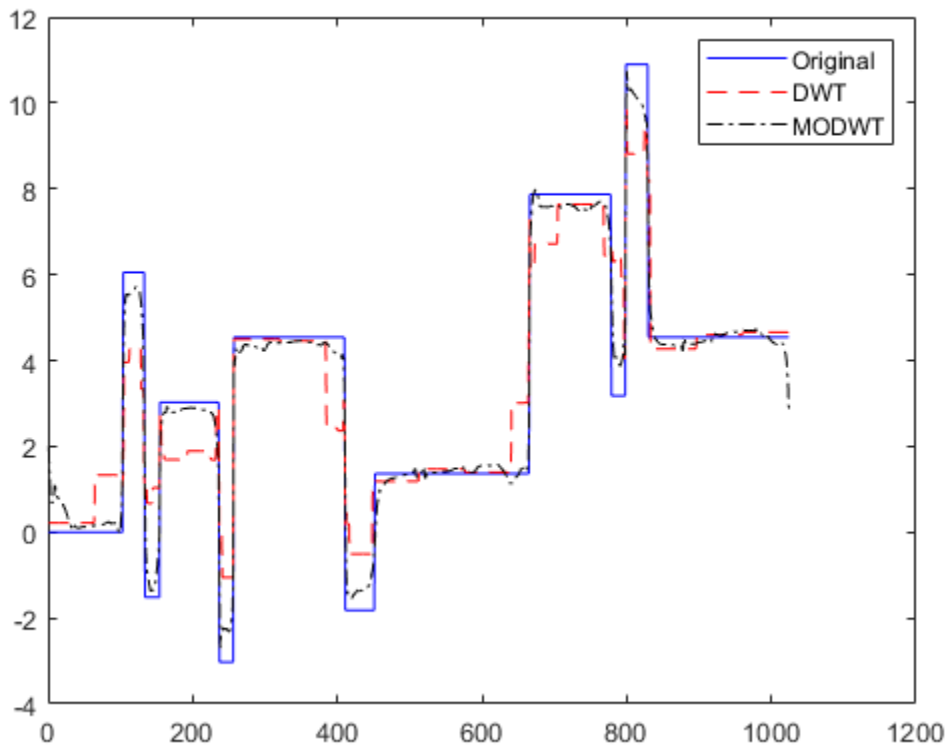


Using the Haar wavelet, perform a level 6 wavelet decomposition of the noisy signal and denoise it using Donoho and Johnstone's universal threshold with level-dependent estimation of the noise. Obtain denoised versions using DWT and MODWT, both with soft thresholding.

```
wname = 'haar';
lev = 6 ;
[xdDWT,c1,l1] = wden(nsig,'sqrtwolog','s','mln',lev,wname);
[xdMODWT,c2] = wden(nsig,'modwtsqrtwolog','s','mln',lev,wname);
```

Plot and compare the original, noise-free version of the signal with the two denoised versions.

```
figure
plot(osig,'b')
hold on
plot(xdDWT,'r--')
plot(xdMODWT,'k-.')
legend('Original','DWT','MODWT')
hold off
```



Calculate the L2 and L-infinity norms of the difference between the original signal and the two denoised versions.

```
L2norm_original_DWT = norm(abs(osig-xdDWT),2)
```

```
L2norm_original_DWT = 36.1194
```

```
L2norm_original_MODWT = norm(abs(osig-xdMODWT),2)
L2norm_original_MODWT = 14.5987
LInfinity_original_DWT = norm(abs(osig-xdDWT),Inf)
LInfinity_original_DWT = 4.7181
LInfinity_original_MODWT = norm(abs(osig-xdMODWT),Inf)
LInfinity_original_MODWT = 2.9655
```

Input Arguments

X — Input data

real-valued vector

Input data to denoise, specified as a real-valued vector.

Data Types: double

C — Wavelet expansion coefficients

real-valued vector

Wavelet expansion coefficients of the data to be denoised, specified as a real-valued vector. C is the output of `wavedec`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: double

L — Size of wavelet expansion coefficients

vector of positive integers

Size of wavelet expansion coefficients of the signal to be denoised, specified as a vector of positive integers. L is the output of `wavedec`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: double

W — Maximal overlap wavelet decomposition structure

real-valued matrix

Maximal overlap wavelet decomposition structure of the signal to denoise, specified as a real-valued matrix. W is the output of `modwt`. You must use the same orthogonal wavelet in both `modwt` and `wden`.

Data Types: `double`

TPTR — Threshold selection rule

character array

Threshold selection rule to apply to the wavelet decomposition structure of X :

- `'rigsure'` — Use the principle of Stein's Unbiased Risk.
- `'heursure'` — Use a heuristic variant of Stein's Unbiased Risk.
- `'sqtwo log'` — Use the universal threshold $\sqrt{2\ln(\text{length}(x))}$.
- `'minimaxi'` — Use minimax thresholding. (See `thselect` for more information.)

SORH — Type of thresholding

`'s'` | `'h'`

Type of thresholding to perform:

- `'s'` — Soft thresholding
- `'h'` — Hard thresholding

SCAL — Multiplicative threshold rescaling

`'one'` | `'sln'` | `'mln'`

Multiplicative threshold rescaling:

- `'one'` — No rescaling
- `'sln'` — Rescaling using a single estimation of level noise based on first-level coefficients
- `'mln'` — Rescaling using a level-dependent estimation of level noise

N — Level of wavelet decomposition

positive integer

Level of wavelet decomposition, specified as a positive integer. Use `wmaxlev` to ensure that the wavelet coefficients are free from boundary effects. If boundary effects are not a concern in your application, a good rule is to set N less than or equal to `fix(log2(length(X)))`.

wname — Name of wavelet

character array

Name of wavelet, specified as a character array, to use for denoising. For DWT denoising, the wavelet must be orthogonal or biorthogonal. For MODWT denoising, the wavelet must be orthogonal. Orthogonal and biorthogonal wavelets are designated as type 1 and type 2 wavelets, respectively, in the wavelet manager, `wavemngr`.

- Valid built-in orthogonal wavelet families begin with `haar`, `dbN`, `fkN`, `coifN`, or `symN`, where `N` is the number of vanishing moments for all families except `fk`. For `fk`, `N` is the number of filter coefficients.
- Valid biorthogonal wavelet families begin with `'biorNr.Nd'` or `'rbioNd.Nr'`, where `Nr` and `Nd` are the number of vanishing moments in the reconstruction (synthesis) and decomposition (analysis) wavelet.

Determine valid values for the vanishing moments by using `waveinfo` with the wavelet family short name. For example, enter `waveinfo('db')` or `waveinfo('bior')`. Use `wavemngr('type',wname)` to determine if a wavelet is orthogonal (returns 1) or biorthogonal (returns 2).

Output Arguments

XD — Denoised signal

real-valued vector

Denoised data, returned as a real-valued vector.

Data Types: `double`**CXD — Denoised wavelet coefficients**

real-valued vector or matrix

Denoised wavelet coefficients, returned as a real-valued vector or matrix. For DWT denoising, `CXD` is a vector (see `wavedec`). For MODWT denoising, `CXD` is a matrix with `N+1` rows (see `modwt`). The number of columns is equal to the length of the input signal `X`.

Data Types: `double`**LXD — Size of denoised wavelet coefficients**

vector of positive integers

Size of denoised wavelet coefficients by level for DWT denoising, returned as a vector of positive integers (see `wavedec`). The LXD output is not supported for MODWT denoising. [CXD, LXD] is the wavelet decomposition structure of the denoised signal XD.

Data Types: `double`

THR — Denoising thresholds

real-valued vector

Denoising thresholds by level, returned as a length N real-valued vector.

Data Types: `double`

Algorithms

The most general model for the noisy signal has the following form:

$$s(n) = f(n) + \sigma e(n),$$

where time n is equally spaced. In the simplest model, suppose that $e(n)$ is a Gaussian white noise $N(0,1)$, and the noise level σ is equal to 1. The denoising objective is to suppress the noise part of the signal s and to recover f .

The denoising procedure has three steps:

- 1** Decomposition — Choose a wavelet, and choose a level N. Compute the wavelet decomposition of the signal s at level N.
- 2** Detail coefficients thresholding — For each level from 1 to N, select a threshold and apply soft thresholding to the detail coefficients.
- 3** Reconstruction — Compute wavelet reconstruction based on the original approximation coefficients of level N and the modified detail coefficients of levels from 1 to N.

More details about threshold selection rules are in “Wavelet Denoising and Nonparametric Function Estimation” and in the help of the `thselect` function. Note that:

- The detail coefficients vector is the superposition of the coefficients of f and the coefficients of e . The decomposition of e leads to detail coefficients that are standard Gaussian white noises.

- Minimax and SURE threshold selection rules are more conservative and more convenient when small details of function f lie in the noise range. The two other rules remove the noise more efficiently. The option 'heursure' is a compromise.

In practice, the basic model cannot be used directly. To deal with model deviations, the remaining parameter `scal` must be specified. It corresponds to threshold rescaling methods.

- The option `scal = 'one'` corresponds to the basic model.
- The option `scal = 'sln'` handles threshold rescaling using a single estimation of level noise based on the first-level coefficients.

In general, you can ignore the noise level that must be estimated. The detail coefficients CD_1 (the finest scale) are essentially noise coefficients with standard deviation equal to σ . The median absolute deviation of the coefficients is a robust estimate of σ . The use of a robust estimate is crucial. If level 1 coefficients contain f details, these details are concentrated in a few coefficients to avoid signal end effects, which are pure artifacts due to computations on the edges.

- The option `scal = 'mln'` handles threshold rescaling using a level-dependent estimation of the level noise.

When you suspect a nonwhite noise e , thresholds must be rescaled by a level-dependent estimation of the level noise. The same kind of strategy is used by estimating σ_{lev} level by level. This estimation is implemented in the file `wnoisest`, which handles the wavelet decomposition structure of the original signal s directly.

References

- [1] Antoniadis, A., and G. Oppenheim, eds. *Wavelets and Statistics*, 103. Lecture Notes in Statistics. New York: Springer Verlag, 1995.
- [2] Donoho, D. L. "Progress in Wavelet Analysis and WVD: A Ten Minute Tour." *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.
- [3] Donoho, D. L., and Johnstone, I. M. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*, Vol. 81, pp. 425-455, 1994.
- [4] Donoho, D. L. "De-noising by Soft-Thresholding." *IEEE Transactions on Information Theory*, Vol. 42, Number 3, pp. 613-627, 1995.

[5] Donoho, D. L., I. M. Johnstone, G. Kerkyacharian, and D. Picard. "Wavelet Shrinkage: Asymptopia?" *Journal of the Royal Statistical Society, series B*. Vol. 57, Number 2, pp. 301-369, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

Functions

`thselect` | `wavedec` | `wavemngr` | `wdencmp` | `wdenoise` | `wfilters` | `wthresh`

Apps

Wavelet Signal Denoiser

Introduced before R2006a

wdencmp

Denoising or compression

Syntax

```
[XC,CXC,LXC,PERF0,PERFL2] = wdencmp('gbl',X,wname,N,THR,SORH,
KEEPAPP)
[ ___ ] = wdencmp('gbl',C,L,wname,N,THR,SORH,KEEPAPP)
[ ___ ] = wdencmp('lvl',X,wname,N,THR,SORH)
[ ___ ] = wdencmp('lvl',C,L,wname,N,THR,SORH)
```

Description

[XC,CXC,LXC,PERF0,PERFL2] = wdencmp('gbl',X,wname,N,THR,SORH,KEEPAPP) returns a denoised or compressed version XC of the input data X obtained by wavelet coefficients thresholding using the global positive threshold THR. X is a real-valued vector or matrix. [CXC,LXC] is the N-level wavelet decomposition structure of XC (see wavedec or wavedec2 for more information). PERF0 and PERFL2 are the L^2 -norm recovery and compression scores in percentages, respectively. If KEEPAPP = 1, the approximation coefficients are kept. If KEEPAPP = 0, the approximation coefficients can be thresholded.

[___] = wdencmp('gbl',C,L,wname,N,THR,SORH,KEEPAPP) uses the wavelet decomposition structure [C,L] of the data to be denoised or compressed.

[___] = wdencmp('lvl',X,wname,N,THR,SORH) uses the level-dependent thresholds THR. The approximation coefficients are kept.

[___] = wdencmp('lvl',C,L,wname,N,THR,SORH) uses the wavelet decomposition structure [C,L].

Examples

Denoise 1-D Signal Using Default Global Threshold

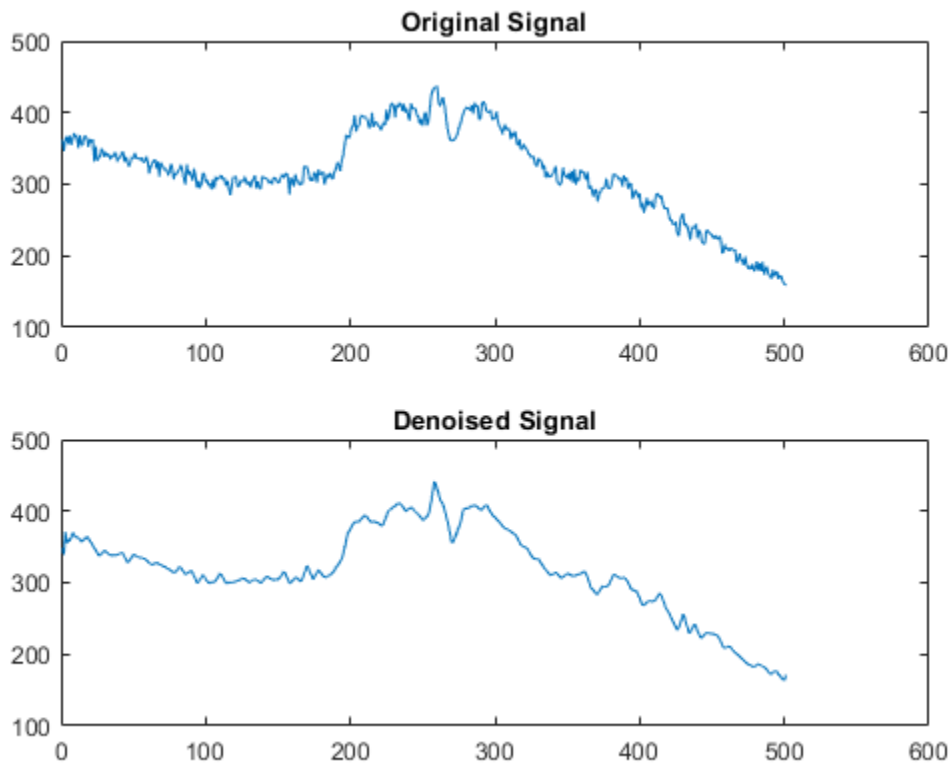
Denoise 1-D electricity consumption data using the Donoho-Johnstone global threshold.

Load the signal and select a segment for denoising.

```
load leleccum; indx = 2600:3100;  
x = leleccum(indx);
```

Use `ddencmp` to determine the default global threshold and denoise the signal. Plot the original and denoised signals.

```
[thr,sorh,keepapp] = ddencmp('den','wv',x);  
xd = wdencomp('gbl',x,'db3',2,thr,sorh,keepapp);  
subplot(211)  
plot(x); title('Original Signal');  
subplot(212)  
plot(xd); title('Denoised Signal');
```



Denoise Image Using Default Global Threshold

Denoise an image in additive white Gaussian noise using the Donoho-Johnstone universal threshold.

Load an image and add white Gaussian noise.

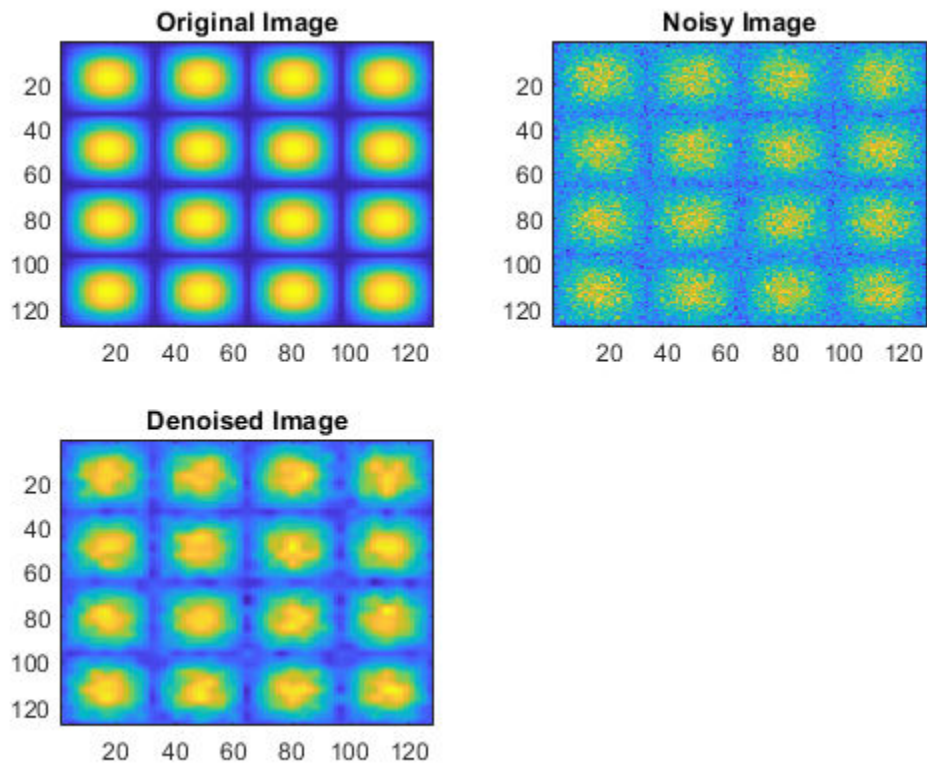
```
load sinsin  
Y = X+18*randn(size(X));
```

Use `ddencomp` to obtain the threshold.

```
[thr,sorh,keepapp] = ddencmp('den','wv',Y);
```

Denoise the image. Use the order 4 Symlet and a two-level wavelet decomposition. Plot the original image, the noisy image, and the denoised result.

```
xd = wdencomp('gbl',Y,'sym4',2,thr,sorh,keepapp);  
subplot(2,2,1)  
imagesc(X)  
title('Original Image')  
subplot(2,2,2)  
imagesc(Y)  
title('Noisy Image')  
subplot(2,2,3)  
imagesc(xd)  
title('Denoised Image')
```

Input Arguments

X – Input data

real-valued vector | real-valued matrix

Input data to denoise or compress, specified by a real-valued vector or matrix.

Data Types: double

C – Wavelet expansion coefficients

real-valued vector

Wavelet expansion coefficients of the data to be compressed or denoised, specified as a real-valued vector. If the data is one-dimensional, `C` is the output of `wavedec`. If the data is two-dimensional, `C` is the output of `wavedec2`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: `double`

L — Size of wavelet expansion coefficients

vector of positive integers | matrix of positive integers

Size of wavelet expansion coefficients of the signal or image to be compressed or denoised, specified as a vector or matrix of positive integers.

For signals, `L` is the output of `wavedec`. For images, `L` is the output of `wavedec2`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: `double`

wname — Name of wavelet

character vector | string scalar

Name of wavelet, specified as a character vector or string scalar, to use for denoising or compression. See `wavemngr` for more information. `wdencmp` uses `wname` to generate the `N`-level wavelet decomposition of `X`.

N — Level of wavelet decomposition

positive integer

Level of wavelet decomposition, specified as a positive integer.

THR — Threshold

scalar | real-valued vector | real-valued matrix

Threshold to apply to the wavelet coefficients, specified as a scalar, real-valued vector, or real-valued matrix.

- For the case `'gbl'`, `THR` is a scalar.
- For the one-dimensional case and `'lvd'` option, `THR` is a length `N` real-valued vector containing the level-dependent thresholds.
- For the two-dimensional case and `'lvd'` option, `THR` is a 3-by-`N` matrix containing the level-dependent thresholds in the three orientations: horizontal, diagonal, and vertical.

Data Types: double

SORH — Type of thresholding

's' | 'h'

Type of thresholding to perform:

- 's' — Soft thresholding
- 'h' — Hard thresholding

See `wthresh` for more information.

KEEPAPP — Threshold approximation setting

0 | 1

Threshold approximation setting, specified as either 0 or 1. If `KEEPAPP = 1`, the approximation coefficients cannot be thresholded. If `KEEPAPP = 0`, the approximation coefficients can be thresholded.

Data Types: double

Output Arguments

XC — Denoised or compressed data

real-valued vector | real-valued matrix

Denoised or compressed data, returned as a real-valued vector or matrix. `XC` and `X` have the same dimensions.

CXC — Wavelet expansion coefficients

real-valued vector

Wavelet expansion coefficients of the denoised or compressed data `XC`, returned as a real-valued vector. `LXC` contains the number of coefficients by level.

LXC — Size of wavelet expansion coefficients

vector of positive integers | matrix of positive integers

Size of wavelet expansion coefficients of the denoised or compressed data `XC`, specified as a vector or matrix of positive integers. If the data is one-dimensional, `LXC` is a vector of positive integers (see `wavedec` for more information). If the data is two-dimensional, `LXC` is a matrix of positive integers (see `wavedec2` for more information).

PERF0 — Compression score

scalar

Compression score, returned as a real number. PERF0 is the percentage of thresholded coefficients that are equal to 0.

PERFL2 — L^2 energy recovery

scalar

$PERFL2 = 100 * (\text{vector-norm of } CXC / \text{vector-norm of } C)^2$ if $[C, L]$ denotes the wavelet decomposition structure of X .

If X is a one-dimensional signal and ' $wname$ ' an orthogonal wavelet, PERFL2 is reduced to

$$\frac{100\|XC\|^2}{\|X\|^2}$$

Algorithms

The denoising and compression procedures contain three steps:

- 1 Decomposition.
- 2 Thresholding.
- 3 Reconstruction.

The two procedures differ in Step 2. In compression, for each level in the wavelet decomposition, a threshold is selected and hard thresholding is applied to the detail coefficients.

References

- [1] DeVore, R. A., B. Jawerth, and B. J. Lucier. "Image Compression Through Wavelet Transform Coding." *IEEE Transactions on Information Theory*. Vol. 38, Number 2, 1992, pp. 719-746.
- [2] Donoho, D. L. "Progress in Wavelet Analysis and WVD: A Ten Minute Tour." *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.

- [3] Donoho, D. L., and I. M. Johnstone. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*. Vol. 81, pp. 425-455, 1994.
- [4] Donoho, D. L., I. M. Johnstone, G. Kerkycharian, and D. Picard. "Wavelet Shrinkage: Asymptopia?" *Journal of the Royal Statistical Society, series B*, Vol. 57, No. 2, pp. 301-369, 1995.
- [5] Donoho, D. L., and I. M. Johnstone. "Ideal denoising in an orthonormal basis chosen from a library of bases." *C. R. Acad. Sci. Paris, Ser. I*, Vol. 319, pp. 1317-1322, 1994.
- [6] Donoho, D. L. "De-noising by Soft-Thresholding." *IEEE Transactions on Information Theory*. Vol. 42, Number 3, pp. 613-627, 1995.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- Variable-size data support must be enabled.

See Also

Functions

ddencmp | wavedec | wavedec2 | wbmphen | wcompress | wdcbm2 | wdenoise |
wpdencmp | wthresh

Apps

Wavelet Signal Denoiser

Introduced before R2006a

wdenoise

Wavelet signal denoising

Syntax

```
XDEN = wdenoise(X)
```

```
XDEN = wdenoise(X,LEVEL)
```

```
XDEN = wdenoise( ____,Name,Value)
```

```
[XDEN,DENOISEDDFS] = wdenoise( ____,Name,Value)
```

```
[XDEN,DENOISEDDFS,ORIGDFS] = wdenoise( ____,Name,Value)
```

Description

`XDEN = wdenoise(X)` denoises the data in `X` using an empirical Bayesian method with a Cauchy prior. By default, the `sym4` wavelet is used with a posterior median threshold rule. Denoising is down to the minimum of `floor(log2N)` and `wmaxlev(N, 'sym4')` where `N` is the number of samples in the data. (For more information, see `wmaxlev`.) `X` is a real-valued vector, matrix, or timetable.

- If `X` is a matrix, `wdenoise` denoises each column of `X`.
- If `X` is a timetable, `wdenoise` must contain real-valued vectors in separate variables, or one real-valued matrix of data.
- `X` is assumed to be uniformly sampled.
- If `X` is a timetable and the timestamps are not linearly spaced, `wdenoise` issues a warning.

`XDEN = wdenoise(X,LEVEL)` denoises `X` down to `LEVEL`. `LEVEL` is a positive integer less than or equal to `floor(log2N)` where `N` is the number of samples in the data. If unspecified, `LEVEL` defaults to the minimum of `floor(log2N)` and `wmaxlev(N, 'sym4')`.

`XDEN = wdenoise(____,Name,Value)` specifies options using name-value pair arguments in addition to any of the input arguments in previous syntaxes.

[XDEN,DENOISEDCFS] = wdenoise(____) returns the denoised wavelet and scaling coefficients in the cell array DENOISEDCFS. The elements of DENOISEDCFS are in order of decreasing resolution. The final element of DENOISEDCFS contains the approximation (scaling) coefficients.

[XDEN,DENOISEDCFS,ORIGCFS] = wdenoise(____) returns the original wavelet and scaling coefficients in the cell array ORIGCFS. The elements of ORIGCFS are in order of decreasing resolution. The final element of ORIGCFS contains the approximation (scaling) coefficients.

Examples

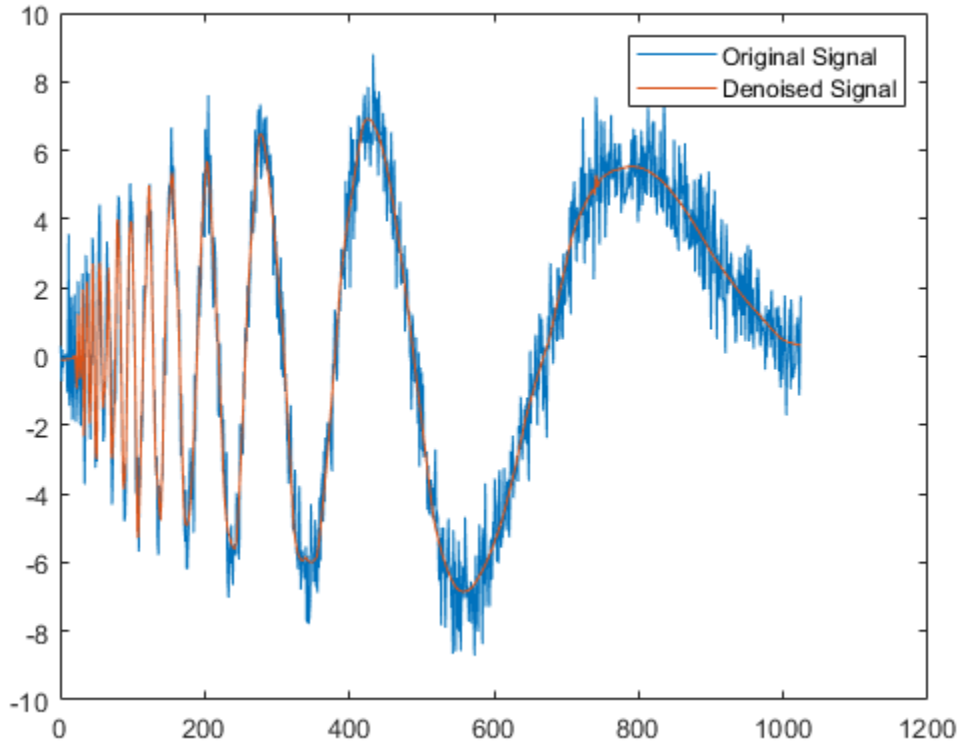
Denoise A Signal Using Default Values

Obtain the denoised version of a noisy signal using default values.

```
load noisdopp
xden = wdenoise(noisdopp);
```

Plot the original and denoised signals.

```
plot([noisdopp' xden'])
legend('Original Signal','Denoised Signal')
```



Denoise a Timetable Using Block Thresholding

Denoise a timetable of noisy data down to level 5 using block thresholding.

Load a noisy dataset.

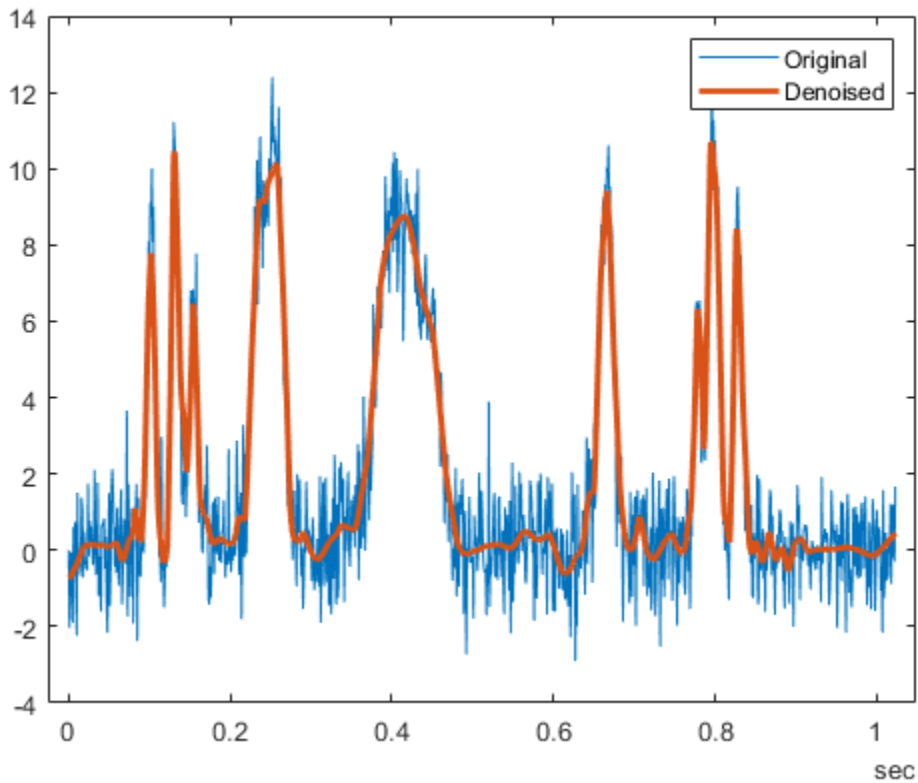
```
load wnoisydata
```

Denoise the data down to level 5 using block thresholding by setting the name-value pair 'DenoisingMethod', 'BlockJS'.

```
xden = wdenoise(wnoisydata,5,'DenoisingMethod','BlockJS');
```


Plot the original data and the denoised data.

```
h1 = plot(wnoisydata.t,[wnoisydata.noisydata(:,1) xden.noisydata(:,1)]);  
h1(2).LineWidth = 2;  
legend('Original','Denoised')
```

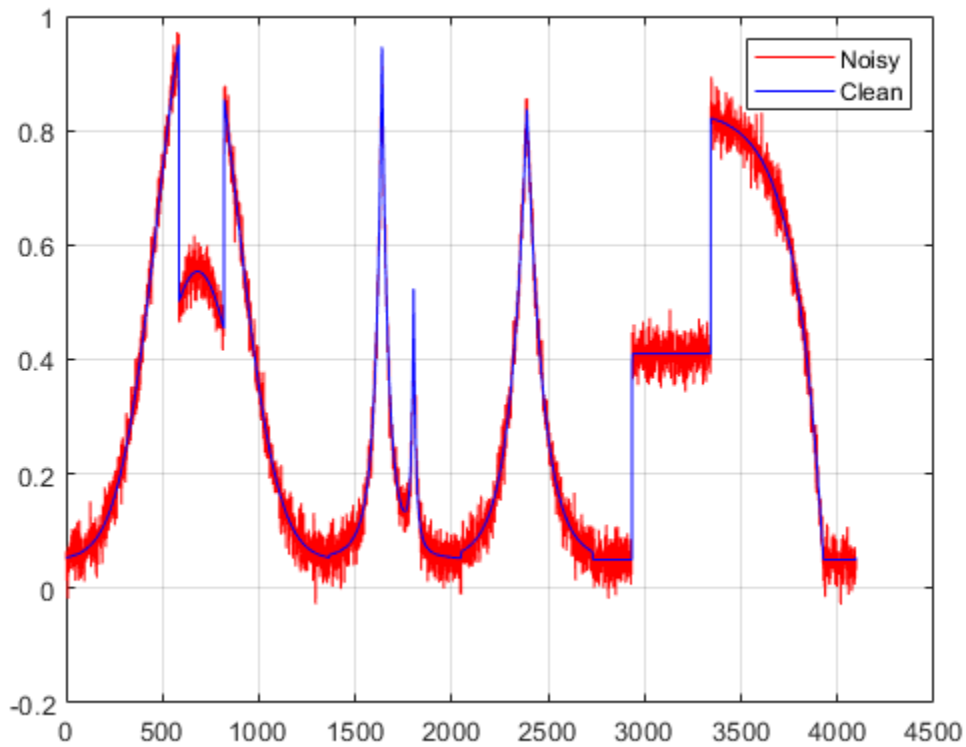


Compare Denoised Signals

Denoise a signal in different ways and compare results.

Load a datafile that contains clean and noisy versions of a signal. Plot the signals.

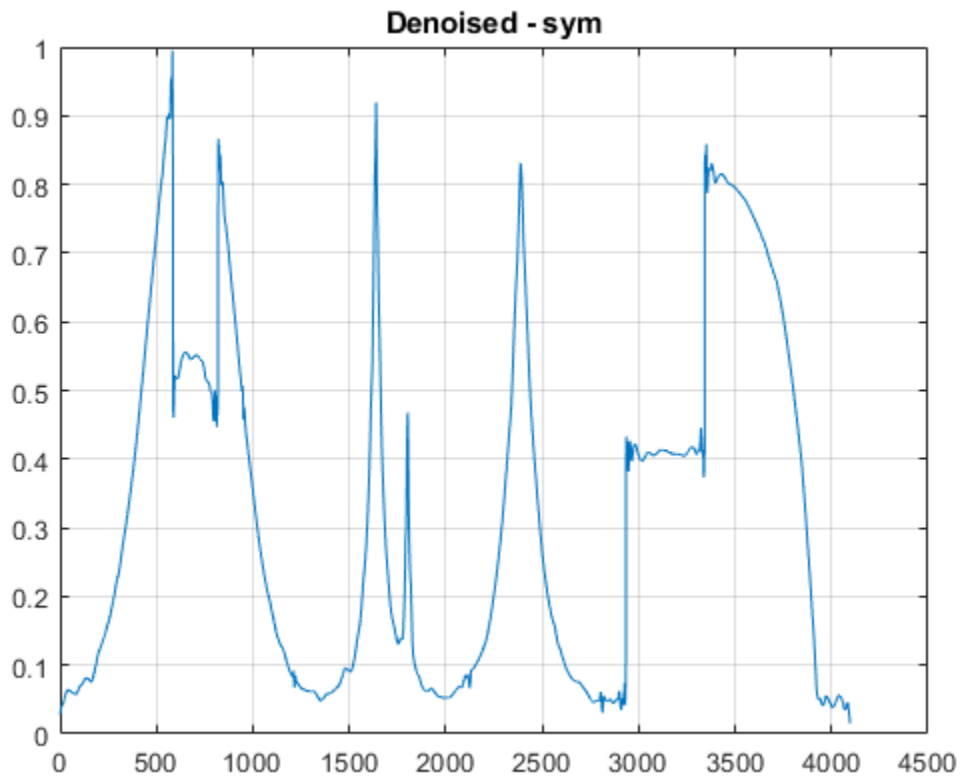
```
load fdata.mat
plot(fNoisy,'r-')
hold on
plot(fClean,'b-')
grid on
legend('Noisy','Clean');
```



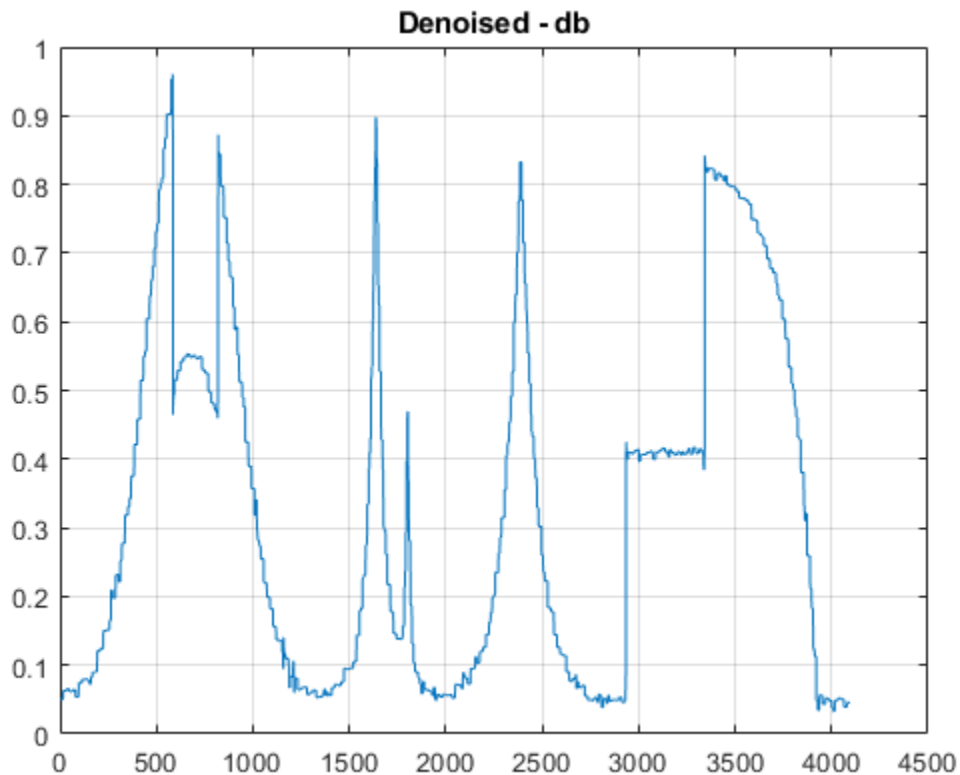
Denoise the signal using the `sym4` and `db1` wavelets, with a nine-level wavelet decomposition. Plot the results.

```
cleansym = wdenoise(fNoisy,9,'Wavelet','sym4');
cleandb = wdenoise(fNoisy,9,'Wavelet','db1');
figure
plot(cleansym)
```

```
title('Denoised - sym')  
grid on
```



```
figure  
plot(cleandb)  
title('Denoised - db')  
grid on
```



Compute the SNR of each denoised signal. Confirm that using the `sym4` wavelet produces a better result.

```
snrsym = -20*log10(norm(abs(fClean-cleansym))/norm(fClean))
```

```
snrsym = 35.6654
```

```
snrdb = -20*log10(norm(abs(fClean-cleandb))/norm(fClean))
```

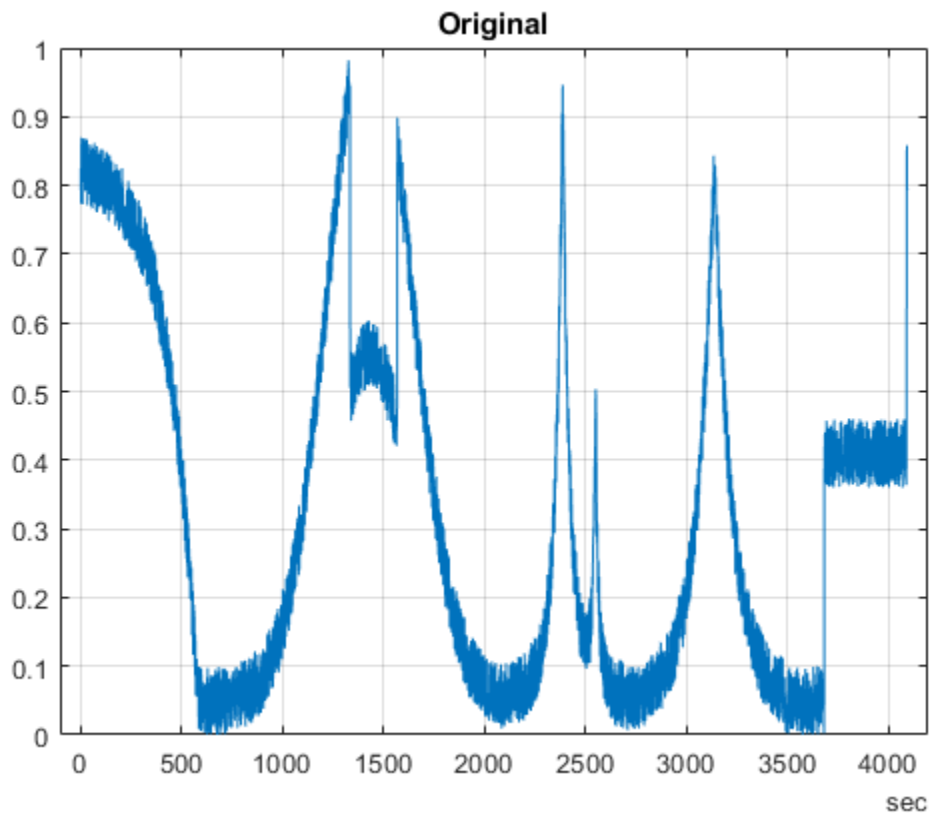
```
snrdb = 32.2672
```

Load in a file which contains noisy data of 100 time series. Every time series is a noisy version of `fClean`. Denoise the time series twice, estimating the noise variance differently in each case.

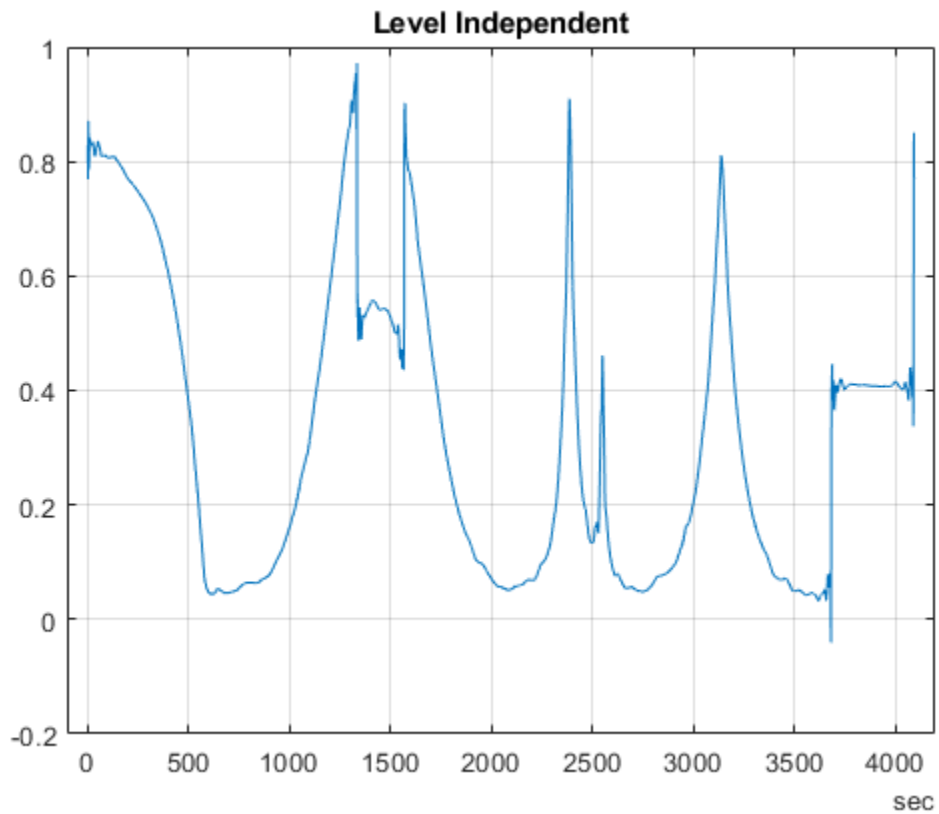
```
load fdataTS.mat
cleanTSld = wdenoise(fdataTS,9,'NoiseEstimate','LevelDependent');
cleanTSli = wdenoise(fdataTS,9,'NoiseEstimate','LevelIndependent');
```

Compare one of the noisy time series with its two denoised versions.

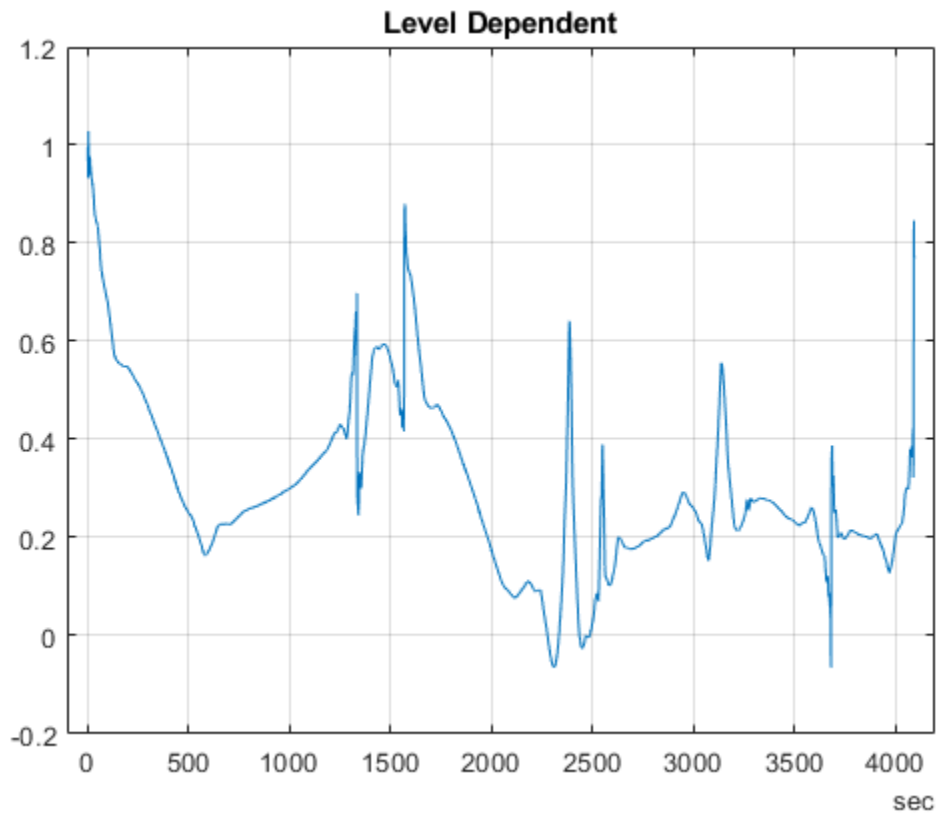
```
figure
plot(fdataTS.Time,fdataTS.fTS15)
title('Original')
grid on
```



```
figure
plot(cleanTSli.Time,cleanTSli.fTS15)
title('Level Independent')
grid on
```



```
figure  
plot(cleanTSl.d.Time,cleanTSl.d.fTS15)  
title('Level Dependent')  
grid on
```



Input Arguments

X — Input data

vector | matrix | timetable

Input data, specified as a matrix, vector, or timetable of real values. If X is a vector, it must have at least two samples. If X is a matrix or timetable, it must have at least two rows.

Data Types: double

LEVEL — Level of wavelet decomposition

positive integer

Level of wavelet decomposition, specified as a positive integer. LEVEL is a positive integer less than or equal to $\text{floor}(\log_2 N)$ where N is the number of samples in the data.

- If unspecified, LEVEL defaults to the minimum of $\text{floor}(\log_2 N)$ and $\text{wmaxlev}(N, 'sym4')$.
- For James-Stein block thresholding, 'BlockJS', there must be $\text{floor}(\log_2 N)$ coefficients at the coarsest resolution level, LEVEL.

Data Types: double

Name-Value Pair Arguments

Specify optional comma-separated pairs of Name, Value arguments. Name is the argument name and Value is the corresponding value. Name must appear inside quotes. You can specify several name and value pair arguments in any order as Name1, Value1, ..., NameN, ValueN.

Example: 'Wavelet', 'db6', 'DenoisingMethod', 'Bayes' denoises using the Daubechies db6 wavelet and the empirical Bayesian method.

Wavelet — Name of wavelet

'sym4' (default) | character array

Name of wavelet, specified as a character array, to use for denoising. The wavelet must be orthogonal or biorthogonal. Orthogonal and biorthogonal wavelets are designated as type 1 and type 2 wavelets respectively in the wavelet manager, `wavemngr`.

- Valid built-in orthogonal wavelet families begin with `haar`, `dbN`, `fkN`, `coifN`, or `symN` where N is the number of vanishing moments for all families except `fk`. For `fk`, N is the number of filter coefficients.
- Valid biorthogonal wavelet families begin with `'biorNr.Nd'` or `'rbioNd.Nr'`, where Nr and Nd are the number of vanishing moments in the reconstruction (synthesis) and decomposition (analysis) wavelet.

Determine valid values for the vanishing moments by using `waveinfo` with the wavelet family short name. For example, enter `waveinfo('db')` or `waveinfo('bior')`. Use `wavemngr('type', WNAME)` to determine if a wavelet is orthogonal (returns 1) or biorthogonal (returns 2).

DenoisingMethod — Denoising method

'Bayes' (default) | 'BlockJS' | 'FDR' | 'Minimax' | 'SURE' | 'UniversalThreshold'

Denoising method, specified as a character array, used to determine the denoising thresholds for the data X .

- **Bayes** - Empirical Bayes

This method uses a threshold rule based on assuming measurements have independent prior distributions given by a mixture model. Because measurements are used to estimate the weight in the mixture model, the method tends to work better with more samples. By default, the posterior median rule is used to measure risk [8].

- **BlockJS** - Block James-Stein

This method is based on determining an `optimal block size and threshold. The resulting block thresholding estimator yields simultaneously optimal global and local adaptivity [3].

- **FDR** - False Discovery Rate

This method uses a threshold rule based on controlling the expected ratio of false positive detections to all positive detections. The FDR method works best with sparse data. Choosing a ratio, or Q -value, less than $1/2$ yields an asymptotically minimax estimator [1].

- **Minimax** - Minimax Estimation

This method uses a fixed threshold chosen to yield minimax performance for mean square error against an ideal procedure. The minimax principle is used in statistics to design estimators. See `thselect` for more information.

- **SURE** - Stein's Unbiased Risk Estimate

This method uses a threshold selection rule based on Stein's Unbiased Estimate of Risk (quadratic loss function). One gets an estimate of the risk for a particular threshold value (t). Minimizing the risks in (t) gives a selection of the threshold value.

- **UniversalThreshold** - Universal Threshold $\sqrt{2\ln(\text{length}(x))}$.

This method uses a fixed-form threshold yielding minimax performance multiplied by a small factor proportional to $\log(\text{length}(X))$.

Note For 'FDR', there is an optional argument for the Q -value, which is the proportion of false positives. Q is a real-valued scalar between 0 and $1/2$, $0 < Q \leq 1/2$. To specify

'FDR' with a Q -value, use a cell array where the second element is the Q -value. For example, 'DenoisingMethod', {'FDR', 0.01}. If unspecified, Q defaults to 0.05.

ThresholdRule — Threshold rule

character array

Threshold rule, specified as a character array, to use to shrink the wavelet coefficients. 'ThresholdRule' is valid for all denoising methods, but the valid options and defaults depend on the denoising method. Rules possible for different denoising methods are specified as follows:

- 'BlockJS': The only supported option is 'James-Stein'. You do not need to specify 'ThresholdRule' for 'BlockJS'.
- 'SURE', 'Minimax', 'UniversalThreshold': Valid options are 'Soft' or 'Hard'. The default is 'Soft'.
- 'Bayes': Valid options are 'Median', 'Mean', 'Soft', or 'Hard'. The default is 'Median'.
- 'FDR': The only supported option is 'Hard'. You do not need to define 'ThresholdRule' for 'FDR'.

NoiseEstimate — Method of estimating variance of noise

'LevelIndependent' (default) | 'LevelDependent'

Method of estimating variance of noise in the data, specified as a character array. Valid options are 'LevelIndependent' and 'LevelDependent'.

- 'LevelIndependent' estimates the variance of the noise based on the finest-scale (highest-resolution) wavelet coefficients.
- 'LevelDependent' estimates the variance of the noise based on the wavelet coefficients at each resolution level.
- For the block James-Stein estimator ('BlockJS'), 'LevelIndependent' is the only supported option.

Output Arguments

XDEN — Denoised data

vector | matrix | timetable

Denoised vector, matrix, or timetable version of X . For timetable input, $XDEN$ has the same variable names and timestamps as the original timetable.

Data Types: `double`

DENOISED_CFS — Denoised wavelet and scaling coefficients

cell array

Denoised wavelet and scaling coefficients of the denoised data $XDEN$, returned in a cell array. The elements of `DENOISED_CFS` are in order of decreasing resolution. The final element of `DENOISED_CFS` contains the approximation (scaling) coefficients.

Data Types: `double`

ORIG_CFS — Original wavelet and scaling coefficients

cell array

Original wavelet and scaling coefficients of the data X , returned in a cell array. The elements of `ORIG_CFS` are in order of decreasing resolution. The final element of `ORIG_CFS` contains the approximation (scaling) coefficients.

Data Types: `double`

Algorithms

The most general model for the noisy signal has the following form:

$$s(n) = f(n) + \sigma e(n),$$

where time n is equally spaced. In the simplest model, suppose that $e(n)$ is a Gaussian white noise $N(0,1)$, and the noise level σ is equal to 1. The denoising objective is to suppress the noise part of the signal s and to recover f .

The denoising procedure has three steps:

- 1** Decomposition — Choose a wavelet, and choose a level N . Compute the wavelet decomposition of the signal s at level N .
- 2** Detail coefficients thresholding — For each level from 1 to N , select a threshold and apply soft thresholding to the detail coefficients.
- 3** Reconstruction — Compute wavelet reconstruction based on the original approximation coefficients of level N and the modified detail coefficients of levels from 1 to N .

More details about threshold selection rules are in “Wavelet Denoising and Nonparametric Function Estimation” and in the help of the `thselect` function.

References

- [1] Abramovich, F., Y. Benjamini, D. L. Donoho, and I. M. Johnstone. “Adapting to Unknown Sparsity by Controlling the False Discovery Rate.” *Annals of Statistics*, Vol. 34, Number 2, pp. 584-653, 2006.
- [2] Antoniadis, A., and G. Oppenheim, eds. *Wavelets and Statistics*. Lecture Notes in Statistics. New York: Springer Verlag, 1995.
- [3] Cai, T. T. “On Block Thresholding in Wavelet Regression: Adaptivity, Block size, and Threshold Level.” *Statistica Sinica*, Vol. 12, pp. 1241-1273, 2002.
- [4] Donoho, D. L. “Progress in Wavelet Analysis and WVD: A Ten Minute Tour.” *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.
- [5] Donoho, D. L., I. M. Johnstone. “Ideal Spatial Adaptation by Wavelet Shrinkage.” *Biometrika*, Vol. 81, pp. 425-455, 1994.
- [6] Donoho, D. L. “De-noising by Soft-Thresholding.” *IEEE Transactions on Information Theory*, Vol. 42, Number 3, pp. 613-627, 1995.
- [7] Donoho, D. L., I. M. Johnstone, G. Kerkycharian, and D. Picard. “Wavelet Shrinkage: Asymptopia?” *Journal of the Royal Statistical Society, series B*, Vol. 57, No. 2, pp. 301-369, 1995.
- [8] Johnstone, I. M., and B. W. Silverman. “Needles and Straw in Haystacks: Empirical Bayes Estimates of Possibly Sparse Sequences.” *Annals of Statistics*, Vol. 32, Number 4, pp. 1594-1649, 2004.

See Also

Wavelet Signal Denoiser | `thselect` | `waveinfo` | `wavemngr` | `wdenoise2` | `wmaxlev`

Introduced in R2017b

wdenoise2

Wavelet image denoising

Syntax

```

IMDEN = wdenoise2(IM)
IMDEN = wdenoise2(IM,LEVEL)
[IMDEN,DENOISED_CFS] = wdenoise2( ___ )

[IMDEN,DENOISED_CFS,ORIG_CFS] = wdenoise2( ___ )
[IMDEN,DENOISED_CFS,ORIG_CFS,S] = wdenoise2( ___ )
[IMDEN,DENOISED_CFS,ORIG_CFS,S,SHIFTS] = wdenoise2( ___ )

[ ___ ] = wdenoise2( ___ ,Name,Value)

wdenoise2( ___ )

```

Description

`IMDEN = wdenoise2(IM)` denoises the grayscale or RGB image `IM` using an empirical Bayesian method. The `bior4.4` wavelet is used with a posterior median threshold rule. Denoising is down to the minimum of `floor(log2([M N]))` and `wmaxlev([M N], 'bior4.4')` where `M` and `N` are the row and column sizes of the image. `IMDEN` is the denoised version of `IM`.

For RGB images, by default, `wdenoise2` projects the image onto its principle component analysis (PCA) color space before denoising. To denoise an RGB image in the original color space, use the `ColorSpace` name-value pair.

`IMDEN = wdenoise2(IM,LEVEL)` denoises the image `IM` down to resolution level `LEVEL`. `LEVEL` is a positive integer less than or equal to `floor(log2(min([M N])))` where `M` and `N` are the row and column sizes of the image. If unspecified, `LEVEL` defaults to the minimum `floor(log2(min([M N])))` and `wmaxlev([M N],wname)` where `wname` is the wavelet used (`'bior4.4'` by default).

[IMDEN,DENOISEDCFS] = `wdenoise2`(___) returns the scaling and denoised wavelet coefficients in DENOISEDCFS using any of the preceding syntaxes.

[IMDEN,DENOISEDCFS,ORIGCFS] = `wdenoise2`(___) returns the scaling and wavelet coefficients of the input image in ORIGCFS using any of the preceding syntaxes.

[IMDEN,DENOISEDCFS,ORIGCFS,S] = `wdenoise2`(___) returns the sizes of the approximation coefficients at the coarsest scale along with the sizes of the wavelet coefficients at all scales. S is a matrix with the same structure as the S output of `wavedec2`.

[IMDEN,DENOISEDCFS,ORIGCFS,S,SHIFTS] = `wdenoise2`(___) returns the shifts along the row and column dimensions for cycle spinning. SHIFTS is 2-by-(`numshifts` + 1)² matrix where each column of SHIFTS contains the shifts along the row and column dimension used in cycle spinning and `numshifts` is the value of `CycleSpinning`.

[___] = `wdenoise2`(___ ,Name,Value) returns the denoised image with additional options specified by one or more Name,Value pair arguments, using any of the preceding syntaxes.

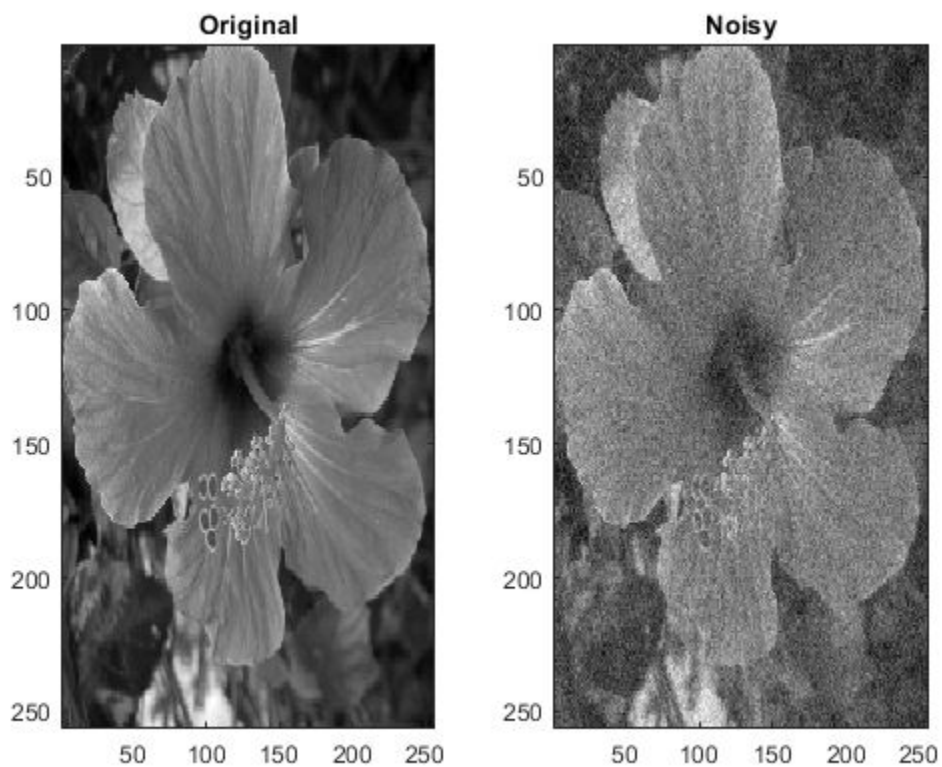
`wdenoise2`(___) with no output arguments plots the original image along with the denoised image in the current figure.

Examples

Denoise Grayscale Image Using Default Settings

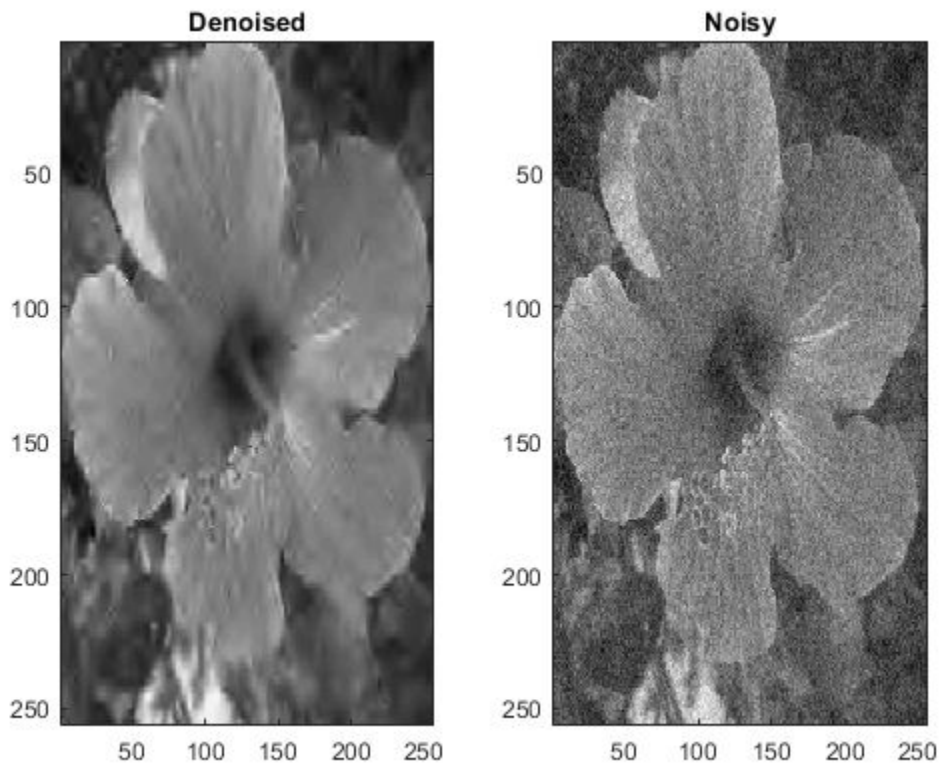
Load the structure `flower`. The structure contains a grayscale image of a flower, and a noisy version of that image. Display the original and noisy images.

```
load flower
subplot(1,2,1)
imagesc(flower.Orig)
title('Original')
subplot(1,2,2)
imagesc(flower.Noisy)
title('Noisy')
colormap gray
```



Denoise the noisy image using the default `wdenoise2` settings. Compare with the original image.

```
imden = wdenoise2(flower.Noisy);  
subplot(1,2,1)  
imagesc(imden)  
title('Denoised')  
subplot(1,2,2)  
imagesc(flower.Noisy)  
title('Noisy')  
colormap gray
```



Note the improvement in SNR before and after denoising.

```
beforeSNR = ...  
    20*log10(norm(flower.Orig(:))/norm(flower.Orig(:)-flower.Noisy(:)))
```

```
beforeSNR = 14.1300
```

```
afterSNR = ...  
    20*log10(norm(flower.Orig(:))/norm(flower.Orig(:)-imden(:)))
```

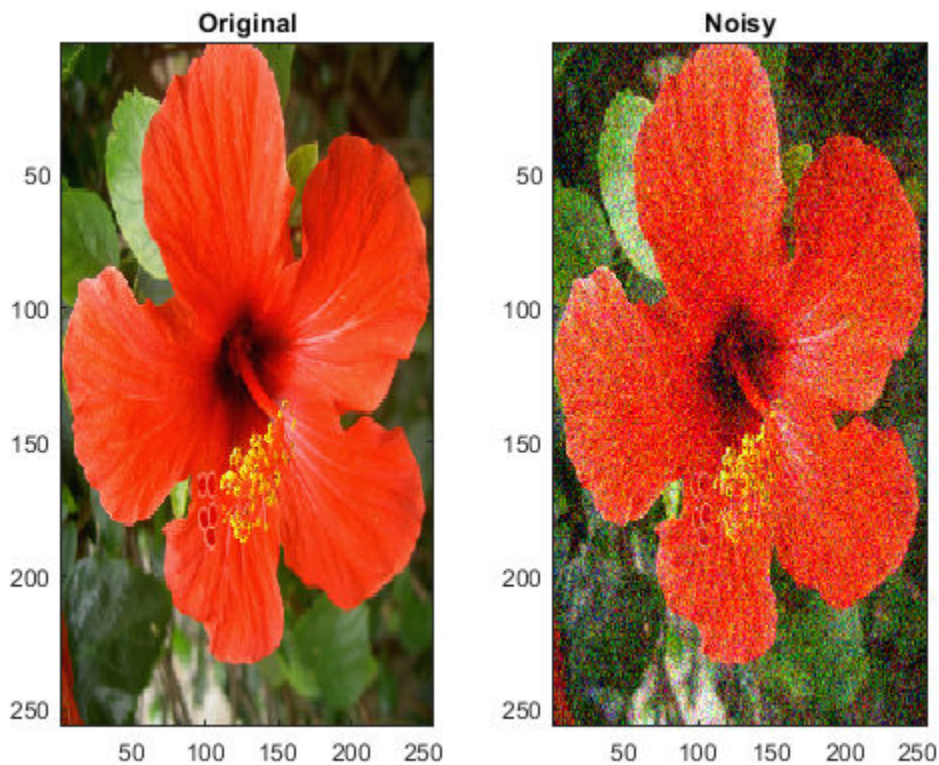
```
afterSNR = 20.1388
```


Denoise Color Image Using Cycle Spinning

This example shows how to denoise a color image using cycle spinning.

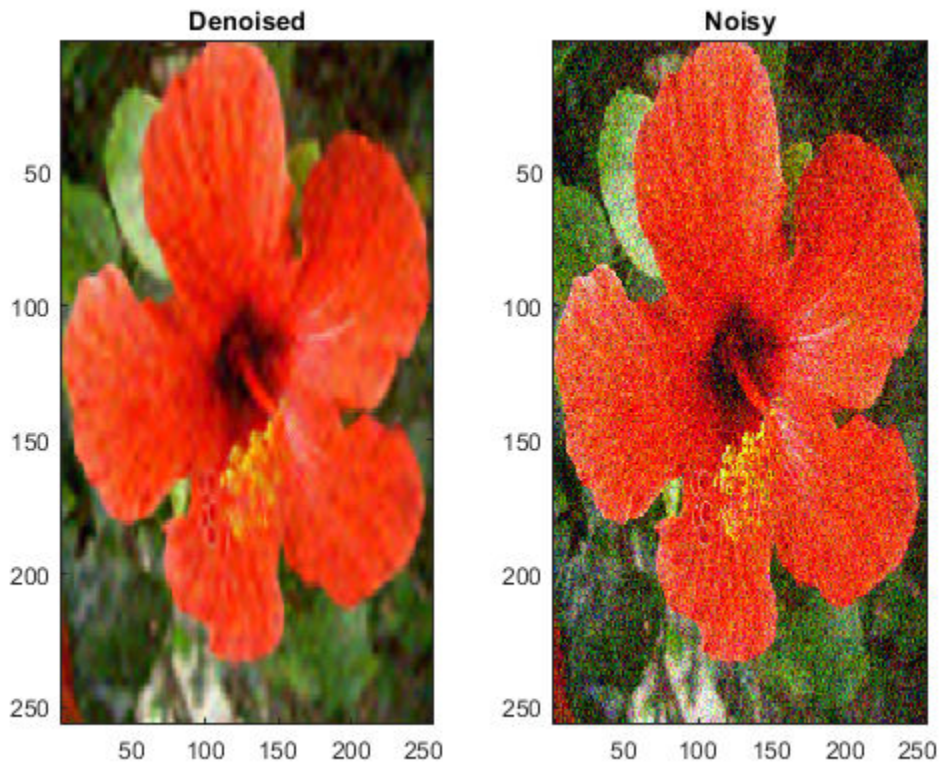
Load the structure `colorflower`. The structure contains the RGB image of a flower, and a noisy version of that image. Display the original and noisy images.

```
load colorflower
subplot(1,2,1)
imagesc(colorflower.Orig)
title('Original')
subplot(1,2,2)
imagesc(colorflower.Noisy)
title('Noisy')
```



Denoise the image down to level 2 using the default Bayesian method and cycle spinning with $(1 + 1)^2$ shifts. Display the noisy and denoised images.

```
imden = wdenoise2(colorflower.Noisy,2,'CycleSpinning',1);  
figure  
subplot(1,2,1)  
imagesc(imden)  
title('Denoised')  
subplot(1,2,2)  
imagesc(colorflower.Noisy)  
title('Noisy')
```



Compute the SNR before and after denoising.

```
beforeSNR = ...
    20*log10(norm(colorflower.Orig(:))/norm(colorflower.Orig(:)-colorflower.Noisy(:)))

beforeSNR = 11.2217

afterSNR = ...
    20*log10(norm(colorflower.Orig(:))/norm(colorflower.Orig(:)-imden(:)))

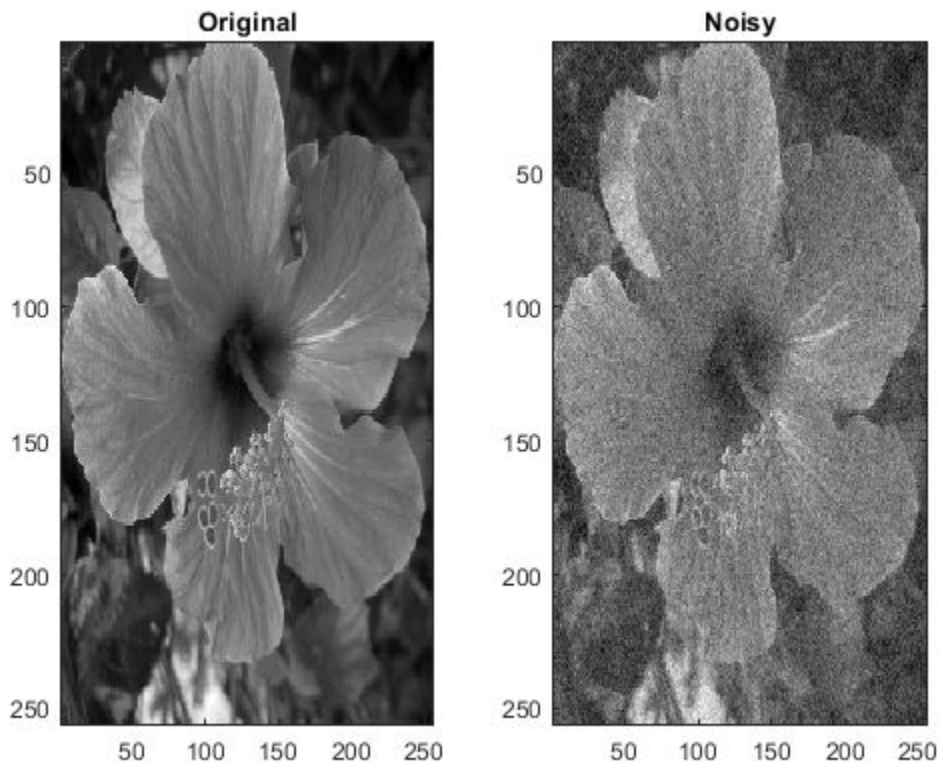
afterSNR = 19.7578
```

Denoise Image Using Specific Subband

This example shows how to denoise an image using a specific subband to estimate the variance of the noise.

Load the structure `flower`. The structure contains the grayscale image of a flower, and a noisy version of that image. Display the original and noisy images.

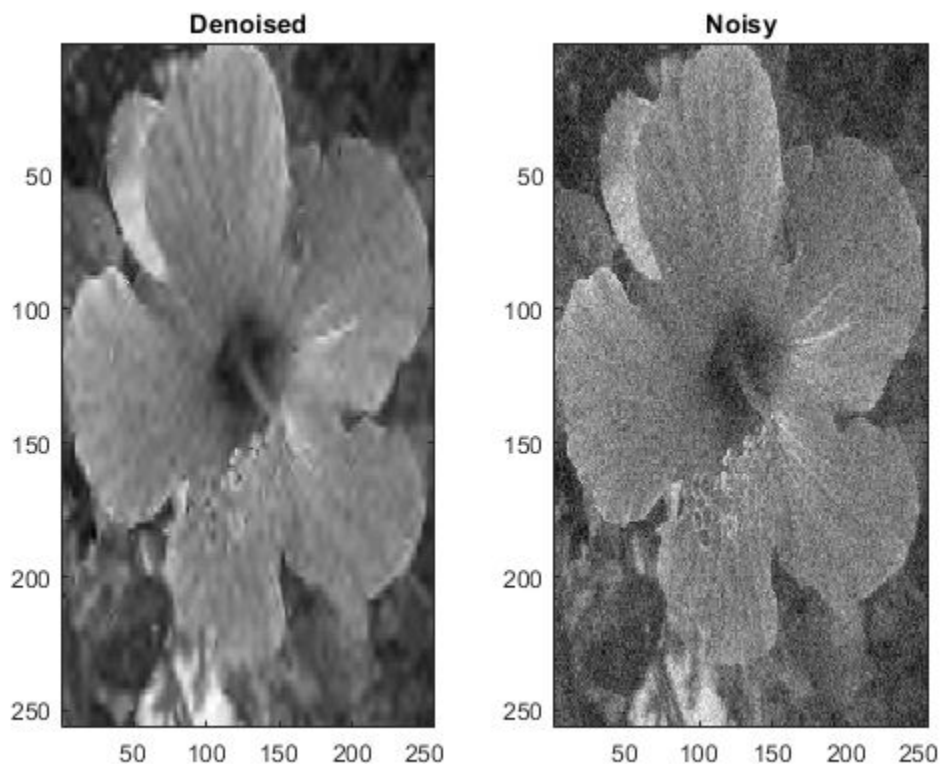
```
load flower
subplot(1,2,1)
imagesc(flower.Orig)
title('Original')
subplot(1,2,2)
imagesc(flower.Noisy)
title('Noisy')
colormap gray
```



Denoise the image down to level 2 using the False Discovery Rate method with a Q-value of 0.01. Denoise only based on the diagonal wavelet coefficients. Display the denoised and noisy images.

```
imden = wdenoise2(flower.Noisy,2,...  
    'DenoisingMethod',{'FDR',0.01},...  
    'NoiseDirection',"d");  
figure  
subplot(1,2,1)  
imagesc(imden)  
title('Denoised')  
subplot(1,2,2)  
imagesc(flower.Noisy)
```

```
title('Noisy')  
colormap gray
```



Compute the SNR before and after denoising.

```
beforeSNR = ...  
    20*log10(norm(flower.Orig(:))/norm(flower.Orig(:)-flower.Noisy(:)))
```

```
beforeSNR = 14.1300
```

```
afterSNR = ...  
    20*log10(norm(flower.Orig(:))/norm(flower.Orig(:)-imden(:)))
```

```
afterSNR = 19.9164
```

Input Arguments

IM — Input image

real-valued 2-D or 3-D matrix

Input image to denoise, specified as a real-valued 2-D or 3-D matrix. If IM is 3-D, IM is assumed to be a color image in the RGB color space and the third dimension of IM must be 3. For RGB images, `wdenoise2` by default projects the image onto its PCA color space before denoising. To denoise an RGB image in the original color space, use the `'ColorSpace'` name-value pair.

LEVEL — Wavelet decomposition level

positive integer

Wavelet decomposition level used for denoising, specified as a positive integer. LEVEL is a positive integer less than or equal to $\text{floor}(\log_2(\min([M \ N])))$ where M and N are the row and column sizes of the image. If unspecified, LEVEL defaults to the minimum $\text{floor}(\log_2(\min([M \ N])))$ and $\text{wmaxlev}([M \ N], \text{wname})$ where *wname* is the wavelet used (`'bior4.4'` by default).

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'NoiseEstimate', 'LevelDependent', 'Wavelet', 'sym6'`

Wavelet — Name of wavelet

`'sym4'` (default) | character array

Name of wavelet, specified as a character array, to use for denoising. The wavelet must be orthogonal or biorthogonal. Orthogonal and biorthogonal wavelets are designated as type 1 and type 2 wavelets respectively in the wavelet manager, `wavemngr`.

- Valid built-in orthogonal wavelet families begin with `haar`, `dbN`, `fkN`, `coifN`, or `symN` where N is the number of vanishing moments for all families except `fk`. For `fk`, N is the number of filter coefficients.

- Valid biorthogonal wavelet families begin with 'biorNr.Nd' or 'rbioNd.Nr', where Nr and Nd are the number of vanishing moments in the reconstruction (synthesis) and decomposition (analysis) wavelet.

Determine valid values for the vanishing moments by using `waveinfo` with the wavelet family short name. For example, enter `waveinfo('db')` or `waveinfo('bior')`. Use `wavemngr('type',WNAME)` to determine if a wavelet is orthogonal (returns 1) or biorthogonal (returns 2).

DenoisingMethod – Denoising method

'Bayes' (default) | 'FDR' | 'Minimax' | 'SURE' | 'UniversalThreshold'

Denoising method to use to determine the denoising thresholds for the image IM.

- Bayes - Empirical Bayes

This method uses a threshold rule based on assuming measurements have independent prior distributions given by a mixture model. Because measurements are used to estimate the weight in the mixture model, the method tends to work better with more samples. By default, the posterior median rule is used to measure risk [7].

- FDR - False Discovery Rate

This method uses a threshold rule based on controlling the expected ratio of false positive detections to all positive detections. The FDR method works best with sparse data. Choosing a ratio, or Q -value, less than $1/2$ yields an asymptotically minimax estimator [1].

Note For 'FDR', there is an optional argument for the Q -value, which is the proportion of false positives. Q is a real-valued scalar between 0 and $1/2$, $0 < Q \leq 1/2$. To specify 'FDR' with a Q -value, use a cell array where the second element is the Q -value. For example, 'DenoisingMethod', {'FDR', 0.01}. If unspecified, Q defaults to 0.05.

- Minimax - Minimax Estimation

This method uses a fixed threshold chosen to yield minimax performance for mean squared error against an ideal procedure. The minimax principle is used in statistics to design estimators. See `thselect` for more information.

- SURE - Stein's Unbiased Risk Estimate

This method uses a threshold selection rule based on Stein's Unbiased Estimate of Risk (quadratic loss function). One gets an estimate of the risk for a particular threshold value (t). Minimizing the risks in (t) gives a selection of the threshold value.

- `UniversalThreshold` - Universal Threshold $\sqrt{2\ln(\text{length}(x))}$.

This method uses a fixed-form threshold yielding minimax performance multiplied by a small factor proportional to $\log(\text{length}(X))$.

ThresholdRule — Threshold rule

'Hard' | 'Soft' | 'Mean' | 'Median'

Threshold rule to use to shrink the wavelet coefficients. 'ThresholdRule' is valid for all denoising methods, but the valid options and defaults depend on the denoising method. Rules possible for different denoising methods are specified as follows:

- 'SURE', 'Minimax', 'UniversalThreshold': Valid options are 'Soft' or 'Hard'. The default is 'Soft'.
- 'Bayes': Valid options are 'Median', 'Mean', 'Soft', or 'Hard'. The default is 'Median'.
- 'FDR': The only supported option is 'Hard'. You do not need to define 'ThresholdRule' for 'FDR'

NoiseEstimate — Method of estimating variance of noise

'LevelIndependent' (default) | 'LevelDependent'

Method of estimating variance of noise in the image. Valid options are 'LevelIndependent' and 'LevelDependent'.

- 'LevelIndependent' estimates the variance of the noise based on the finest-scale (highest-resolution) wavelet coefficients.
- 'LevelDependent' estimates the variance of the noise based on the wavelet coefficients at each resolution level.

There are three wavelet subbands: horizontal, vertical, and diagonal. The value of 'NoiseDirection' specifies which subbands to use in estimating the variance.

NoiseDirection — Wavelet subbands

["h", "v", "d"] (default) | character array | scalar string

Wavelet subbands to use to estimate the variance of the noise, specified as a string vector or scalar string. Valid entries are "h", "v", or "d", for the horizontal, vertical, and diagonal subbands, respectively.

Example: 'NoiseDirection', ["h" "v"] specifies the horizontal and vertical subbands.

CycleSpinning — Number of circular shifts

0 (default) | nonnegative integer

Number of circular shifts in both the row and column directions to use for denoising IM with cycle spinning. In cycle spinning, circular shifts of the image along the row and column dimensions are denoised, shifted back, and averaged together to provide the final result.

Generally, SNR improvements are observed with cycle spinning up to 3-4 shifts and asymptote after that. Because of the asymptotic effect on SNR and the fact that $(\text{CycleSpinning}+1)^2$ images are being denoised, it is recommended to start with CycleSpinning equal to 0. Then gradually increase the number of shifts to determine if there is any improvement in SNR to justify the computational expense.

For example, specifying 'CycleSpinning', 1 results in four copies of IM being denoised:

- The original image (unshifted)
- IM circularly shifted a single-element along the row dimension
- IM circularly shifted a single-element along the column dimension
- IM circularly shifted a single-element along both the row and column dimensions

The four denoised copies of IM are denoised, reconstructed, shifted back to their original positions, and averaged together. The value of CycleSpinning represents the maximum shift along both the row and column dimensions. For RGB images, there are no shifts applied along the color space dimension.

ColorSpace — Color space

'PCA' (default) | 'Original'

Color space used for denoising an RGB image. Valid options are 'PCA' and 'Original'.

- 'PCA': The RGB image is first projected onto its PCA color space, denoised in the PCA color space, and returned to the original color space after denoising.

- `'Original'`: Denoising is done in the same color space as the input image.

`ColorSpace` is valid only for RGB images.

Output Arguments

IMDEN — Denoised image

real-valued matrix

Denoised image, returned as a matrix. The dimensions of `IM` and `IMDEN` are equal.

DENOISEDCFS — Scaling and denoised wavelet coefficients

real-valued matrix

Scaling and denoised wavelet coefficients of the denoised image, returned as a real-valued matrix. `DENOISEDCFS` is a $(\text{numshifts}+1)^2$ -by-`N` matrix where `N` is the number of wavelet coefficients in the decomposition of `IM` and `numshifts` is the value of `'CycleSpinning'`. Each row of `DENOISEDCFS` contains the denoised wavelet coefficients for one of $(\text{numshifts}+1)^2$ shifted versions of `IM`. For RGB images, `DENOISEDCFS` are the denoised coefficients in the specified color space.

The i^{th} row of `DENOISEDCFS` contains the denoised wavelet coefficients of the image circularly shifted by the amount returned in the i^{th} column of `SHIFTS`. For example, if the second column of `SHIFTS` is `[1 ; 1]`, the second row of `DENOISEDCFS` contains the denoised coefficients of the image circularly shifted by a single element in the row direction and a single element in the column direction.

ORIGCFS — Scaling and wavelet coefficients

real-valued matrix

Scaling and wavelet coefficients of the input image, returned as a real-valued 2-D matrix. `ORIGCFS` is a $(\text{numshifts}+1)^2$ -by-`N` matrix where `N` is the number of wavelet coefficients in the decomposition of `IM` and `numshifts` is the value of the `'CycleSpinning'`. Each row of `ORIGCFS` contains the wavelet coefficients for one of $(\text{numshifts}+1)^2$ shifted versions of `IM`. For RGB images, `ORIGCFS` are the original coefficients in the specified color space.

The i^{th} row of `ORIGCFS` contains the wavelet coefficients of the image circularly shifted by the amount returned in the i^{th} column of `SHIFTS`. For example, if the second column of `SHIFTS` is `[1 ; 1]`, the second row of `ORIGCFS` contains the coefficients of the image

circularly shifted by a single element in the row direction and a single element in the column direction.

S — Bookkeeping matrix

integer-valued matrix

Bookkeeping matrix. The matrix S contains the dimensions of the approximation coefficients at the coarsest scale, the sizes of the wavelet coefficients at all scales, and the size of the original input image. S is a matrix with the same structure as the S output of `wavedec2`.

SHIFTS — Image shifts

integer-valued matrix

Image shifts used in cycle spinning, returned as an integer-valued matrix. `SHIFTS` is $2\text{-by-}(\text{numshifts}+1)^2$ matrix where each column of `SHIFTS` contains the shifts along the row and column dimension used in cycle spinning.

References

- [1] Abramovich, F., Y. Benjamini, D. L. Donoho, and I. M. Johnstone. "Adapting to Unknown Sparsity by Controlling the False Discovery Rate." *Annals of Statistics*, Vol. 34, Number 2, pp. 584-653, 2006.
- [2] Antoniadis, A., and G. Oppenheim, eds. *Wavelets and Statistics*. Lecture Notes in Statistics. New York: Springer Verlag, 1995.
- [3] Donoho, D. L. "Progress in Wavelet Analysis and WVD: A Ten Minute Tour." *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.
- [4] Donoho, D. L., I. M. Johnstone. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*, Vol. 81, pp. 425-455, 1994.
- [5] Donoho, D. L. "De-noising by Soft-Thresholding." *IEEE Transactions on Information Theory*, Vol. 42, Number 3, pp. 613-627, 1995.
- [6] Donoho, D. L., I. M. Johnstone, G. Kerkycharian, and D. Picard. "Wavelet Shrinkage: Asymptopia?" *Journal of the Royal Statistical Society, series B*, Vol. 57, No. 2, pp. 301-369, 1995.

[7] Johnstone, I. M., and B. W. Silverman. "Needles and Straw in Haystacks: Empirical Bayes Estimates of Possibly Sparse Sequences." *Annals of Statistics*, Vol. 32, Number 4, pp. 1594-1649, 2004.

See Also

wavedec2 | wdenoise

Introduced in R2019a

wenergy

Energy for 1-D wavelet or wavelet packet decomposition

Syntax

```
[Ea,Ed] = wenergy(C,L)
E = wenergy(T)
```

Description

For a one-dimensional wavelet decomposition $[C, L]$ (see `wavedec` for details), $[Ea, Ed] = wenergy(C, L)$ returns Ea , which is the percentage of energy corresponding to the approximation and Ed , which is the vector containing the percentages of energy corresponding to the details.

For a wavelet packet tree T (see `wptree`, `wpdec`, `wpdec2`), $E = wenergy(T)$ returns a vector E , which contains the percentages of energy corresponding to the terminal nodes of the tree T . In this case, `wenergy` is a method of the `wptree` object T , which overloads the previous `wenergy` function.

Examples

```
% Example 1: 1-D wavelet decomposition
%-----
load noisbump
[C,L] = wavedec(noisbump,4,'sym4');
[Ea,Ed] = wenergy(C,L)

Ea =

    88.2860

Ed =

    2.1560    1.2286    1.4664    6.8630
```

```
% Example 2: 1-D wavelet packet decomposition
%-----
load noisbump
T = wpdec(noisbump,3,'sym4');
E = wenergy(T)

E =

95.0329  1.4664  0.6100  0.6408  0.5935  0.5445  0.5154
0.5965
```

Introduced before R2006a

wenergy2

Energy for 2-D wavelet decomposition

Syntax

```
[Ea,Eh,Ev,Ed] = wenergy2(C,S)  
[Ea,EDetail] = wenergy2(C,S)
```

Description

For a two-dimensional wavelet decomposition $[C, S]$ (see `wavedec2` for details), $[Ea, Eh, Ev, Ed] = wenergy2(C, S)$ returns Ea , which is the percentage of energy corresponding to the approximation, and vectors Eh, Ev, Ed , which contain the percentages of energy corresponding to the horizontal, vertical, and diagonal details, respectively.

$[Ea, EDetail] = wenergy2(C, S)$ returns Ea , and $EDetail$, which is the sum of vectors Eh, Ev , and Ed .

Examples

```
load detail  
[C,S] = wavedec2(X,2,'sym4');  
[Ea,Eh,Ev,Ed] = wenergy2(C,S)
```

```
Ea =  
    89.3520
```

```
Eh =  
    1.8748    2.7360
```

```
Ev =  
    1.5860    2.6042
```

```
Ed =
```

```
    0.7539    1.0932
[Ea,EDetails] = wenergy2(C,S)
Ea =
    89.3520
EDetails =
    4.2147    6.4334
```

Introduced before R2006a

wentropy

Entropy (wavelet packet)

Syntax

```
E = wentropy(X,T)
E = wentropy(X,T,P)
```

Description

`E = wentropy(X,T)` returns the entropy specified by `T` of the vector or matrix `X`.

`E = wentropy(X,T,P)` returns the entropy where `P` is a parameter depending on `T`.

`E = wentropy(X,T,θ)` is equivalent to `E = wentropy(X,T)`.

Examples

Signal Entropy

This example shows the different values of entropy of a random signal.

For purposes of reproducibility, reset the random seed and generate a random signal.

```
rng default
x = randn(1,200);
```

Compute the Shannon entropy of `x`.

```
e = wentropy(x, 'shannon')
e = -224.5551
```

Compute the log energy entropy of `x`.

```
e = wentropy(x, 'log energy')
```

```
e = -229.5183
```

Compute the threshold entropy of x with the threshold entropy equal to 0.2.

```
e = wentropy(x, 'threshold', 0.2)
```

```
e = 168
```

Compute the Sure entropy of x with the threshold equal to 3.

```
e = wentropy(x, 'sure', 3)
```

```
e = 35.7962
```

Compute the norm entropy of x with power equal to 1.1

```
e = wentropy(x, 'norm', 1.1)
```

```
e = 173.6578
```

You can use your own entropy function `ABC` with `wentropy`. Your function must be defined in a `.m` file, and the first line must be of the form:

```
function e = ABC(x)
```

where x is a vector and e is a real number. The new entropy can be used by typing

```
e = wentropy(x, 'user', 'ABC')
```

or more directly

```
e = wentropy(x, 'ABC')
```

The function file `myEntropy.m` returns the normalized Shannon entropy of a signal. Compute the normalized Shannon entropy of x .

```
w = wentropy(x, 'myEntropy')
```

```
w = -1.1228
```

Input Arguments

X — Input data

real-valued vector or matrix

Input data, specified as a real-valued vector or matrix.

T — Entropy type

'shannon' | 'log energy' | 'threshold' | 'sure' | 'norm' | 'user' | '*FunName*'

Entropy type, specified as one of the following:

Entropy Type (T)	Threshold Parameter (P)	Comments
'shannon'		P is not used.
'log energy'		P is not used.
'threshold'	$0 \leq P$	P is the threshold.
'sure'	$0 \leq P$	P is the threshold.
'norm'	$1 \leq P$	P is the power.
'user'	Character vector	P is a character vector containing the file name of your own entropy function, with a single input x.
' <i>FunName</i> '	No constraints on P	FunName is any character vector other than the previous entropy types listed. FunName contains the file name of your own entropy function, with x as input and P as an additional parameter to your entropy function.

T and the threshold parameter P together define the entropy criterion.

Note The 'user' option is historical and still kept for compatibility, but it is obsoleted by the last option described in the table above. The *FunName* option do the same as the 'user' option and in addition gives the possibility to pass a parameter to your own entropy function.

P — Threshold parameter

real number | character vector | string scalar

Threshold parameter, specified by a real number, character vector, or string scalar. P and the entropy type T together define the entropy criterion.

Output Arguments

E — Entropy

real number

Entropy of X, returned as a real number.

Definitions

Entropy

Functionals verifying an additive-type property are well suited for efficient searching of binary-tree structures and the fundamental splitting property of the wavelet packets decomposition. Classical entropy-based criteria match these conditions and describe information-related properties for an accurate representation of a given signal. Entropy is a common concept in many fields, mainly in signal processing. The following example lists different entropy criteria. Many others are available and can be easily integrated. In the following expressions, s is the signal and $(s_i)_i$ the coefficients of s in an orthonormal basis.

The entropy E must be an additive cost function such that $E(0) = 0$ and

$$E(s) = \sum_i E(s_i)$$

- The (nonnormalized) Shannon entropy.

$$E1(s_i) = s_i^2 \log(s_i^2)$$

so,

$$E1(s) = -\sum_i s_i^2 \log(s_i^2)$$

with the convention $0 \log(0) = 0$.

- The concentration in l^p norm entropy with $1 \leq p$.

$$E2(s_i) = |s_i|^p \text{ so } E2(s) = \sum_i |s_i|^p = \|s\|_p^p$$

- The “log energy” entropy.

$$E3(s_i) = \log(s_i^2)$$

so,

$$E3(s) = \sum_i \log(s_i^2)$$

with the convention $\log(0) = 0$.

- The threshold entropy.

$E4(s_i) = 1$ if $|s_i| > p$ and 0 elsewhere so $E4(s) = \#\{i \text{ such that } |s_i| > p\}$ is the number of time instants when the signal is greater than a threshold p .

- The “SURE” entropy.

$E5(s) = n - \#\{i \text{ such that}$

$$|s_i| \leq p\} + \sum_i \min(s_i^2, p^2)$$

For more information, see “Wavelet Packets for Compression and Denoising”.

References

- [1] Coifman, R. R., and M. V. Wickerhauser. "Entropy-based Algorithms for best basis selection." *IEEE Transactions on Information Theory*. Vol. 38, Number 2, March 1992, pp. 713-718.
- [2] Donoho, D. L., and I. M. Johnstone. "Ideal denoising in an orthonormal basis chosen from a library of bases." *Comptes Rendus Acad. Sci. Paris, Ser. I*. Vol. 319, 1994, pp. 1317-1322.

See Also

wpdec | wpdec2 | wpdencmp

Introduced before R2006a

wextend

Extend vector or matrix

Syntax

```
YEXT= wextend(TYPE,MODE,X,LEN)  
YEXT = wextend( ____,LOC)
```

Description

YEXT= wextend(TYPE,MODE,X,LEN) extends real-valued input vector or matrix X by length LEN, using the TYPE method and MODE extension. The TYPE specifies the dimension of the extension. The MODE specifies the rule to apply to fill in values in the extension.

YEXT = wextend(____,LOC) also specifies the location of the extension.

Examples

Extending Vectors and Matrices

Extend Vector

Extend a vector using a number of different methods.

Create a vector and set the extension length to 2.

```
len = 2;  
x = [1 2 3]
```

```
x = 1×3
```

```
    1    2    3
```

Perform a zero-pad extension. To verify that different forms of the input arguments are possible, perform this extension twice. The result is the same both times.

```
xextzpd1 = wextend('1', 'zpd', x, len)
```

```
xextzpd1 = 1×7
```

```
0 0 1 2 3 0 0
```

```
xextzpd2 = wextend('1D', 'zpd', x, len, 'b')
```

```
xextzpd2 = 1×7
```

```
0 0 1 2 3 0 0
```

Perform a half-point symmetric extension.

```
xextsym = wextend('1D', 'sym', x, len)
```

```
xextsym = 1×7
```

```
2 1 1 2 3 3 2
```

Perform a periodic extension. Since the input vector is of odd length, `wextend` appends an extra example to the end before extending using the 'ppd' mode. This sample is equal to the last value on the right.

```
xextper = wextend('1D', 'per', x, len)
```

```
xextper = 1×8
```

```
3 3 1 2 3 3 1 2
```

Extend Matrix

Extend a small matrix using a number of different methods.

Create a matrix and set the extension length to 2.

```
len = 2;
```

```
X = [1 2 3; 4 5 6]
```



```
X = 2×3
```

```
  1   2   3
  4   5   6
```

Perform a zero-pad extension of the array.

```
Xextzpd = wextend(2, 'zpd', X, len)
```

```
Xextzpd = 6×7
```

```
  0   0   0   0   0   0   0
  0   0   0   0   0   0   0
  0   0   1   2   3   0   0
  0   0   4   5   6   0   0
  0   0   0   0   0   0   0
  0   0   0   0   0   0   0
```

Perform a half-point symmetric extension of the array.

```
Xextsym = wextend('2D', 'sym', X, len)
```

```
Xextsym = 6×7
```

```
  5   4   4   5   6   6   5
  2   1   1   2   3   3   2
  2   1   1   2   3   3   2
  5   4   4   5   6   6   5
  5   4   4   5   6   6   5
  2   1   1   2   3   3   2
```

Extend uint8 Data Beyond Range Limits

Observe the effects of symmetric, antisymmetric, and smooth extensions on a `uint8` vector when values are at or near the limits of the data type's range.

Symmetric Extensions

The smallest `uint8` integer is 0, and the largest is 255. Create a vector of `uint8` integers that includes those limits.

```
dataVector = uint8([0 1 2 253 254 255])
```

```
dataVector = 1x6 uint8 row vector
```

```
    0     1     2  253  254  255
```

Obtain whole-point and half-point symmetric extensions of the vector. Extend the vector by two values on the left and right.

```
wholePointSym = wextend('1','symw',dataVector,2)
```

```
wholePointSym = 1x10 uint8 row vector
```

```
    2     1     0     1     2  253  254  255  254  253
```

```
halfPointSym = wextend('1','symh',dataVector,2)
```

```
halfPointSym = 1x10 uint8 row vector
```

```
    1     0     0     1     2  253  254  255  255  254
```

Extending symmetrically never results in values outside the `uint8` range.

Antisymmetric Extensions

Create a type `double` copy of the vector, and then obtain a whole-point antisymmetric extension of the copy. The extension includes negative values and values greater than 255.

```
dataVectorDouble = double(dataVector);
```

```
wholePointAsymDouble = wextend('1','asymw',dataVectorDouble,2)
```

```
wholePointAsymDouble = 1x10
```

```
   -2   -1    0    1    2  253  254  255  256  257
```

Obtain a whole-point antisymmetric extension of the original `uint8` vector. Values outside the `uint8` range are mapped to the closest `uint8` integer, which is 0 for negative values and 255 for values greater than 255.

```
wholePointAsym = wextend('1','asymw',dataVector,2)
```

```
wholePointAsym = 1x10 uint8 row vector
```

```
    0    0    0    1    2  253  254  255  255  255
```

Now obtain half-point antisymmetric extensions of the `double` copy and the original `uint8` vector.

```
halfPointAsymDouble = wextend('1','asymh',dataVectorDouble,2)
```

```
halfPointAsymDouble = 1x10
```

```
   -1    0    0    1    2  253  254  255  -255  -254
```

```
halfPointAsym = wextend('1','asymh',dataVector,2)
```

```
halfPointAsym = 1x10 uint8 row vector
```

```
    0    0    0    1    2  253  254  255    0    0
```

As with the whole-point antisymmetric extension, negative values in the extended `uint8` data are mapped to 0.

Smooth Extensions

Obtain order-0 smooth extensions of the `double` copy and the original `uint8` vector.

```
smooth0Double = wextend('1','sp0',dataVectorDouble,2)
```

```
smooth0Double = 1x10
```

```
    0    0    0    1    2  253  254  255  255  255
```

```
smooth0 = wextend('1','sp0',dataVector,2)
```

```
smooth0 = 1x10 uint8 row vector
```

```
    0    0    0    1    2  253  254  255  255  255
```

Results are identical. Next, obtain an order-1 smooth extension of each vector.

```
smooth1Double = wextend('1','sp1',dataVectorDouble,2)
```

```
smooth1Double = 1x10
    -2    -1     0     1     2  253  254  255  256  257

smooth1 = wextend('1','sp1',dataVector,2)
smooth1 = 1x10 uint8 row vector
     0     0     0     1     2  253  254  255  255  255
```

The values in the `double` result that are outside the `uint8` range are mapped to the closest `uint8` values in the `uint8` extension.

Extend int8 Data Beyond Range Limits

Observe the effects of symmetric, antisymmetric, and smooth extensions of `int8` data when values are at or near the limits of the data type's range.

Symmetric Extensions

The smallest `int8` integer is `-128`, and the largest is `127`. Create a vector of `int8` integers that includes those limits.

```
dataVector = int8([-128 -127 -126 125 126 127])
dataVector = 1x6 int8 row vector
    -128    -127    -126     125     126     127
```

Obtain whole-point and half-point symmetric extensions of the data. Extend the vector by two values on the left and right.

```
wholePointSym = wextend('1','symw',dataVector,2)
wholePointSym = 1x10 int8 row vector
    -126    -127    -128    -127    -126     125     126     127     126     125

halfPointSym = wextend('1','symh',dataVector,2)
```

```
halfPointSym = 1x10 int8 row vector
    -127    -128    -128    -127    -126    125    126    127    127    126
```

Extending symmetrically never results in values outside the `int8` range.

Antisymmetric Extensions

Create a type `double` copy of the vector, and then obtain a whole-point antisymmetric extension of the copy. The extension includes negative values less than `-128` and values greater than `127`.

```
dataVectorDouble = double(dataVector);
wholePointsAsymDouble = wextend('1','asymw',dataVectorDouble,2)

wholePointsAsymDouble = 1x10
    -130    -129    -128    -127    -126    125    126    127    128    129
```

Obtain a whole-point antisymmetric extension of the original `int8` vector. Values outside the `int8` range are mapped to the closest `int8` integer, which is `-128` for values less than `-128` and `127` for values greater than `127`.

```
wholePointAsym = wextend('1','asymw',dataVector,2)

wholePointAsym = 1x10 int8 row vector
    -128    -128    -128    -127    -126    125    126    127    127    127
```

Now obtain half-point antisymmetric extensions of the `double` copy and the original `int8` vector.

```
halfPointAsymDouble = wextend('1','asymh',dataVectorDouble,2)

halfPointAsymDouble = 1x10
    127    128    -128    -127    -126    125    126    127    -127    -126

halfPointAsym = wextend('1','asymh',dataVector,2)

halfPointAsym = 1x10 int8 row vector
```

```
127    127   -128   -127   -126    125    126    127   -127   -126
```

In the double result, the first value is 127, which can be represented as an `int8` integer. The second value is 128, which cannot be represented as an `int8` integer. Therefore, in the `int8` result, it is being mapped to 127. The remaining values in the type double result can all be represented as `int8` integers.

Smooth Extensions

Obtain order-0 smooth extensions of the double copy and the original `int8` vector.

```
smooth0Double = wextend('1','sp0',dataVectorDouble,2)
```

```
smooth0Double = 1x10
```

```
-128 -128 -128 -127 -126 125 126 127 127 127
```

```
smooth0 = wextend('1','sp0',dataVector,2)
```

```
smooth0 = 1x10 int8 row vector
```

```
-128 -128 -128 -127 -126 125 126 127 127 127
```

The results are identical. Now obtain an order-1 smooth extension of each vector.

```
smooth1Double = wextend('1','sp1',dataVectorDouble,2)
```

```
smooth1Double = 1x10
```

```
-130 -129 -128 -127 -126 125 126 127 128 129
```

```
smooth1 = wextend('1','sp1',dataVector,2)
```

```
smooth1 = 1x10 int8 row vector
```

```
-128 -128 -128 -127 -126 125 126 127 127 127
```

The values in the `double` result outside the `int8` range are mapped to the closest `int8` values in the `int8` extension.

Input Arguments

TYPE — Extension method

1 | '1' | '1d' | '1D' | 2 | '2' | '2d' | '2D' | 'ar' | 'addrow' | 'ac' | 'addcol'

Extension method used on the input, specified as one of the values listed here.

TYPE	Description
1, '1', '1d', or '1D'	1-D extension
2, '2', '2d', or '2D'	2-D extension
'ar' or 'addrow'	Add rows
'ac' or 'addcol'	Add columns

Data Types: `double` | `char`

MODE — Specific extension

'zpd' | 'sp0' | 'spd' | 'sp1' | 'sym' | 'symh' | 'symw' | 'asym' | 'asymh' | 'asymw' | 'ppd' | 'per'

Specific extension method to use to extend the input, specified as one of the values listed here.

MODE	Description
'zpd'	Zero extension
'sp0'	Smooth extension of order 0
'spd' (or 'sp1')	Smooth extension of order 1
'sym' or 'symh'	Symmetric padding (half point): boundary value symmetric replication
'symw'	Symmetric padding (whole point): boundary value symmetric replication
'asym' or 'asymh'	Antisymmetric padding (half point): boundary value antisymmetric replication

MODE	Description
'asymw'	Antisymmetric padding (whole point): boundary value antisymmetric replication
'ppd'	Periodized extension (1)
'per'	Periodized extension (2) If the signal length is odd, <code>wextend</code> appends on the right a copy of the last value, and performs the extension using the 'ppd' mode. Otherwise, 'per' reduces to 'ppd'. This rule also applies to images.

For more information on symmetric extension modes, see [1].

Note The extension modes 'sp0' and 'spd' (or 'sp1') cast the data internally to double precision before performing the extension. For integer data types, `wextend` warns if one of the following occurs.

- The conversion to double causes a loss of precision.
 - The requested extension results in integers beyond the range where double precision numbers can represent consecutive integers exactly.
-

Data Types: char

X — Input data

real-valued vector or matrix

Input data, specified as a real-valued vector or matrix.

Data Types: single | double | int8 | int16 | int32 | int64 | uint8 | uint16 | uint32 | uint64

LEN — Length of extension

nonnegative integer | two-element vector of nonnegative integers

Length of extension, specified as a nonnegative integer or two-element vector of nonnegative integers. You can extend a matrix by expressing LEN as [LROW, LCOL], where LROW is the number of rows to add and LCOL is the number of columns to add. You

can perform a 2-D extension of a matrix by the same amount in both directions by specifying LEN as single integer.

An extension of length 0 is equivalent to the null extension.

Example: `wextend('2D', 'sym', [1 2 3 4; 5 6 7 8], [2 0])` extends only two rows up and two rows down.

LOC — Location of extension

'l' | 'u' | 'r' | 'd' | 'b' | 'n' | two-character array

Location of extension, specified as one or a pair of the following:

- 'l' — Extension left
- 'u' — Extension up
- 'r' — Extension right
- 'd' — Extension down
- 'b' — Extension on both sides
- 'n' — Null extension

The valid and default values for LOC, and the behavior of LEN, depend on the specified TYPE.

TYPE	LOC
1, '1', '1d' or '1D'	'l', 'u', 'r', 'd', 'b', or 'n' Example: <code>wextend('1D', 'zpd', X, 3, 'r')</code> extends input vector X three elements to the right. Default: 'b' LEN is the length of the extension.
2, '2', '2d' or '2D'	[LOCROW, LOCCOL], where LOCROW and LOCCOL are 1-D extension locations or 'n' (none). Example: <code>wextend('2D', 'zpd', X, [2 3], 'ub')</code> extends input vector or matrix X two rows up and three columns on both sides. Default: 'bb' LEN, specified as [LROW, LCOL], is the number of rows and columns to add.

TYPE	LOC
'ar' or 'addrow'	'l', 'u', 'r', 'd', 'b', or 'n' Example: <code>wextend('addrow','zpd',X,4,'d')</code> extends input vector or matrix X four rows down. Default: 'b' LEN is the number of rows to add.
'ac' or 'addcol'	'l', 'u', 'r', 'd', 'b', or 'n' Example: <code>wextend('addcol','zpd',X,1,'l')</code> extends input vector or matrix X one column to the left. Default: 'b' LEN is the number of columns to add.

Tips

For most wavelet applications, either a periodic extension or symmetric extension works fine.

Algorithms

When a value is outside the input data type's range, `wextend` maps it to the closest value of the input data type. For examples of data being extended beyond a data type's range, see “Extend uint8 Data Beyond Range Limits” on page 1-1435 and “Extend int8 Data Beyond Range Limits” on page 1-1438.

References

[1] Strang, G., and T. Nguyen. *Wavelets and Filter Banks*. Wellesley, MA: Wellesley-Cambridge Press, 1996.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

Usage notes and limitations:

- The generated code can return a column vector when MATLAB returns a row vector if all of the following conditions are true:
 - TYPE specifies a 1-D extension.
 - Input X is a variable-size vector.
 - Input X is not a variable-length row vector (1-by-:).

Code generation does not produce a warning or error message about the shape mismatch. In the output vector that the generated code returns, the values match the values in the output vector that MATLAB returns.

In this case, to generate code that returns a row vector, pass `X(:)'` instead of `X`.

- Input X must be of type `double`.

See Also

`dwtmode`

Introduced before R2006a

wfbm

Fractional Brownian motion synthesis

Syntax

```
FBM = wfbm(H,L)
FBM = wfbm(H,L,'plot')
FBM = wfbm(H,L,NS,W)
FBM = wfbm(H,L,W,NS)
wfbm(H,L,'plot',NS)
wfbm(H,L,'plot',W)
wfbm(H,L,'plot',NS,W)
wfbm(H,L,'plot',W,NS)
```

Description

`FBM = wfbm(H,L)` returns a fractional Brownian motion signal FBM of the Hurst parameter H ($0 < H < 1$) and length L , following the algorithm proposed by Abry and Sellan.

`FBM = wfbm(H,L,'plot')` generates and plots the FBM signal.

`FBM = wfbm(H,L,NS,W)` or `FBM = wfbm(H,L,W,NS)` returns the FBM using NS reconstruction steps and the sufficiently regular orthogonal wavelet W .

`wfbm(H,L,'plot',NS)` or `wfbm(H,L,'plot',W)` or `wfbm(H,L,'plot',NS,W)` or `wfbm(H,L,'plot',W,NS)` generates and plots the FBM signal.

`wfbm(H,L)` is equivalent to `WFBM(H,L,6,'db10')`.

`wfbm(H,L,NS)` is equivalent to `WFBM(H,L,NS,'db10')`.

`wfbm(H,L,W)` is equivalent to `WFBM(H,L,W,6)`.

A fractional Brownian motion (fBm) is a continuous-time Gaussian process depending on the Hurst parameter $0 < H < 1$. It generalizes the ordinary Brownian motion

corresponding to $H = 0.5$ and whose derivative is the white noise. The fBm is self-similar in distribution and the variance of the increments is given by

$$\text{Var}(fBm(t) - fBm(s)) = v |t-s|^{2H}$$

where v is a positive constant.

Examples

According to the value of H , the fBm exhibits for $H > 0.5$, long-range dependence and for $H < 0.5$, short or intermediate dependence. This example shows each situation using the wfbm file, which generates a sample path of this process.

```
% Generate fBm for H = 0.3 and H = 0.7

% Set the parameter H and the sample length
H = 0.3; lg = 1000;
% Generate and plot wavelet-based fBm for H = 0.3
fBm03 = wfbm(H,lg,'plot');

H = 0.7;
% Generate and plot wavelet-based fBm for H = 0.7
fBm07 = wfbm(H,lg,'plot');

% The last step is equivalent to
% Define wavelet and level of decomposition
% w = 'db10'; ns = 6;
% Generate
% fBm07 = wfbm(H,lg,'plot',w,ns);
```

fBm07 clearly exhibits a stronger low-frequency component and has, locally, less irregular behavior.

Algorithms

Starting from the expression of the fBm process as a fractional integral of the white noise process, the idea of the algorithm is to build a biorthogonal wavelet depending on a given orthogonal one and adapted to the parameter H .

Then the generated sample path is obtained by the reconstruction using the new wavelet starting from a wavelet decomposition at a given level designed as follows: details

coefficients are independent random Gaussian realizations and approximation coefficients come from a fractional ARIMA process.

This method was first proposed by Meyer and Sellan and implementation issues were examined by Abry and Sellan.

Nevertheless, the samples generated following this original scheme exhibit too many high-frequency components. To circumvent this undesirable behavior Bardet et al. propose downsampling the obtained sample by a factor 10.

Two internal parameters `delta = 10` (the downsampling factor) and a threshold `prec = 1E-4`, to evaluate series by truncated sums, can be modified by the user for extreme values of `H`.

A complete overview of long-range dependence process generators is available in Bardet et al.

References

Abry, P.; F. Sellan (1996), "The wavelet-based synthesis for the fractional Brownian motion proposed by F. Sellan and Y. Meyer: Remarks and fast implementation," *Appl. and Comp. Harmonic Anal.*, 3(4), pp. 377-383.

Bardet, J.-M.; G. Lang, G. Oppenheim, A. Philippe, S. Stoev, M.S. Taqqu (2003), "Generators of long-range dependence processes: a survey," *Theory and applications of long-range dependence*, Birkhäuser, pp. 579-623.

See Also

wfbmesti

Introduced before R2006a

wfbmesti

Parameter estimation of fractional Brownian motion

Syntax

```
HEST = wfbmesti(X)
```

Description

`HEST = wfbmesti(X)` returns a one-by-three vector `HEST` which contains three estimates of the fractal index H of the input signal X . The signal X is assumed to be a realization of fractional Brownian motion with Hurst index H .

The first two elements of the vector are estimates based on the second derivative with the second computed in the wavelet domain.

The third estimate is based on the linear regression in loglog plot, of the variance of detail versus level.

A fractional Brownian motion (fBm) is a continuous-time Gaussian process depending on the so-called Hurst parameter $0 < H < 1$. It generalizes the ordinary Brownian motion corresponding to $H = 0.5$ and whose derivative is the white noise. The fBm is self-similar in distribution and the variance of the increments is

$$\text{Var}(fBm(t) - fBm(s)) = v |t-s|^{2H}$$

where v is a positive constant.

This special form of the variance of the increments suggests various ways to estimate the parameter H . One can find in Bardet et al. a survey of such methods. The `wfbmesti` file provides three different estimates. The first one, due to Istas and Lang, is based on the discrete second-order derivative. The second one is a wavelet-based adaptation and has similar properties. The third one, proposed by Flandrin, estimates H using the slope of the loglog plot of the detail variance versus the level. A more recent extension can be found in Abry et al.

Examples

Hurst Parameter Estimation

This example shows how to estimate the Hurst index of a fractional Brownian motion. The example simulates 1,000 realizations of fractional Brownian motion with $H=0.6$. Each realization consists of 10,000 samples. At the end of the simulation, the three estimates of the Hurst index are compared.

Initialize the random number generator for repeatable results. Set the Hurst index equal to 0.6 and the length of the realizations to be 10,000.

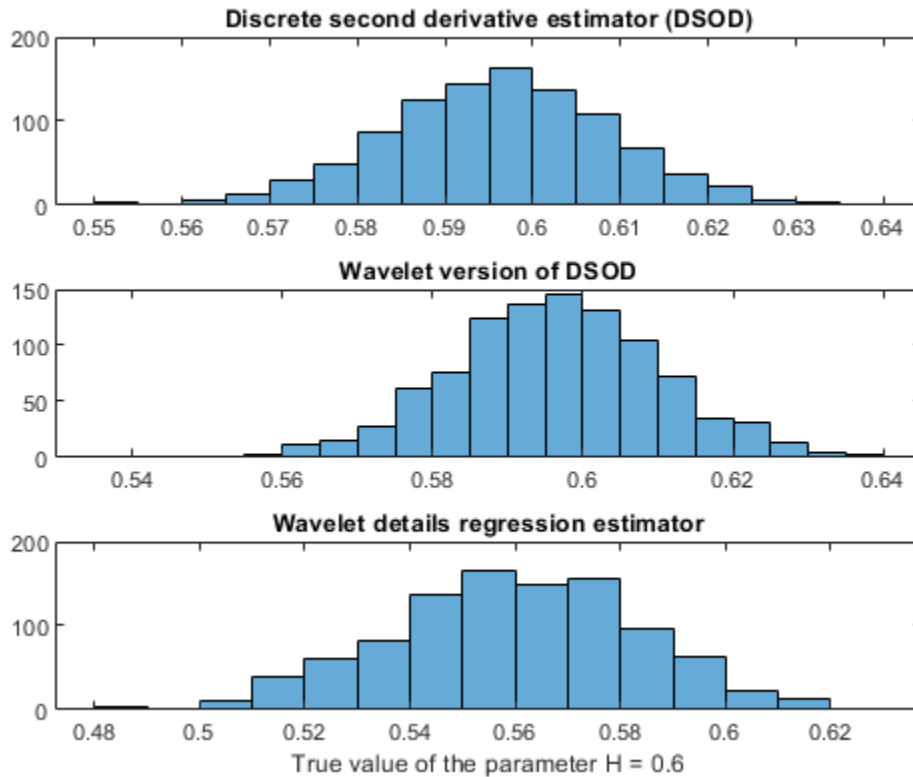
```
rng default;  
H = 0.6;  
len = 10000;
```

Generate 1,000 realizations of fractional Brownian motion and compute the estimates of the Hurst parameter.

```
n = 1000;  
Hest = zeros(n,3);  
for ii = 1:n  
    fBm06 = wfbm(H,len);  
    Hest(ii,:) = wfbmesti(fBm06);  
end
```

Compare the estimates.

```
subplot(311), histogram(Hest(:,1));  
title('Discrete second derivative estimator (DSOD)')  
subplot(312), histogram(Hest(:,2));  
title('Wavelet version of DSOD')  
subplot(313), histogram(Hest(:,3));  
title('Wavelet details regression estimator')  
xlabel('True value of the parameter H = 0.6')
```

References

Abry, P.; P. Flandrin, M.S. Taqqu, D. Veitch (2003), "Self-similarity and long-range dependence through the wavelet lens," Theory and applications of long-range dependence, Birkhäuser, pp. 527-556.

Bardet, J.-M.; G. Lang, G. Oppenheim, A. Philippe, S. Stoev, M.S. Taqqu (2003), "Semi-parametric estimation of the long-range dependence parameter: a survey," Theory and applications of long-range dependence, Birkhäuser, pp. 557-577.

Flandrin, P. (1992), "Wavelet analysis and synthesis of fractional Brownian motion," *IEEE Trans. on Inf. Th.*, 38, pp. 910-917.

Istas, J.; G. Lang (1994), "Quadratic variations and estimation of the local Hölder index of a Gaussian process," *Ann. Inst. Poincaré*, 33, pp. 407-436.

See Also

wfbm

Introduced before R2006a

wfilters

Wavelet filters

Syntax

```
[LoD,HiD,LoR,HiR] = wfilters(wname)
[F1,F2] = wfilters(wname,type)
```

Description

`[LoD,HiD,LoR,HiR] = wfilters(wname)` returns the four lowpass and highpass, decomposition and reconstruction filters associated with the orthogonal or biorthogonal wavelet `wname`.

`[F1,F2] = wfilters(wname,type)` returns the pair of `type` filters associated with the orthogonal or biorthogonal wavelet `wname`. For example, `wfilters('db6','h')` returns the pair of highpass filters `HiD` and `HiR` associated with the `db6` wavelet.

Examples

Compute Four Filters

Set the wavelet name.

```
wname = 'db5';
```

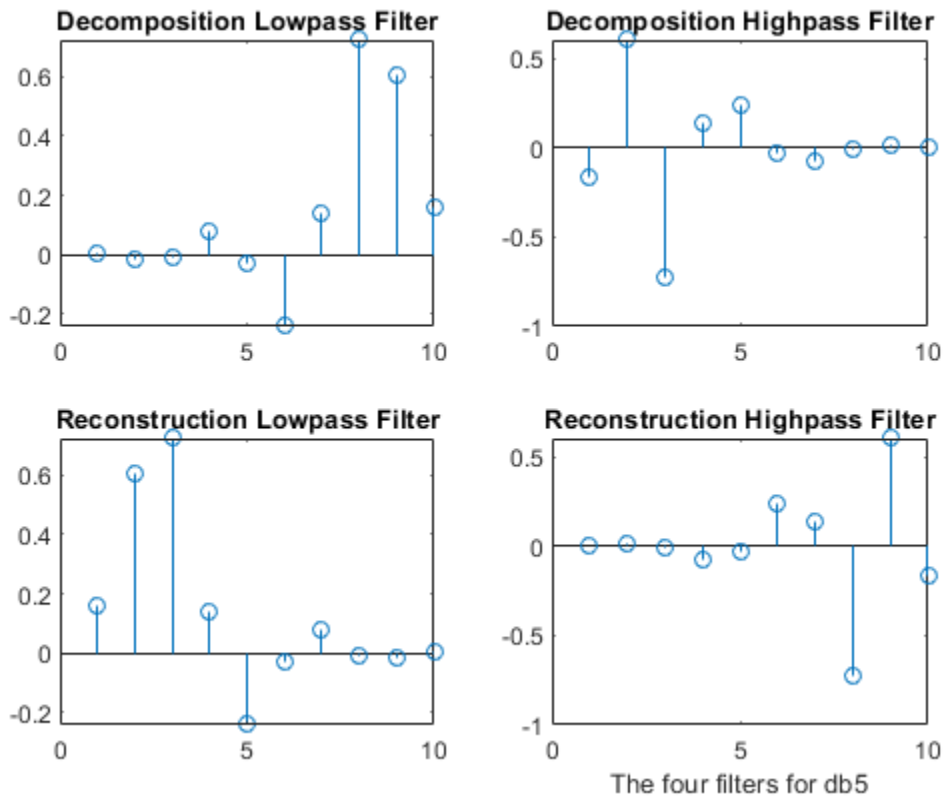
Compute the four filters associated with wavelet name specified by `wname` and plot the results.

```
[LoD,HiD,LoR,HiR] = wfilters(wname);
subplot(2,2,1)
stem(LoD)
title('Decomposition Lowpass Filter')
subplot(2,2,2)
```

```

stem(HiD)
title('Decomposition Highpass Filter')
subplot(2,2,3)
stem(LoR)
title('Reconstruction Lowpass Filter')
subplot(2,2,4)
stem(HiR)
title('Reconstruction Highpass Filter')
xlabel(['The four filters for ',wname])

```



Input Arguments

wname — Name of orthogonal or biorthogonal wavelet

'haar' | 'db1' | 'db2' | 'coif1' | 'coif2' | ...

Name of orthogonal or biorthogonal wavelet, specified as one of the values listed here.

Wavelet Families	Wavelets
Daubechies	'db1' or 'haar', 'db2', ..., 'db10', ..., 'db45'
Coiflets	'coif1', ..., 'coif5'
Symlets	'sym2', ..., 'sym8', ..., 'sym45'
Fejer-Korovkin filters	'fk4', 'fk6', 'fk8', 'fk14', 'fk22'
Discrete Meyer	'dmey'
Biorthogonal	'bior1.1', 'bior1.3', 'bior1.5' 'bior2.2', 'bior2.4', 'bior2.6', 'bior2.8' 'bior3.1', 'bior3.3', 'bior3.5', 'bior3.7' 'bior3.9', 'bior4.4', 'bior5.5', 'bior6.8'
Reverse Biorthogonal	'rbio1.1', 'rbio1.3', 'rbio1.5' 'rbio2.2', 'rbio2.4', 'rbio2.6', 'rbio2.8' 'rbio3.1', 'rbio3.3', 'rbio3.5', 'rbio3.7' 'rbio3.9', 'rbio4.4', 'rbio5.5', 'rbio6.8'

type — Type of filter pair

'd' | 'r' | 'l' | 'h'

Type of filter pair to return, specified as one of the values listed here.

type	Description
'd'	Decomposition filters (LoD and HiD)
'r'	Reconstruction filters (LoR and HiR)
'l'	Lowpass filters (LoD and LoR)
'h'	Highpass filters (HiD and HiR)

Output Arguments

LoD — Decomposition lowpass filter

real-valued vector

Decomposition lowpass filter, returned as a real-valued vector, associated with the wavelet `wname`.

HiD — Decomposition highpass filter

real-valued vector

Decomposition highpass filter, returned as a real-valued vector, associated with the wavelet `wname`.

LoR — Reconstruction lowpass filter

real-valued vector

Reconstruction lowpass filter, returned as a real-valued vector, associated with the wavelet `wname`.

HiR — Reconstruction highpass filter

real-valued vector

Reconstruction highpass filter, returned as a real-valued vector, associated with the wavelet `wname`.

F1, F2 — Filter pair

real-valued vectors

Filter pair of requested type, returned, specified as one of the pairs of filters listed here.

type	Description	Filter Pair
'd'	Decomposition filters	LoD and HiD
'r'	Reconstruction filters	LoR and HiR
'l'	Lowpass filters	LoD and LoR
'h'	Highpass filters	HiD and HiR

References

- [1] Daubechies, I. *Ten Lectures on Wavelets*. CBMS-NSF Regional Conference Series in Applied Mathematics. Philadelphia, PA: Society for Industrial and Applied Mathematics, 1992.
- [2] Mallat, S. G. "A Theory for Multiresolution Signal Decomposition: The Wavelet Representation." *IEEE Transactions on Pattern Analysis and Machine Intelligence*. Vol. 11, Issue 7, July 1989, pp. 674-693.

See Also

`biorfilt` | `orthfilt` | `waveinfo` | `wavemngr`

Introduced before R2006a

wfusimg

Fusion of two images

Syntax

```
XFUS = wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH)
[XFUS,TFUS,TX1,TX2] = wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH)
wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH,FLAGPLOT)
```

Description

The principle of image fusion using wavelets is to merge the wavelet decompositions of the two original images using fusion methods applied to approximations coefficients and details coefficients (see Zeeuw and Misiti et al.).

`XFUS = wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH)` returns the fused image `XFUS` obtained by fusion of the two original images `X1` and `X2`. Each fusion method, defined by `AFUSMETH` and `DFUSMETH`, merges in a specific way detailed below, the decompositions of `X1` and `X2`, at level `LEVEL` and using wavelet `WNAME`.

`AFUSMETH` and `DFUSMETH` define the fusion method for approximations and details, respectively.

`[XFUS,TFUS,TX1,TX2] = wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH)` returns, in addition to matrix `XFUS`, three objects of the class `WDECTREE` associated with `XFUS`, `X1`, and `X2` respectively (see `@WDECTREE`).

`wfusimg(X1,X2,WNAME,LEVEL,AFUSMETH,DFUSMETH,FLAGPLOT)` also plots the objects `TFUS`, `TX1`, and `TX2`.

`Fusmeth` denotes `AFUSMETH` or `DFUSMETH`. Available fusion methods are

- Simple — `Fusmeth` can be 'max', 'min', 'mean', 'img1', 'img2' or 'rand', which merges the two approximations or details structures obtained from `X1` and `X2` element-wise by taking the maximum, the minimum, the mean, the first element, the second element, or a randomly chosen element

- Parameter-dependent — Fusedmeth is of the following form

```
Fusedmeth = struct('name',nameMETH,'param',paramMETH)
```

where nameMETH can be

'linear'	
'UD_fusion'	Up-down fusion
'DU_fusion'	Down-up fusion
'RL_fusion'	Right-left fusion
'UserDEF'	User-defined fusion

For the description of these options and the paramMETH parameter, see wfulmat.

Examples

The following three examples examine the process of image fusion

- The first example merges two different images leading to a new image
- The second example restores an image from two fuzzy versions of an original image.
- The third example shows how to make an image fusion using a user defined fusion method.

```
% Example 1: Fusion of two different images

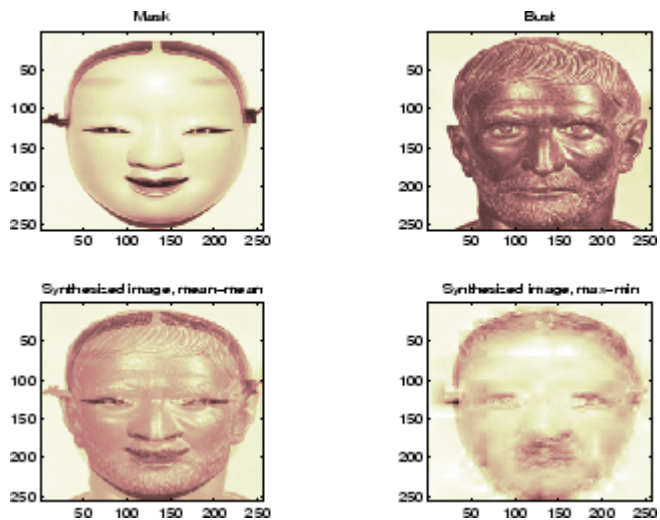
% Load two original images: a mask and a bust
load mask; X1 = X;
load bust; X2 = X;

% Merge the two images from wavelet decompositions at level 5
% using db2 by taking two different fusion methods

% fusion by taking the mean for both approximations and details
XFUSmean = wfulimg(X1,X2,'db2',5,'mean','mean');

% fusion by taking the maximum for approximations and the
% minimum for the details
XFUSmaxmin = wfulimg(X1,X2,'db2',5,'max','min');
```

```
% Plot original and synthesized images
colormap(map);
subplot(221), image(X1), axis square, title('Mask')
subplot(222), image(X2), axis square, title('Bust')
subplot(223), image(XFUSmean), axis square,
title('Synthesized image, mean-mean')
subplot(224), image(XFUSmaxmin), axis square,
title('Synthesized image, max-min')
```



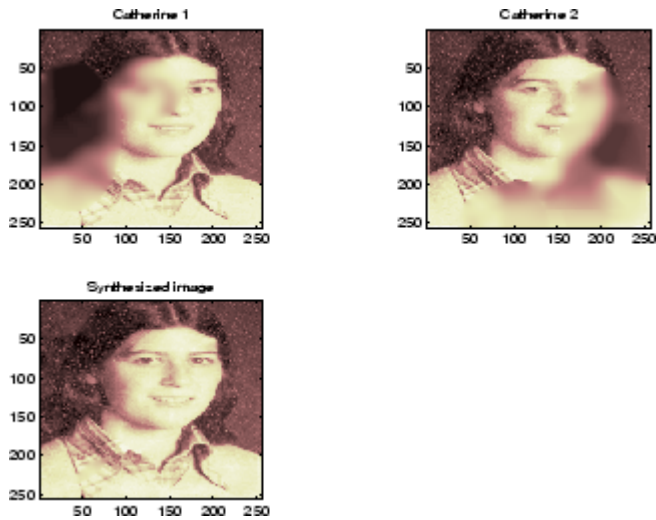
```
% Example 2: Restoration by fusion of fuzzy images
```

```
% Load two fuzzy versions of an original image
load cathe_1; X1 = X;
load cathe_2; X2 = X;
```

```
% Merge the two images from wavelet decompositions at level 5
% using sym4 by taking the maximum of absolute value of the
% coefficients for both approximations and details
XFUS = wfusing(X1,X2,'sym4',5,'max','max');
```

```
% Plot original and synthesized images
colormap(map);
subplot(221), image(X1), axis square,
title('Catherine 1')
subplot(222), image(X2), axis square,
title('Catherine 2')
```

```
subplot(223), image(XFUS), axis square,
title('Synthesized image')
```



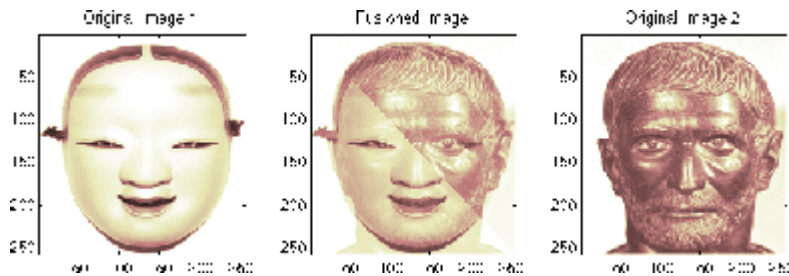
```
% The synthesized image is a restored version of good
% quality of the common underlying original image.
```

```
% Example 3: Fusion using a user defined fusion method.
% This example calls a user fusion method defined by the
% file myfus_FUN.m which is listed below at the end of
% the example.
```

```
% load two images of the same size.
load mask; A = X;
load bust; B = X;
```

```
% Define the fusion method and call the fusion function
Fus_Method = struct('name','userDEF','param','myfus_FUN');
C = wfusmat(A,B,Fus_Method);
```

```
figure;
colormap(pink(220))
subplot(1,3,1), image(A), title('Original Image 1'), axis square
subplot(1,3,2), image(C), title('Fused Image'), axis square
subplot(1,3,3), image(B), title('Original Image 2'), axis square
```



```
%*****  
% User defined fusion method. *  
%*****  
function C = myfus_FUN(A,B)  
  
D = logical(triu(ones(size(A)))); t = 0.3;  
C = A;  
C(D) = t*A(D)+(1-t)*B(D);  
C(~D) = t*B(~D)+(1-t)*A(~D);
```

Tips

X1 and X2 must be of same size (see `wextend` to resize images) and represent indexed images or truecolor images, which are m-by-n matrices or m-by-n-by-3 arrays, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

References

Zeeuw, P.M. (1998), "Wavelet and image fusion," CWI, Amsterdam, March 1998, <http://www.cwi.nl/~pauldz/>

Misiti, M.; Y. Misiti, G. Oppenheim, J.-M. Poggi (2003), "Les ondelettes et leurs applications," Hermes.

See Also

`wextend` | `wfusmat`

Introduced before R2006a

wfusmat

Fusion of two matrices or arrays

Syntax

`C = wfusmat(A,B,METHOD)`

Description

`C = wfusmat(A,B,METHOD)` returns the fused matrix `C` obtained from the matrices `A` and `B` using the fusion method defined by `METHOD`.

The matrices `A` and `B` must be of the same size. The output matrix `C` is of the same size as `A` and `B`.

Available fusion methods are

- Simple, where `METHOD` is
 - 'max' : $D = \text{abs}(A) \geq \text{abs}(B)$; $C = A(D) + B(\sim D)$
 - 'min' : $D = \text{abs}(A) \leq \text{abs}(B)$; $C = A(D) + B(\sim D)$
 - 'mean' : $C = (A+B) / 2$; $D = \text{ones}(\text{size}(A))$
 - 'rand' : $C = A(D) + B(\sim D)$; D is a Boolean random matrix
 - 'img1' : $C = A$
 - 'img2' : $C = B$
- Parameter-dependent, where `METHOD` is of the following form:

```
METHOD = struct('name',nameMETH,'param',paramMETH)
```

where `nameMETH` can be

- 'linear' : $C = A*\text{paramMETH} + B*(1-\text{paramMETH})$,

where $0 \leq \text{paramMETH} \leq 1$

- 'UD_fusion': Up-down fusion, with $\text{paramMETH} \geq 0$

```
x = linspace(0,1,size(A,1));  
P = x.^paramMETH;
```

Then each row of C is computed with

```
C(i,:) = A(i,)*(1-P(i)) + B(i,)*P(i);  
So C(1,:) = A(1,:) and C(end,:) = B(end,);
```

- 'DU_fusion': Down-up fusion
- 'LR_fusion': Left-right fusion (column-wise fusion)
- 'RL_fusion': Right-left fusion (column-wise fusion)
- 'UserDEF': User-defined fusion, `paramMETH` is a character vector or string scalar 'userFUNCTION' containing a function name such that `C = userFUNCTION(A,B)`.

In addition, `[C,D] = wfusmat(A,B,METHOD)` returns the Boolean matrix D when defined, or an empty matrix otherwise.

Introduced before R2006a

wkeep

Keep part of vector or matrix

Syntax

```
Y = wkeep(X,L,OPT)
Y = wkeep(X,L,FIRST)
Y = wkeep(X,L)
Y = wkeep(X,L,'c')
Y = wkeep(X,S,[FIRSTR FIRSTC])
```

Description

wkeep is a general utility.

For a vector, $Y = \text{wkeep}(X,L,OPT)$ extracts the vector Y from the vector X . The length of Y is L .

If OPT is equal to 'c', 'l', or 'r', Y is the central, left, or right part of X .

$Y = \text{wkeep}(X,L,FIRST)$ returns the vector $X(FIRST:FIRST+L-1)$.

$Y = \text{wkeep}(X,L)$ is equivalent to $Y = \text{wkeep}(X,L,'c')$.

For a matrix, $Y = \text{wkeep}(X,S)$ extracts the central part of the matrix X . The size of Y is S .

$Y = \text{wkeep}(X,S,[FIRSTR FIRSTC])$ extracts the submatrix of matrix X , of size S and starting from $X(FIRSTR,FIRSTC)$.

Examples

```
% For a vector.
x = 1:10;
y = wkeep(x,6,'c')
```



```
y =  
 3    4    5    6    7    8  
  
y = wkeep(x,6)  
y =  
 3    4    5    6    7    8  
  
y = wkeep(x,7,'c')  
y =  
 2    3    4    5    6    7    8  
y = wkeep(x,6,'l')  
y =  
 1    2    3    4    5    6  
  
y = wkeep(x,6,'r')  
y =  
 5    6    7    8    9    10  
  
% For a matrix.  
x = magic(5)  
x =  
 17    24     1     8    15  
 23     5     7    14    16  
  4     6    13    20    22  
 10    12    19    21     3  
 11    18    25     2     9  
  
y = wkeep(x,[3 2])  
y =  
 5     7  
 6    13  
 12    19
```

Introduced before R2006a

wmaxlev

Maximum wavelet decomposition level

Syntax

```
L = wmaxlev(S,wname)
```

Description

`L = wmaxlev(S,wname)` returns the maximum level `L` possible for a wavelet decomposition of a signal or image of size `S` using the wavelet specified by `wname` (see `wfilters` for more information). The maximum level is the last level for which at least one coefficient is correct.

`wmaxlev` returns the maximum allowed level decomposition, but in a general, a smaller value is taken.

Examples

Maximum Levels of Decomposition for a Signal and Image

Return the maximum level of decomposition of a 1-D signal with 1024 samples using the Haar wavelet.

```
s = 1024;  
wv = 'haar';  
l = wmaxlev(s,wv)
```

```
l = 10
```

Return the maximum level using the db7 wavelet.

```
wv = 'db7';  
l = wmaxlev(s,wv)
```

```
l = 6
```

Return the maximum level of decomposition for a 2-D signal of dimension 512-by-128 using the Haar wavelet.

```
s = [512 128];  
wv = 'haar';  
l = wmaxlev(s,wv)
```

```
l = 7
```

Observe the maximum level is the same when taking the minimum of the two dimensions.

```
l = wmaxlev(min(s),wv)
```

```
l = 7
```

Return the maximum level using the db7 wavelet.

```
wv = 'db7';  
l = wmaxlev(s,wv)
```

```
l = 3
```

Input Arguments

S — Size of signal or image

positive integer | two-element vector of positive integers

Size of signal or image, specified as a positive integer for a signal, or two-element vector of positive integers for an image.

Data Types: double

wname — Wavelet

character vector | string scalar

Wavelet used to determine maximum level of wavelet decomposition. The wavelet is from one of the following wavelet families: Daubechies, Coiflets, Symlets, Fejér-Korovkin, Discrete Meyer, Biorthogonal, and Reverse Biorthogonal. See `wfilters` for the wavelets available in each family.

See Also

wavedec | wavedec2 | wpdec | wpdec2

Introduced before R2006a

wmpalg

Matching pursuit

Syntax

```

YFIT = wmpalg(MPALG,Y,MPDICT)
[YFIT,R] = wmpalg(...)
[YFIT,R,COEFF] = wmpalg(...)
[YFIT,R,COEFF,IOPT] = wmpalg(...)
[YFIT,R,COEFF,IOPT,QUAL] = wmpalg(...)
[YFIT,R,COEFF,IOPT,QUAL,X] = wmpalg(...)
[YFIT,R,COEFF,IOPT,QUAL,X] = wmpalg(...,Name,Value)

```

Description

`YFIT = wmpalg(MPALG,Y,MPDICT)` returns an adaptive greedy approximation, `YFIT`, of the input signal, `Y`, in the dictionary, `MPDICT`. The adaptive greedy approximation uses the matching pursuit algorithm, `MPALG`. The dictionary, `MPDICT`, is typically an overcomplete set of vectors constructed using `wmpdictionary`.

`[YFIT,R] = wmpalg(...)` returns the residual, `R`, which is the difference vector between `Y` and `YFIT` at the termination of the matching pursuit.

`[YFIT,R,COEFF] = wmpalg(...)` returns the expansion coefficients, `COEFF`. The number of expansion coefficients depends on the number of iterations in the matching pursuit.

`[YFIT,R,COEFF,IOPT] = wmpalg(...)` returns the column indices of the retained atoms, `IOPT`. The length of `IOPT` equals the length of `COEFF` and is determined by the number of iterations in the matching pursuit.

`[YFIT,R,COEFF,IOPT,QUAL] = wmpalg(...)` returns the proportion of retained signal energy, `QUAL`, for each iteration of the matching pursuit. `QUAL` is the ratio of the ℓ^2 squared norm of the expansion coefficient vector, `COEFF`, to the ℓ^2 squared norm of the input signal, `Y`.

[YFIT,R,COEFF,IOPT,QUAL,X] = `wmpalg(...)` returns the normalized dictionary, X. X contains the unit vectors in the ℓ^2 norm corresponding to the columns of MPDICT.

[YFIT,R,COEFF,IOPT,QUAL,X] = `wmpalg(...,Name,Value)` returns an adaptive greedy approximation with additional options specified by one or more Name, Value pair arguments.

Input Arguments

MPALG

Matching pursuit algorithm as a character vector or string scalar. Valid entries are:

- 'BMP' — Basic matching pursuit
- 'OMP' — Orthogonal matching pursuit
- 'WMP' — Weak orthogonal matching pursuit

See “Matching Pursuit Algorithms”.

Default: 'BMP'

MPDICT

Matching pursuit dictionary. MPDICT is a N-by-P matrix where N is equal to the length of the input signal, Y. You can construct MPDICT using `wmpdictionary`. In matching pursuit, MPDICT is commonly a frame, or overcomplete set of vectors. You may use the Name-Value pair 'lscpt' to specify a dictionary instead of using MPDICT. If you specify a value for 'lscpt', `wmpalg` calls `wmpdictionary`.

Y

Signal for matching pursuit. Y is 1-D, real-valued row or column vector. The row dimension of MPDICT must match the length of Y.

Name-Value Pair Arguments

Specify optional comma-separated pairs of Name, Value arguments. Name is the argument name and Value is the corresponding value. Name must appear inside quotes. You can specify several name and value pair arguments in any order as Name1, Value1, ..., NameN, ValueN.

itermax

Positive integer fixing the maximum number of iterations of the matching pursuit algorithm. If you do not specify a 'maxerr' value, the number of expansion coefficients, COEFF, the number of dictionary vector indices, IOPT, and the length of the QUAL vector equal the value of 'itermax'.

Default: 25

lstcpt

A cell array of cell arrays with valid subdictionaries. This name-value pair is only valid if you do not input a dictionary in MPDICT. Each cell array describes one subdictionary. Valid subdictionaries are:

- A valid Wavelet Toolbox orthogonal or biorthogonal wavelet family short name with the number of vanishing moments and an optional decomposition level and extension mode. For example, { 'sym4' , 5} denotes the Daubechies least-asymmetric wavelet with 4 vanishing moments at level 5 and the default extension mode 'per'. If you do not specify the optional number level and extension mode, the decomposition level defaults to 5 and the extension mode to 'per'.
- A valid Wavelet Toolbox orthogonal or biorthogonal wavelet family short name preceded by wp with the number of vanishing moments and an optional decomposition level and extension mode. For example, { 'wpsym4' , 5} denotes the Daubechies least-asymmetric wavelet packet with 4 vanishing moments at level 5. If you do not specify the optional number level and extension mode, the decomposition level defaults to 5 and the extension mode to 'per'.
- 'dct' Discrete cosine transform-II basis. The DCT-II orthonormal basis is:

$$\phi_k(n) = \begin{cases} \frac{1}{\sqrt{N}} & k = 0 \\ \sqrt{\frac{2}{N}} \cos\left(\frac{\pi}{N}\left(n + \frac{1}{2}\right)k\right) & k = 1, 2, \dots, N - 1 \end{cases}$$

- 'sin' Sine subdictionary. The sine subdictionary is:

$$\phi_k(t) = \sin(2\pi kt) \quad k = 1, 2, \dots, \left\lfloor \frac{N}{2} \right\rfloor \quad 0 \leq t \leq 1$$

- 'cos' Cosine subdictionary. The cosine subdictionary is

$$\phi_k(t) = \cos(2\pi kt) \quad k = 1, 2, \dots, \left\lfloor \frac{N}{2} \right\rfloor \quad 0 \leq t \leq 1$$

- 'poly' Polynomial subdictionary. The polynomial subdictionary is:

$$p_n(t) = t^{n-1} \quad n = 1, 2, \dots, 20 \quad 0 \leq t \leq 1$$

- 'RnIdent' The shifted Kronecker delta subdictionary. The shifted Kronecker delta subdictionary is:

$$\phi_k(n) = \delta(n - k) \quad k = 0, 1, \dots, N$$

If you use the 'lstcpt' name-value pair to generate your dictionary, you can use the additional 'addbeg' and 'addend' name-value pairs to append and addend dictionary atoms. See `wmpdictionary` for details.

maxerr

Cell array containing the name of the norm and the maximum relative error in the norm expressed as a percentage. Valid norms are 'L1', 'L2', and 'Linf'. The relative error expressed as a percentage is

$$100 \frac{\|R\|}{\|Y\|}$$

where R is the residual at each iteration and Y is the input signal. For example, {'L1', 10} sets maximum acceptable ratio of the L1 norms of the residual to the input signal to 0.10.

If you specify 'maxerr', the matching pursuit terminates when the first of the following conditions is satisfied:

- The number of iterations reaches the minimum of the length of the input signal, Y , or 500:
`min(length(Y), 500)`
- The relative error falls below the percentage you specify with the 'maxerr' name-value pair.

stepplot

Number of iterations between successive plots. 'stepplot' requires a positive integer. This name-value pair is only valid when 'typeplot' is 2 or 3 ('movie' or 'stepwise').

typeplot

Type of plot to produce during the progression of matching pursuit. Valid entries for 'typeplot' are: 0 or 'none', 1 or 'one', 2 or 'movie', 3 or 'stepwise'. When 'typeplot' is 'movie' or 'stepwise', the plot updates based on the value of 'stepplot'.

Default: 0 or 'none'

wmpcfs

Optimality factor for weak orthogonal matching pursuit. The optimality factor is a real number in the interval (0,1]. This name-value pair is only valid when MPALG is 'WMP'.

Default: 0.6

Output Arguments

YFIT

Adaptive greedy approximation of the input signal, Y, in the dictionary

R

Residual after matching pursuit terminates

COEFF

Expansion coefficients in the dictionary. The selected dictionary atoms weighted by the expansion coefficients yield the approximated signal, YFIT.

IOPT

Column indices of the selected dictionary atoms. Using the column indices in IOPT with the expansion coefficients in COEFF, you can form the approximated signal, YFIT.

QUAL

Proportion of retained signal energy for each iteration in the matching pursuit. QUAL is a vector with each element equal to

$$\frac{\|\alpha_k\|_2^2}{\|Y\|_2^2}$$

where α_k is the vector of expansion coefficients after the k -th iteration.

X

The normalized matching pursuit dictionary. X is an N -by- P matrix where N is the length of the input signal, Y . The columns of X have unit norm.

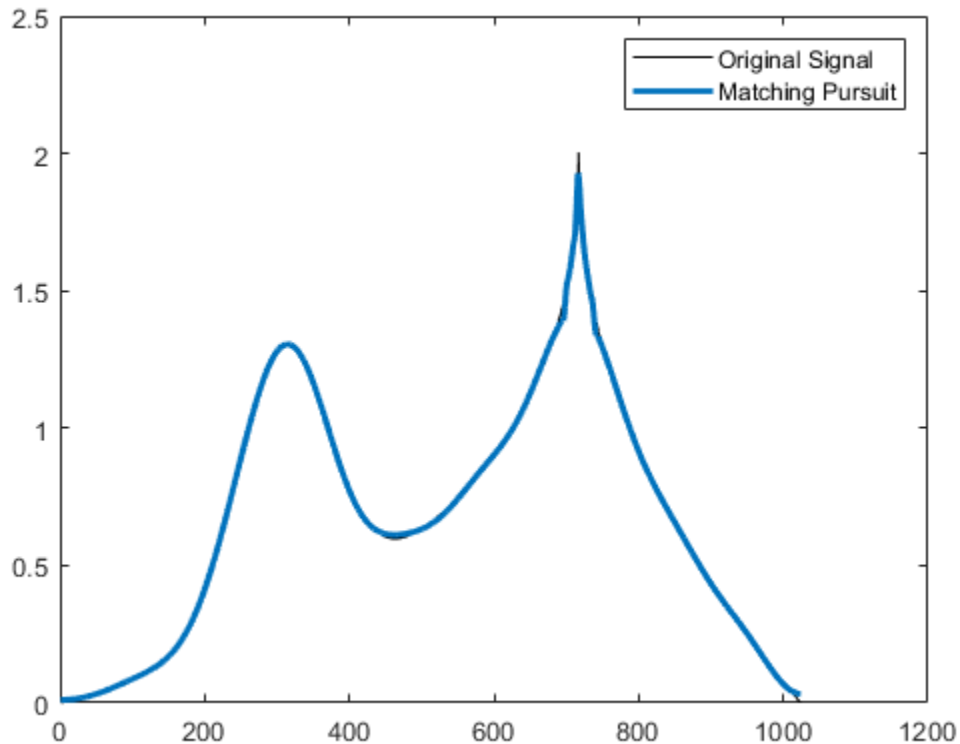
Examples

Adaptive Approximation using Orthogonal Matching Pursuit

Approximate the `cuspamax` signal with the dictionary using orthogonal matching pursuit.

Use a dictionary consisting of `sym4` wavelet packets and the DCT-II basis.

```
load cuspamax;
mpdict = wmpdictionary(length(cuspamax), 'LstCpt', ...
    {'wpsym4',2}, 'dct');
yfit = wmpalg('OMP',cuspamax,mpdict);
plot(cuspamax,'k'); hold on;
plot(yfit,'linewidth',2); legend('Original Signal',...
    'Matching Pursuit');
```



Return Residual, Expansion Coefficients, Selected Atoms, and Approximation Quality

Obtain the expansion coefficients in the dictionary, the column indices of the selected dictionary atoms, and the proportion of retained signal energy.

Create a dictionary consisting of `sym4` wavelet packets and the DCT-II basis. Approximate the `cuspmamax` signal with the dictionary using orthogonal matching pursuit.

```
load cuspmamax;
mpdict = wmpdictionary(length(cuspmamax), 'LstCpt', ...
```

```
    {'wpsym4',2}, 'dct'});  
[yfit,r,coeff,iopt,qual] = wmpalg('OMP',cuspamax,mpdict);
```

Specify the Maximum Number of Iterations

This example shows how to set the maximum number of iterations of the orthogonal matching pursuit to 50.

```
load cuspamax;  
lstcpt = {'wpsym4',1},{'wpsym4',2}, 'dct'};  
mpdict = wmpdictionary(length(cuspamax), 'LstCpt',lstcpt);  
[yfit,r,coeff,iopt,qual] = wmpalg('OMP',cuspamax,mpdict,...  
    'itermax',50);
```

Stepwise Plot of Weak Orthogonal Matching Pursuit

This example shows how to allow for a suboptimal choice in the update of the orthogonal matching pursuit. Relax the requirement to be 0.8 times the optimal assignment. Plot the results stepwise and update the plot every 5 iterations.

```
load cuspamax;  
lstcpt = {'wpsym4',1},{'wpsym4',2}, 'dct'};  
mpdict = wmpdictionary(length(cuspamax), 'LstCpt',lstcpt);  
[yfit,r,coeff,iopt,qual] = wmpalg('WMP',cuspamax,...  
    mpdict,'wmpcfs',0.8, 'typeplot','stepwise','stepplot',3);
```

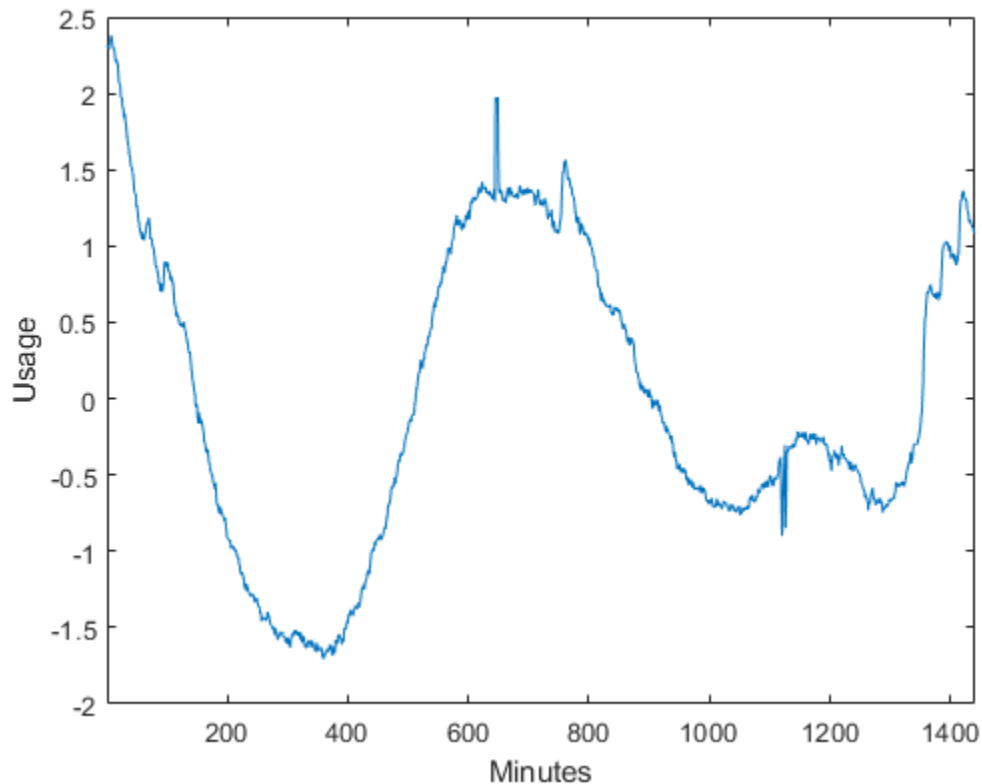
Matching Pursuit of Electricity Consumption Data

Obtain a matching pursuit of electricity consumption measured every minute over a 24-hour period.

Load and plot data. The data shows electricity consumption sampled every minute over a 24-hour period. Because the data is centered, the actual usage values are not interpretable.

```
load elec35_nor;  
y = signals(32,:);
```

```
plot(y); xlabel('Minutes'); ylabel('Usage');
set(gca,'xlim',[1 1440]);
```

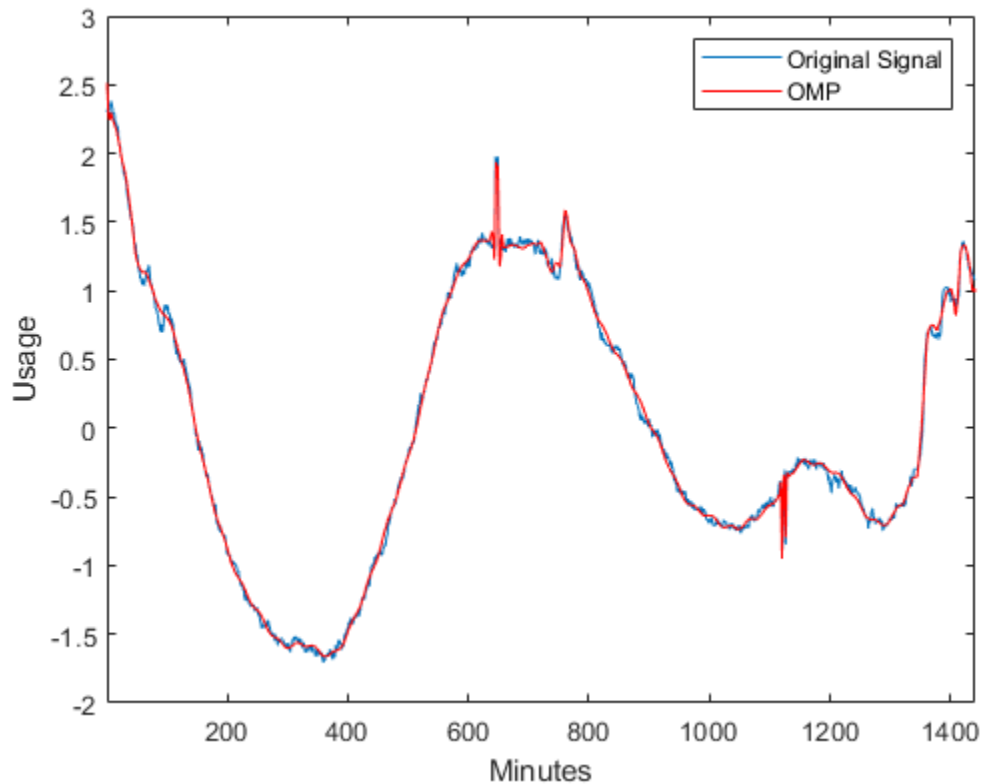


Construct a dictionary for matching pursuit consisting of the Daubechies' extremal-phase wavelet with 2 vanishing moments at level 2, the Daubechies' least-asymmetric wavelet with 4 vanishing moments at levels 1 and 4, the discrete cosine transform-II basis, and the sine basis.

```
dictionary = {'db4',2},{'dct','sin',{'sym4',1},{'sym4',4}};
[mpdict,nbvect] = wmpdictionary(length(y),'lstcpt',dictionary);
```

Implement orthogonal matching pursuit to obtain a signal approximation in the dictionary. Use 35 iterations. Plot the result.

```
[yfit,r,coef,iopt,qual] = wmpalg('OMP',y,mpdict,'itermax',35);  
plot(y); hold on;  
plot(yfit,'r'); xlabel('Minutes'); ylabel('Usage');  
legend('Original Signal','OMP','Location','NorthEast');  
set(gca,'xlim',[1 1440]);
```



Using the expansion coefficients in `coef` and the atom indices in `iopt`, construct the signal approximation, `yhat`, directly from the dictionary. Compare `yhat` with `yfit` returned by `wmpalg`.

```
[~,I] = sort(iopt);  
X = mpdict(:,iopt(I));  
yhat = X*coef(I);  
max(abs(yfit-yhat))
```

ans = 2.2204e-15

References

- [1] Cai, T.T. and Wang,L. “Orthogonal Matching Pursuit for Sparse Signal Recovery with Noise”. *IEEE Transactions on Information Theory*, vol. 57, 7, 4680–4688, 2011.
- [2] Donoho, D., Elad, M., and Temlyakov, V. “Stable Recovery of Sparse Overcomplete Representations in the Presence of Noise”. *IEEE Transactions on Information Theory*. Vol. 52, 1, 6–18, 2004.
- [3] Mallat, S. and Zhang, Z. “Matching Pursuits with Time-Frequency Dictionaries”. *IEEE Transactions on Signal Processing*, vol. 41, 12, 3397–3415, 1993
- [4] Tropp, J.A. “Greed is good: Algorithmic results for sparse approximation”. *IEEE Transactions on Information Theory*, 50, pp. 2231–2242, 2004.

See Also

Wavelet Analyzer | [wmpdictionary](#)

Topics

“Matching Pursuit”
“Matching Pursuit Using Wavelet Analyzer App”
“Matching Pursuit Algorithms”

Introduced in R2012a

wmpdictionary

Dictionary for matching pursuit

Syntax

```
MPDICT = wmpdictionary(N)
[MPDICT,NBVECT] = wmpdictionary(N)
[MPDICT,NBVECT]= wmpdictionary(N,Name,Value)
[MPDICT,NBVECT,LST] = wmpdictionary(N,Name,Value)
[MPDICT,NBVECT,LST,LONGS] = wmpdictionary(N,Name,Value)
```

Description

`MPDICT = wmpdictionary(N)` returns the N-by-P dictionary, `MPDICT`, for the default subdictionaries `{'sym4',5},{'wpsym4',5},'dct','sin'}`. The column dimension of `MPDICT` depends on `N`.

`[MPDICT,NBVECT] = wmpdictionary(N)` returns the row vector, `NBVECT`, which contains the number of vectors in each subdictionary. The order of the elements in `NBVECT` corresponds to the order of the subdictionaries and any prepended or appended subdictionaries. The sum of the elements in `NBVECT` is the column dimension of `MPDICT`.

`[MPDICT,NBVECT]= wmpdictionary(N,Name,Value)` returns the dictionary, `MPDICT`, using additional options specified by one or more `Name,Value` pair arguments.

`[MPDICT,NBVECT,LST] = wmpdictionary(N,Name,Value)` returns the cell array, `LST`, with descriptions of the subdictionaries.

`[MPDICT,NBVECT,LST,LONGS] = wmpdictionary(N,Name,Value)` returns the cell array, `LONGS`, containing the number of vectors in each subdictionary. `LONGS` is only useful for wavelet subdictionaries. In wavelet subdictionaries, the corresponding element in `LONGS` gives the number of scaling functions at the coarsest level and wavelet functions by level. See “Visualize Haar Wavelet Dictionary” on page 1-1486 for an example using `LONGS`.

Input Arguments

N

A positive integer equal to the length of your input signal. The dictionary atoms are constructed to have N elements. N equals the row dimension of the dictionary, `MPDICT`.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

addbeg

Prepended subdictionary. The prepended subdictionary is an N -by- M matrix where N is the length of the input signal. `wmpdictionary` does not check that the M column vectors of the prepended dictionary form a basis. If you do not specify a value for `lstcpt`, the subdictionary is prepended to the default dictionary. The column vectors in the prepended subdictionary do not have to be unit-norm.

addend

Appended subdictionary. The appended subdictionary is a N -by- M matrix where N is the length of the input signal. `wmpdictionary` does not check that the M column vectors of the prepended dictionary form a basis. If you do not specify a value for `lstcpt`, the subdictionary is appended to the default dictionary. The column vectors in the appended subdictionary do not have to be unit-norm.

lstcpt

A cell array of cell arrays with valid subdictionaries. Each cell array describes one subdictionary. Valid subdictionaries are:

- A valid Wavelet Toolbox orthogonal or biorthogonal wavelet family short name with the number of vanishing moments and an optional decomposition level and extension mode. For example, `{ 'sym4' , 5 }` denotes the Daubechies least-asymmetric wavelet with 4 vanishing moments at level 5 and the default extension mode `'per'`. If you do not specify the optional level and extension mode, the decomposition level defaults to 5 and the extension mode to `'per'`.

- A valid Wavelet Toolbox orthogonal or biorthogonal wavelet family short name preceded by `wp` with the number of vanishing moments and an optional decomposition level and extension mode. For example, `{'wpsym4', 5}` denotes the Daubechies least-asymmetric wavelet packet with 4 vanishing moments at level 5. If you do not specify the optional level and extension mode, the decomposition level defaults to 5 and the extension mode to `'per'`.

- `'dct'` Discrete cosine transform-II basis. The DCT-II orthonormal basis is:

$$\phi_k(n) = \begin{cases} \frac{1}{\sqrt{N}} & k = 0 \\ \sqrt{\frac{2}{N}} \cos\left(\frac{\pi}{N}\left(n + \frac{1}{2}\right)k\right) & k = 1, 2, \dots, N - 1 \end{cases}$$

- `'sin'` Sine subdictionary. The sine subdictionary is

$$\phi_k(t) = \sin(2\pi kt) \quad k = 1, 2, \dots, \left\lfloor \frac{N}{2} \right\rfloor \quad 0 \leq t \leq 1$$

where t is a linearly-spaced N -point vector.

- `'cos'` Cosine subdictionary. The cosine subdictionary is

$$\phi_k(t) = \cos(2\pi kt) \quad k = 1, 2, \dots, \left\lfloor \frac{N}{2} \right\rfloor \quad 0 \leq t \leq 1$$

where t is a linearly-spaced N -point vector.

- `'poly'` Polynomial subdictionary. The polynomial subdictionary is:

$$p_n(t) = t^{n-1} \quad n = 1, 2, \dots, 20 \quad 0 \leq t \leq 1$$

where t is a linearly-spaced N -point vector.

- `'RnIdent'` The shifted Kronecker delta subdictionary. The shifted Kronecker delta subdictionary is:

$$\phi_k(n) = \delta(n - k) \quad k = 0, 1, \dots, N$$

Default: `{'sym4', 5}, {'wpsym4', 5}, 'dct', 'sin'`

Output Arguments

MPDICT

Matching pursuit dictionary. MPDICT is an N-by-P matrix with the row dimension, N, equal to the length of the input signal. The column dimension of the matrix depends on the size of the concatenated subdictionaries.

NBVECT

Number of vectors in subdictionaries. NBVECT is a row vector containing the number of elements in each subdictionary. The order of the elements in NBVECT corresponds to the order of the subdictionaries and any prepended or appended subdictionaries.

LST

Cell array describing the dictionary. LST is a 1-by-N cell array where N is the number of subdictionaries. Each element of the cell array contains a description of a subdictionary. If you specify a prepended or appended subdictionary, the first element of LST is 'AddBeg' or 'AddEnd'. If you specify a level for the wavelet or wavelet packet, the corresponding element of LST is a 1-by-2 cell array containing the wavelet or wavelet packet name in the first element and the level in the second element.

LONGS

Cell array containing the number of elements for each subdictionary. LONGS is useful only for wavelet subdictionaries. If you specify a wavelet subdictionary, the corresponding element of LONGS provides the number of scaling functions at the coarsest level and the number of wavelets at each level. See “Visualize Haar Wavelet Dictionary” on page 1-1486 for an example using LONGS.

Examples

Default Dictionary

Create the default dictionary to represent a signal of length 100.

```
mpdict = wmpdictionary(100);
```

Discrete Cosine Transform and Kronecker Delta Dictionary

Create a DCT and shifted Kronecker delta dictionary to represent a signal of length 100.

```
mpdict = wmpdictionary(100, 'lstcpt', {'dct', 'RnIdent'});
```

Haar Wavelet Packets and Discrete Cosine Transform Dictionary

Create a Haar wavelet packet (level 2) and DCT dictionary. Return the number of atoms in each subdictionary.

```
[mpdict, nbvect] = wmpdictionary(100, 'lstcpt', {'wphaar', 2}, 'dct');
```

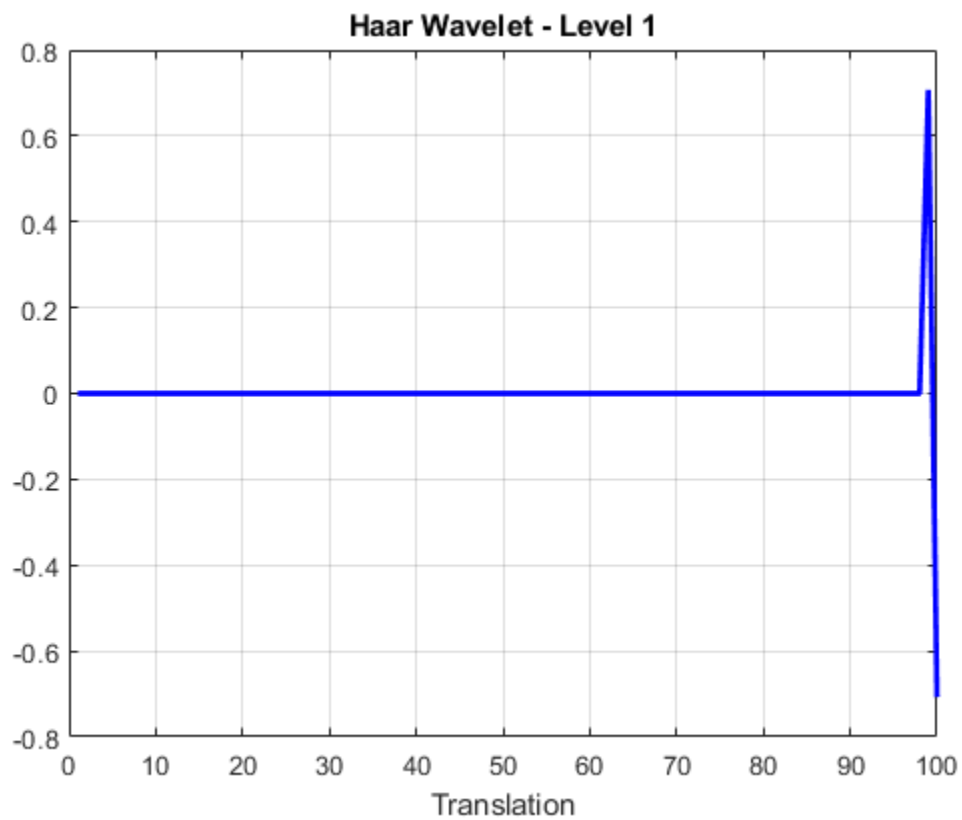
Visualize Haar Wavelet Dictionary

Use the `longs` output argument to visualize a dictionary. Create a Haar wavelet dictionary consisting of level-2 scaling functions, and level-1 and level-2 wavelet functions. Step through a plot of the translated scaling functions and wavelets by level.

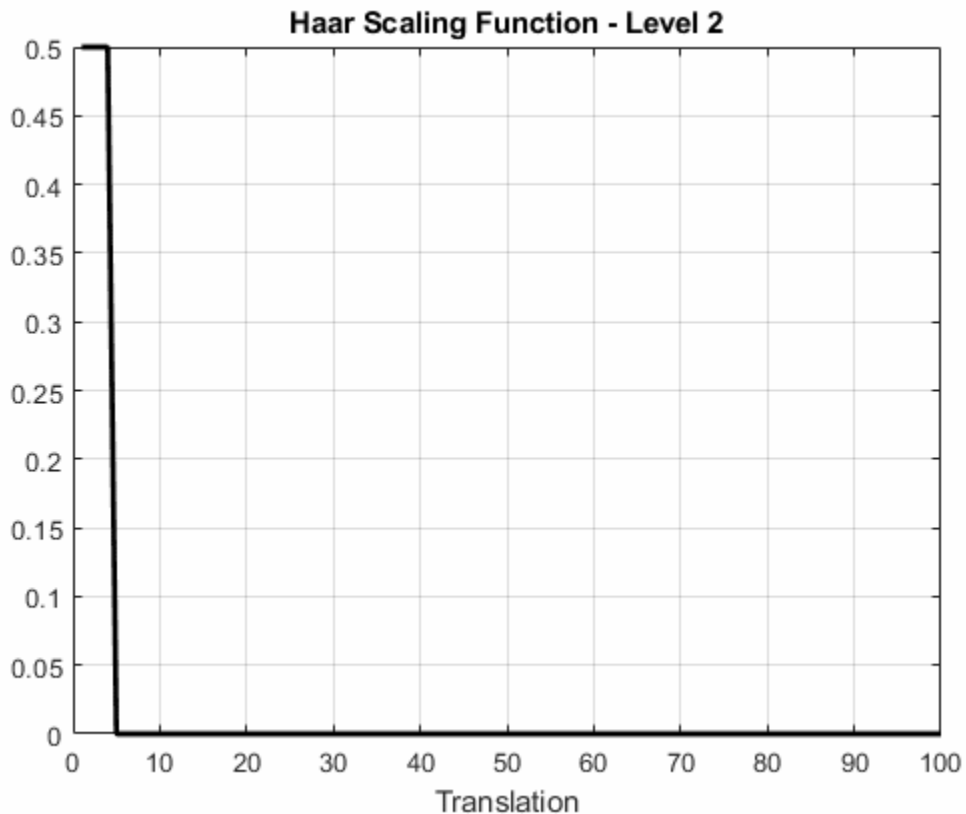
```
[mpdict, ~, ~, longs] = wmpdictionary(100, 'lstcpt', {'haar', 2});
```

```
for nn = 1:size(mpdict,2)
    if (nn<=longs{1}(1))
        plot(mpdict(:,nn), 'k', 'linewidth', 2)
        grid on
        xlabel('Translation')
        title('Haar Scaling Function - Level 2')
    elseif (nn>longs{1}(1) && nn<=longs{1}(1)+longs{1}(2))
        plot(mpdict(:,nn), 'r', 'linewidth', 2)
        grid on
        xlabel('Translation')
        title('Haar Wavelet - Level 2')
    else
        plot(mpdict(:,nn), 'b', 'linewidth', 2)
        grid on
        xlabel('Translation')
        title('Haar Wavelet - Level 1')
```

```
end  
pause(0.2)  
end
```



This animation infinitely loops through all the plots generated.



Definitions

Matching Pursuit

Matching pursuit refers to a number of greedy or weak-greedy algorithms for computing an adaptive nonlinear expansion of a signal in a *dictionary*. In the majority of matching pursuit applications, a dictionary is an overcomplete set of vectors. The elements of the dictionary are referred to as *atoms* and are typically constructed to have certain time/frequency or time/scale properties. Matching pursuit takes the NP-hard problem of finding the best nonlinear expansion in a dictionary and implements it in an energy-

preserving formulation that guarantees convergence. See “Matching Pursuit Algorithms” for more details.

References

- [1] Cai, T.T. and L. Wang “Orthogonal Matching Pursuit for Sparse Signal Recovery with Noise”. *IEEE Transactions on Information Theory*, vol. 57, 7, 4680–4688, 2011.
- [2] Donoho, D., M. Elad, and V. Temlyakov “Stable Recovery of Sparse Overcomplete Representations in the Presence of Noise”. *IEEE Transactions on Information Theory*, 52,1, 6–18, 2004.
- [3] Mallat, S. and Z. Zhang “Matching Pursuits with Time-Frequency Dictionaries”. *IEEE Transactions on Signal Processing*, vol. 41, 12, 3397–3415, 1993
- [4] Tropp, J.A. “Greed is good: Algorithmic results for sparse approximation”. *IEEE Transactions on Information Theory*, 50, pp. 2231–2242, 2004.

See Also

Wavelet Analyzer | `wmpalg`

Topics

“Matching Pursuit”
“Matching Pursuit Using Wavelet Analyzer App”
“Matching Pursuit Algorithms”

Introduced in R2012a

wmspca

Multiscale Principal Component Analysis

Syntax

```
[X_SIM, QUAL, NPC, DEC_SIM, PCA_Params] = wmspca(X, LEVEL, WNAME, NPC)  
[...] = wmspca(X, LEVEL, WNAME, 'mode', EXTMODE, NPC)  
[...] = wmspca(DEC, NPC)  
[...] = wmspca(X, LEVEL, WNAME, 'mode', EXTMODE, NPC)
```

Description

[X_SIM, QUAL, NPC, DEC_SIM, PCA_Params] = wmspca(X, LEVEL, WNAME, NPC) or
[...] = wmspca(X, LEVEL, WNAME, 'mode', EXTMODE, NPC) returns a simplified
version X_SIM of the input matrix X obtained from the wavelet-based multiscale principal
component analysis (PCA).

The input matrix X contains P signals of length N stored column-wise ($N > P$).

Wavelet Decomposition Parameters

The wavelet decomposition is performed using the decomposition level LEVEL and the
wavelet WNAME.

EXTMODE is the extended mode for the DWT (See dwtmode).

If a decomposition DEC obtained using mdwtdec is available, you can use

```
[...] = wmspca(DEC, NPC) instead of  
[...] = wmspca(X, LEVEL, WNAME, 'mode', EXTMODE, NPC).
```


Principal Components Parameter: NPC

If NPC is a vector, then it must be of length LEVEL+2. It contains the number of retained principal components for each PCA performed:

- NPC (d) is the number of retained noncentered principal components for details at level d, for $1 \leq d \leq \text{LEVEL}$.
- NPC (LEVEL+1) is the number of retained non-centered principal components for approximations at level LEVEL.
- NPC (LEVEL+2) is the number of retained principal components for final PCA after wavelet reconstruction.

NPC must be such that $0 \leq \text{NPC}(d) \leq P$ for $1 \leq d \leq \text{LEVEL}+2$.

If NPC = 'kais' (respectively, 'heur'), then the number of retained principal components is selected automatically using Kaiser's rule (or the heuristic rule).

- Kaiser's rule keeps the components associated with eigenvalues greater the mean of all eigenvalues.
- The heuristic rule keeps the components associated with eigenvalues greater than 0.05 times the sum of all eigenvalues.

If NPC = 'nodet', then the details are "killed" and all the approximations are retained.

Output Parameters

X_SIM is a simplified version of the matrix X.

QUAL is a vector of length P containing the quality of column reconstructions given by the relative mean square errors in percent.

NPC is the vector of selected numbers of retained principal components.

DEC_SIM is the wavelet decomposition of X_SIM

PCA_Params is a structure array of length LEVEL+2 such that:

- PCA_Params (d) . pc is a P-by-P matrix of principal components.

The columns are stored in descending order of the variances.

- `PCA_Params(d).variances` is the principal component variances vector.
- `PCA_Params(d).npc = NPC`

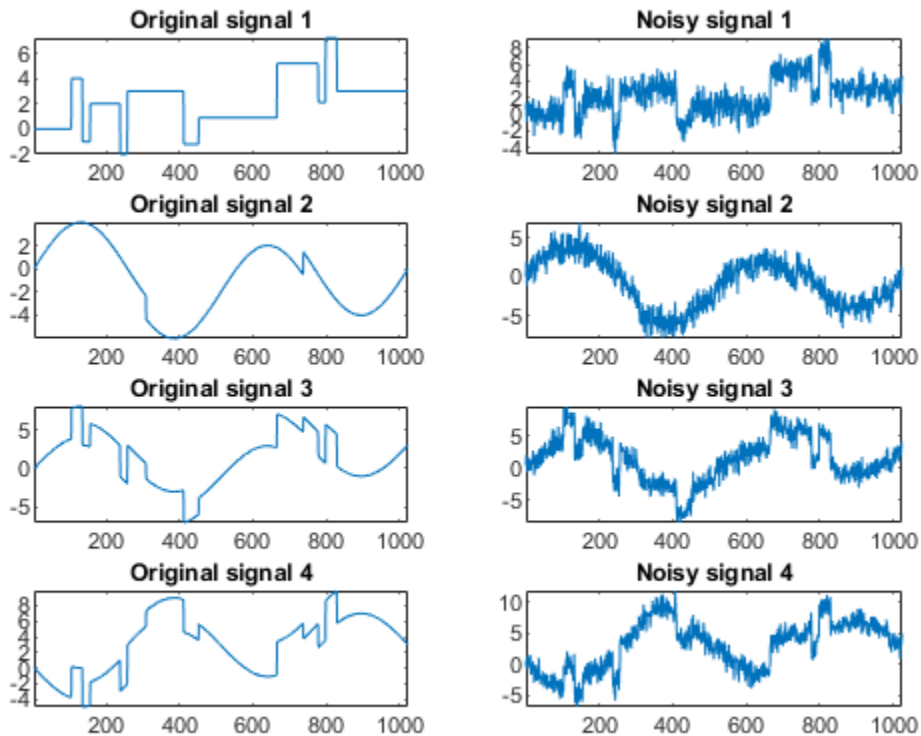
Examples

Wavelet Principal Component Analysis of Noisy Multivariate Signal

Use wavelet multiscale principal component analysis to denoise a multivariate signal.

Load the dataset consisting of four signals of length 1024. Plot the original signals and the signals with additive noise.

```
load ex4mwden;
kp = 0;
for i = 1:4
    subplot(4,2,kp+1)
    plot(x_orig(:,i))
    axis tight
    title(['Original signal ',num2str(i)])
    subplot(4,2,kp+2)
    plot(x(:,i))
    axis tight
    title(['Noisy signal ',num2str(i)])
    kp = kp + 2;
end
```



Perform the first multiscale wavelet PCA using the Daubechies least-asymmetric wavelet with four vanishing moments, `sym4`. Obtain the multiresolution decomposition down to level 5. Use the heuristic rule to decide how many principal components to retain.

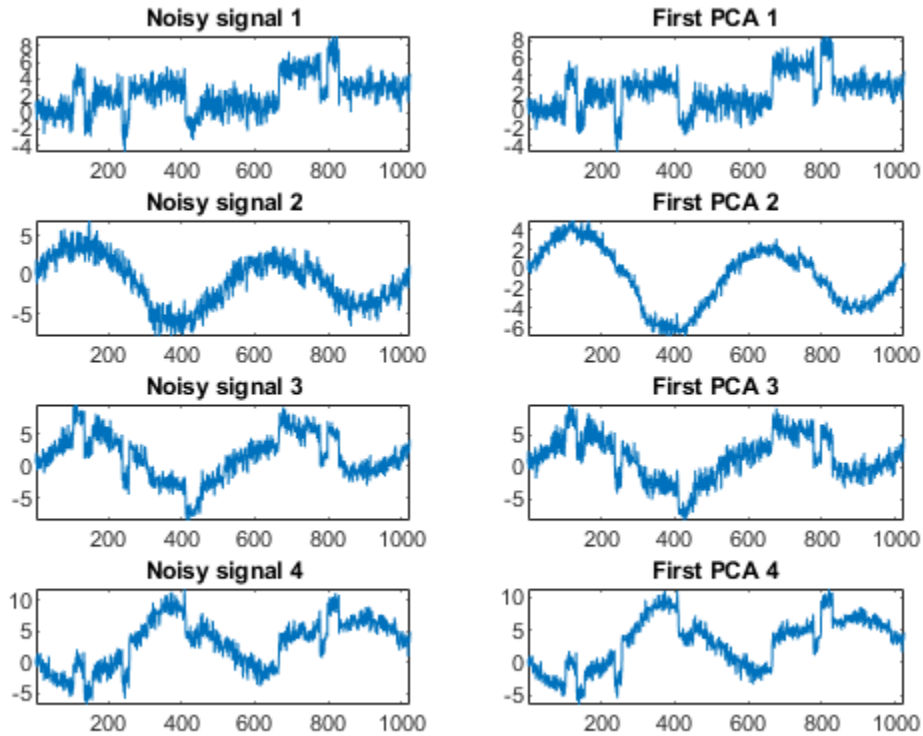
```
level = 5;
wname = 'sym4';
npc = 'heur';
[x_sim, qual, npc] = wmspca(x, level, wname, npc);
```

Plot the result and examine the quality of the approximation.

```
qual
qual = 1x4
```

97.4064 94.6863 97.8333 99.5441

```
kp = 0;
for i = 1:4
    subplot(4,2,kp+1)
    plot(x(:,i))
    axis tight
    title(['Noisy signal ',num2str(i)])
    subplot(4,2,kp+2)
    plot(x_sim(:,i))
    axis tight
    title(['First PCA ',num2str(i)])
    kp = kp+2;
end
```



The quality results are all close to 100%. The npc vector gives the number of principal components retained at each level.

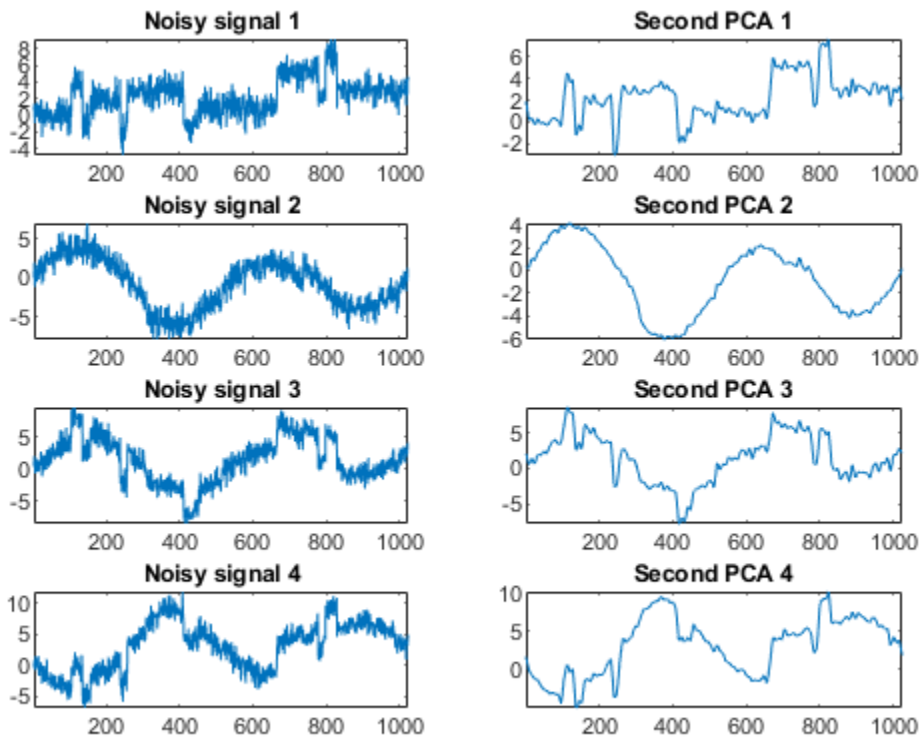
Suppress the noise by removing the principal components at levels 1-3. Perform the multiscale PCA again.

```
npc(1:3) = zeros(1,3);
[x_sim, qual, npc] = wmspca(x, level, wname, npc);
```

Plot the result.

```
kp = 0;
for i = 1:4
    subplot(4,2,kp+1)
```

```
plot(x(:,i))
axis tight
title(['Noisy signal ', num2str(i)])
subplot(4,2,kp+2)
plot(x_sim(:,i))
axis tight
title(['Second PCA ', num2str(i)])
kp = kp+2;
end
```



Algorithms

The multiscale principal components generalizes the usual PCA of a multivariate signal seen as a matrix by performing simultaneously a PCA on the matrices of details of different levels. In addition, a PCA is performed also on the coarser approximation coefficients matrix in the wavelet domain as well as on the final reconstructed matrix. By selecting conveniently the numbers of retained principal components, interesting simplified signals can be reconstructed.

References

Aminghafari, M.; Cheze, N.; Poggi, J-M. (2006), "Multivariate de-noising using wavelets and principal component analysis," *Computational Statistics & Data Analysis*, 50, pp. 2381-2398.

Bakshi, B. (1998), "Multiscale PCA with application to MSPC monitoring," *AIChE J.*, 44, pp. 1596-1610.

See Also

wmulden

Introduced in R2006b

wmulden

Wavelet multivariate denoising

Syntax

```
[X_DEN,NPC,NESTCOV,DEC_DEN,PCA_Params,DEN_Params] = ...  
wmulden(X,LEVEL,WNAME,NPC_APP,NPC_FIN,TPTR,SORH)  
[...] = wmulden(X,LEVEL,WNAME,'mode',EXTMODE,NPC_APP,...)  
[...] = wmulden(DEC,NPC_APP)  
[...] = wmulden(X,LEVEL,WNAME,'mode',EXTMODE,NPC_APP)  
[DEC,PCA_Params] = wmulden('estimate',DEC,NPC_APP,NPC_FIN)  
[X_DEN,NPC,DEC_DEN,PCA_Params] = wmulden('execute',DEC,PCA_Params)
```

Description

```
[X_DEN,NPC,NESTCOV,DEC_DEN,PCA_Params,DEN_Params] = ...  
wmulden(X,LEVEL,WNAME,NPC_APP,NPC_FIN,TPTR,SORH) or  
[...] = wmulden(X,LEVEL,WNAME,'mode',EXTMODE,NPC_APP,...) returns a  
denoised version X_DEN of the input matrix X. The strategy combines univariate wavelet  
denoising in the basis where the estimated noise covariance matrix is diagonal with  
noncentered Principal Component Analysis (PCA) on approximations in the wavelet  
domain or with final PCA.
```

The input matrix X contains P signals of length N stored column-wise where $N > P$.

Wavelet Decomposition Parameters

The wavelet decomposition is performed using the decomposition level `LEVEL` and the wavelet `WNAME`.

`EXTMODE` is the extended mode for the DWT (See `dwtmode`).

If a decomposition `DEC` obtained using `mdwtdec` is available, you can use

```
[...] = wmulden(DEC,NPC_APP) instead of
```


[...] = wmulden(X, LEVEL, WNAME, 'mode', EXTMODE, NPC_APP).

Principal Components Parameters: NPC_APP and NPC_FIN

The input selection methods NPC_APP and NPC_FIN define the way to select principal components for approximations at level LEVEL in the wavelet domain and for final PCA after wavelet reconstruction, respectively.

If NPC_APP (or NPC_FIN) is an integer, it contains the number of retained principal components for approximations at level LEVEL (or for final PCA after wavelet reconstruction).

NPC_XXX must be such that $0 \leq \text{NPC_XXX} \leq P$

NPC_APP or NPC_FIN = 'kais' or 'heur' selects the number of retained principal components using Kaiser's rule or the heuristic rule automatically.

- Kaiser's rule keeps the components associated with eigenvalues greater than the mean of all eigenvalues.
- The heuristic rule keeps the components associated with eigenvalues greater than 0.05 times the sum of all eigenvalues.

NPC_APP or NPC_FIN = 'none' is equivalent to NPC_APP or NPC_FIN = P.

Denosing Parameters: TPTR and SORH

The default values for the denosing parameters TPTR and SORH are:

TPTR = 'sqrtwolog' and SORH = 's'

- Valid values for TPTR are
'rigsure', 'heursure', 'sqrtwolog', 'minimaxi',
'penalhi', 'penalme', 'penallo'
- Valid values for SORH are:
's' (soft) or 'h' (hard)

For additional information, see wden and wbmpen.

Output Parameters

`X_DEN` is a denoised version of the input matrix `X`.

`NPC` is the vector of selected numbers of retained principal components.

`NESTCOV` is the estimated noise covariance matrix obtained using the minimum covariance determinant (MCD) estimator.

`DEC_DEN` is the wavelet decomposition of `X_DEN`.

`PCA_Params` is a structure such that:

```
PCA_Params.NEST = {pc_NEST, var_NEST, NESTCOV}
PCA_Params.APP  = {pc_APP, var_APP, npc_APP}
PCA_Params.FIN  = {pc_FIN, var_FIN, npc_FIN}
```

where:

- `pc_XXX` is a P-by-P matrix of principal components.
The columns are stored in descending order of the variances.
- `var_XXX` is the principal component variances vector.
- `NESTCOV` is the covariance matrix estimate for detail at level 1.

`DEN_Params` is a structure such that:

- `DEN_Params.thrVAL` is a vector of length `LEVEL` which contains the threshold values for each level.
- `DEN_Params.thrMETH` is a character vector containing the name of the denoising method (TPTR).
- `DEN_Params.thrTYPE` is a character variable containing the type of the thresholding (SORH).

Special Cases

`[DEC, PCA_Params] = wmulden('estimate', DEC, NPC_APP, NPC_FIN)` returns the wavelet decomposition `DEC` and the Principal Components Estimates `PCA_Params`.

`[X_DEN, NPC, DEC_DEN, PCA_Params] = wmulden('execute', DEC, PC_Params)` uses the principal components estimates `PCA_Params` previously computed.

The input value DEC can be replaced by X, LEVEL, and WNAME.

Examples

```
% Load a multivariate signal x together with
% the original signals (x_orig) and true noise
% covariance matrix (covar).

load ex4mwden

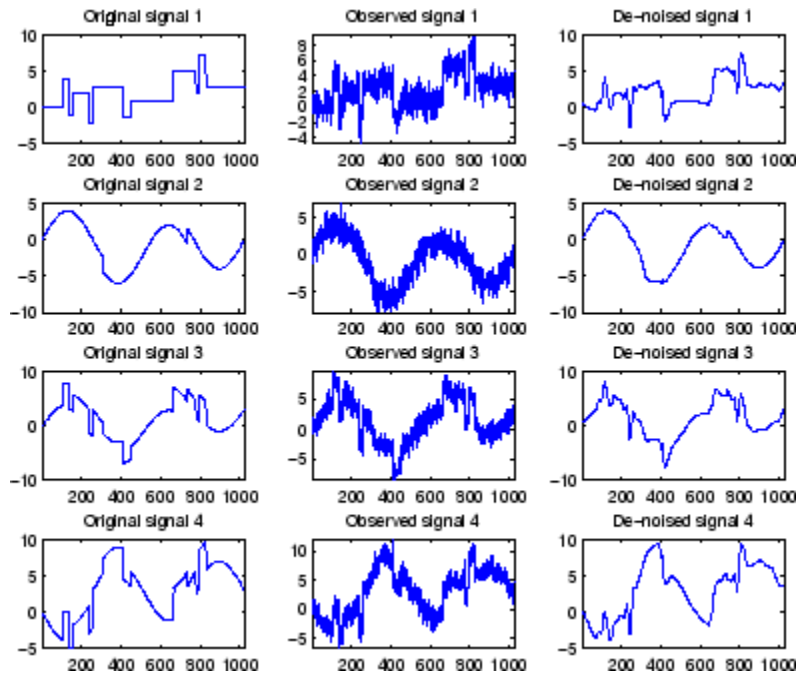
% Set the denoising method parameters.
level = 5;
wname = 'sym4';
tptr = 'sqrtwolog';
sorh = 's';

% Set the PCA parameters to select the number of
% retained principal components automatically by
% Kaiser's rule.

npc_app = 'kais';
npc_fin = 'kais';

% Perform multivariate denoising.
[x_den, npc, nestco] = wmulden(x, level, wname, npc_app, ...
                             npc_fin, tptr, sorh);

% Display the original and denoised signals.
kp = 0;
for i = 1:4
    subplot(4,3,kp+1), plot(x_orig(:,i));
    title(['Original signal ',num2str(i)])
    subplot(4,3,kp+2), plot(x(:,i));
    title(['Observed signal ',num2str(i)])
    subplot(4,3,kp+3), plot(x_den(:,i));
    title(['Denoised signal ',num2str(i)])
    kp = kp + 3;
end
```



```
% The results are good: the first function, which is
% irregular, is correctly recovered while the second
% function, more regular, is well denoised.
```

```
% The second output argument gives the numbers
% of retained principal components for PCA for
% approximations and for final PCA.
```

```
npc
```

```
npc =
```

```
    2    2
```

```
% The third output argument contains the estimated
% noise covariance matrix using the MCD based
% on the matrix of finest details.
```

```
nestco
```

```
nestco =  
  
    1.0784    0.8333    0.6878    0.8141  
    0.8333    1.0025    0.5275    0.6814  
    0.6878    0.5275    1.0501    0.7734  
    0.8141    0.6814    0.7734    1.0967  
  
% The estimation is satisfactory since the values are close  
% to the true values given by covar.  
  
covar  
  
covar =  
  
    1.0000    0.8000    0.6000    0.7000  
    0.8000    1.0000    0.5000    0.6000  
    0.6000    0.5000    1.0000    0.7000  
    0.7000    0.6000    0.7000    1.0000
```

Algorithms

The multivariate denoising procedure is a generalization of the one-dimensional strategy. It combines univariate wavelet denoising in the basis where the estimated noise covariance matrix is diagonal and non-centered Principal Component Analysis (PCA) on approximations in the wavelet domain or with final PCA.

The robust estimate of the noise covariance matrix given by the minimum covariance determinant estimator based on the matrix of finest details.

References

Aminghafari, M.; Cheze, N.; Poggi, J-M. (2006), "Multivariate de-noising using wavelets and principal component analysis," *Computational Statistics & Data Analysis*, 50, pp. 2381-2398.

Rousseeuw, P.; Van Driessen, K. (1999), "A fast algorithm for the minimum covariance determinant estimator," *Technometrics*, 41, pp. 212-223.

See Also

Functions

wdenoise | wmspca

Apps

Wavelet Signal Denoiser

Introduced in R2006b

wnoise

Noisy wavelet test data

Syntax

```
x = wnoise(fun,n)
[x,xn] = wnoise(fun,n,sqrtsnr)
[x,xn] = wnoise( ____,init)
```

Description

`x = wnoise(fun,n)` returns values `x` of the test signal `fun` evaluated at 2^n linearly spaced points from 0 to 1.

`[x,xn] = wnoise(fun,n,sqrtsnr)` returns `x` rescaled such that the standard deviation of `x` is `sqrtsnr`. `xn` is `x` corrupted by additive Gaussian white noise $N(0,1)$ and has a signal-to-noise ratio (SNR) of `sqrtsnr`².

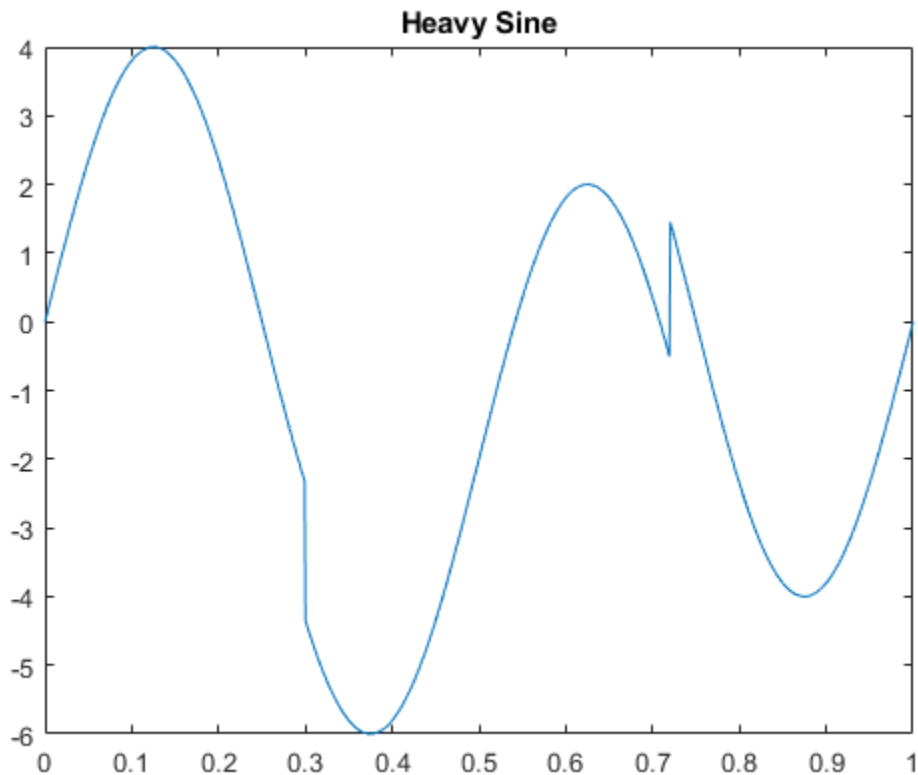
`[x,xn] = wnoise(____,init)` sets the generator seed to `init` before generating additive Gaussian white noise $N(0,1)$.

Examples

Plot Wavelet Test Signals

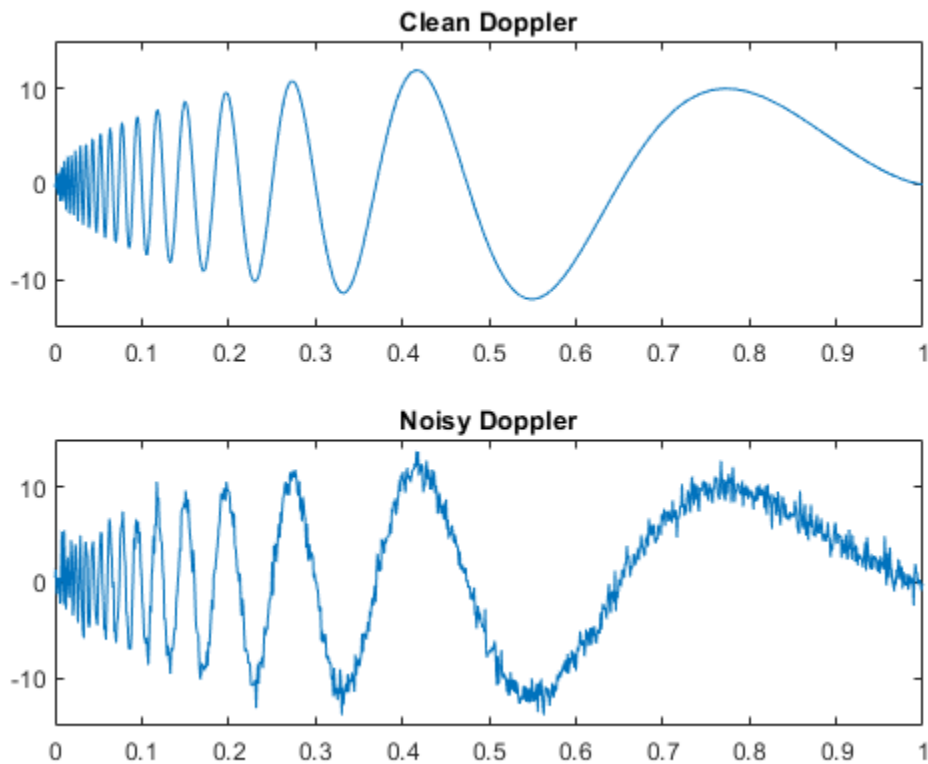
There are six test signals. Generate and plot 2^{10} samples of the third test signal, heavy sine.

```
loc = linspace(0,1,2^10);
x = wnoise(3,10);
plot(loc,x)
title('Heavy Sine')
```



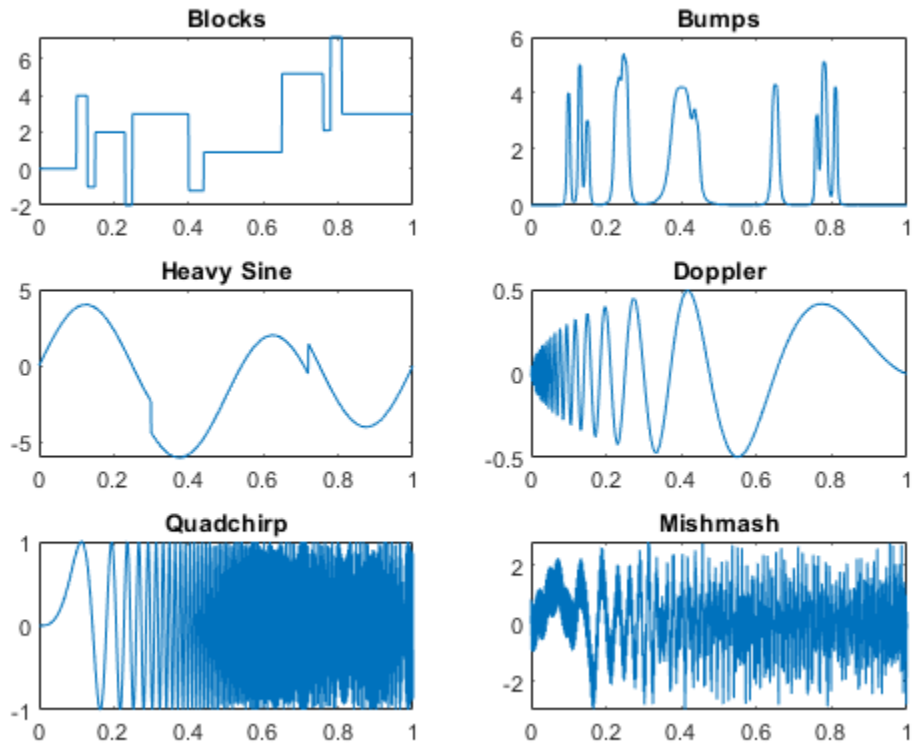
Generate and plot 2^{10} samples of the `doppler` test signal and a noisy version of `doppler` with a square root of the signal-to-noise ratio equal to 7.

```
[x,noisyx] = wnoise('doppler',10,7);  
subplot(2,1,1)  
plot(loc,x)  
title('Clean Doppler')  
ylim([-15 15])  
subplot(2,1,2)  
plot(loc,noisyx)  
title('Noisy Doppler')  
ylim([-15 15])
```

Plot all the test functions.

```
testFunctions = {'Blocks', 'Bumps', 'Heavy Sine', 'Doppler', 'Quadchirp', 'Mishmash'};
for i=1:6
    x = wnoise(lower(testFunctions{i}),10);
    subplot(3,2,i)
    plot(loc,x)
    title(testFunctions{i})
end
```



Input Arguments

fun — Wavelet test function
positive integer | character array

Wavelet test function, specified as one of the values listed here. The six test functions are due to Donoho and Johnstone [1], [2].

- 1 or 'blocks'
- 2 or 'bumps'

- 3 or 'heavy sine'
- 4 or 'doppler'
- 5 or 'quadchirp'
- 6 or 'mishmash'

n — Exponent

positive integer

Exponent used to determine the number of linearly spaced points from 0 to 1 to evaluate the test function, specified as a positive integer. The number of linearly spaced points is 2^n .

sqrtsnr — Square root of SNR

positive real number

Square root of SNR, specified by a positive real number. The test values x are rescaled such that the standard deviation of x is `sqrtsnr`. x_n is equal to x corrupted by additive Gaussian white noise $N(0,1)$ and has an SNR of `sqrtsnr`².

init — Seed

nonnegative integer

Seed used to initialize the random number generator, specified as a nonnegative integer. `init` is used to generate additive Gaussian white noise.

Example: `[a,b] = wnoise(4,10,7,2055415866)`; returns a noisy version of the fourth test signal using the seed `init = 2055415866`.

Output Arguments

x — Test signal

real-valued vector

Test signal, returned as a real-valued vector of length 2^n . x are the values of the test function specified by `fun` evaluated at the 2^n evenly spaced points from 0 to 1. If `sqrtsnr` is set, the standard deviation of x is `sqrtsnr`.

xn — Noisy test signal

real-valued vector

Noisy test signal, returned as a real-valued vector of length 2^n . x_n is x corrupted by additive Gaussian white noise $N(0,1)$ and has an SNR of $\sqrt{2} \text{snr}^2$.

References

- [1] Donoho, D. L., and I. M. Johnstone. "Ideal spatial adaptation by wavelet shrinkage." *Biometrika*. Vol. 81, Issue 3, 1994, pp. 425-455.
- [2] Donoho, D. L., and I. M. Johnstone. "Adapting to unknown smoothness via wavelet shrinkage." *Journal of the American Statistical Association*. Vol. 90, 1995, pp. 1200-1224.

See Also

`wden` | `wdenoise`

Introduced before R2006a

wnoisest

Estimate noise of 1-D wavelet coefficients

Syntax

```
STDC = wnoisest(C,L,S)
STDC = wnoisest(C)
STDC = wnoisest(C)
```

Description

`STDC = wnoisest(C,L,S)` returns estimates of the detail coefficients' standard deviation for levels contained in the input vector `S`. `[C,L]` is the input wavelet decomposition structure (see `wavedec` for more information).

If `C` is a one dimensional cell array, `STDC = wnoisest(C)` returns a vector such that `STDC(k)` is an estimate of the standard deviation of `C{k}`.

If `C` is a numeric array, `STDC = wnoisest(C)` returns a vector such that `STDC(k)` is an estimate of the standard deviation of `C(k,:)`.

The estimator used is Median Absolute Deviation / 0.6745, well suited for zero mean Gaussian white noise in the de-noising one-dimensional model (see `thselect` for more information).

Examples

Estimate Noise Standard Deviation in The Presence of Outliers

Estimate of the noise standard deviation in an $N(0,1)$ white Gaussian noise vector with outliers.

Create an $N(0,1)$ noise vector with 10 randomly-placed outliers.

```
rng default;  
x = randn(1000,1);  
P = randperm(length(x));  
indices = P(1:10);  
x(indices(1:5)) = 10;  
x(indices(6:end)) = -10;
```

Obtain the discrete wavelet transform down to level 2 using the Daubechies' extremal phase wavelet with 3 vanishing moments.

```
[c,l] = wavedec(x,2,'db3');  
stdc = wnoisest(c,l,1:2)
```

```
stdc = 1×2
```

```
    0.9559    1.0556
```

In spite of the outliers, `wnoisest` provides a robust estimate of the standard deviation.

References

Donoho, D.L.; I.M. Johnstone (1994), "Ideal spatial adaptation by wavelet shrinkage," *Biometrika*, vol 81, pp. 425-455.

Donoho, D.L.; I.M. Johnstone (1995), "Adapting to unknown smoothness via wavelet shrinkage via wavelet shrinkage," *JASA*, vol 90, 432, pp. 1200-1224.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

Functions

thselect | wavedec | wden | wdenoise

Apps

Wavelet Signal Denoiser

Introduced before R2006a

wp2wtree

Extract wavelet tree from wavelet packet tree

Syntax

```
T = wp2wtree(T)
```

Description

`wp2wtree` is a one- or two-dimensional wavelet packet analysis function.

`T = wp2wtree(T)` computes the modified wavelet packet tree T corresponding to the wavelet decomposition tree.

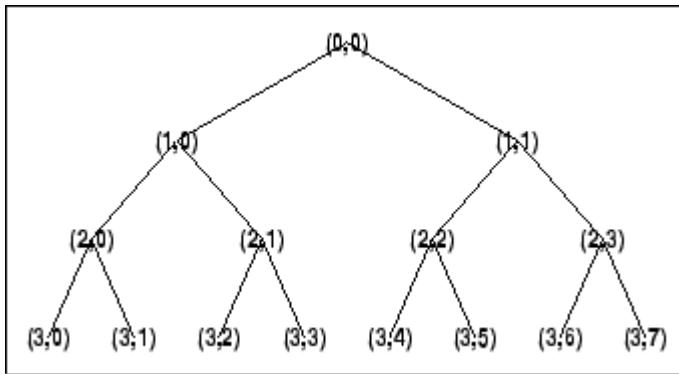
Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load signal.  
load noisdopp; x = noisdopp;
```

```
% Decompose x at depth 3 with db1 wavelet packets  
% using shannon entropy.  
wpt = wpdec(x,3,'db1');
```

```
% Plot wavelet packet tree wpt.  
plot(wpt)
```

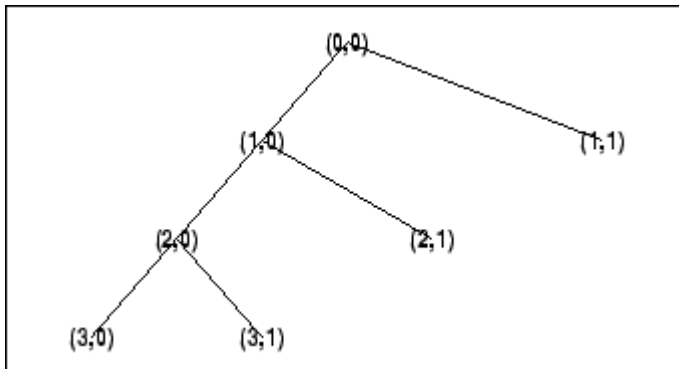



```

% Compute wavelet tree.
wt = wp2wtree(wpt);

% Plot wavelet tree wt.
plot(wt)

```



See Also

wpdec | wpdec2

Introduced before R2006a

wpbmpen

Penalized threshold for wavelet packet denoising

Syntax

`THR = wpbmpen(T, SIGMA, ALPHA)`
`wpbmpen(T, SIGMA, ALPHA, ARG)`

Description

`THR = wpbmpen(T, SIGMA, ALPHA)` returns a global threshold `THR` for denoising. `THR` is obtained by a wavelet packet coefficients selection rule using a penalization method provided by Birgé-Massart.

`T` is a wavelet packet tree corresponding to the wavelet packet decomposition of the signal or image to be denoised.

`SIGMA` is the standard deviation of the zero mean Gaussian white noise in the denoising model (see `wnoiset` for more information).

`ALPHA` is a tuning parameter for the penalty term. It must be a real number greater than 1. The sparsity of the wavelet packet representation of the denoised signal or image grows with `ALPHA`. Typically `ALPHA = 2`.

`THR` minimizes the penalized criterion given by

let t^* be the minimizer of

$$\text{crit}(t) = -\sum(c(k)^2, k \leq t) + 2 * \text{SIGMA}^2 * t * (\text{ALPHA} + \log(n/t))$$

where $c(k)$ are the wavelet packet coefficients sorted in decreasing order of their absolute value and n is the number of coefficients, then $\text{THR} = |c(t^*)|$.

`wpbmpen(T, SIGMA, ALPHA, ARG)` computes the global threshold and, in addition, plots three curves:

- $2 \cdot \text{SIGMA}^2 \cdot t \cdot (\text{ALPHA} + \log(n/t))$
- $\text{sum}(c(k)^2, k \neq t)$
- $\text{crit}(t)$

Examples

```
% Example 1: Signal denoising.
% Load noisy chirp signal.
load nois chir; x = nois chir;

% Perform a wavelet packet decomposition of the signal
% at level 5 using sym6.
wname = 'sym6'; lev = 5;
tree = wpdec(x,lev,wname);

% Estimate the noise standard deviation from the
% detail coefficients at level 1,
% corresponding to the node index 2.
det1 = wpccoef(tree,2);
sigma = median(abs(det1))/0.6745;

% Use wpbmpen for selecting global threshold
% for signal denoising, using the recommended parameter.
alpha = 2;
thr = wpbmpen(tree,sigma,alpha)

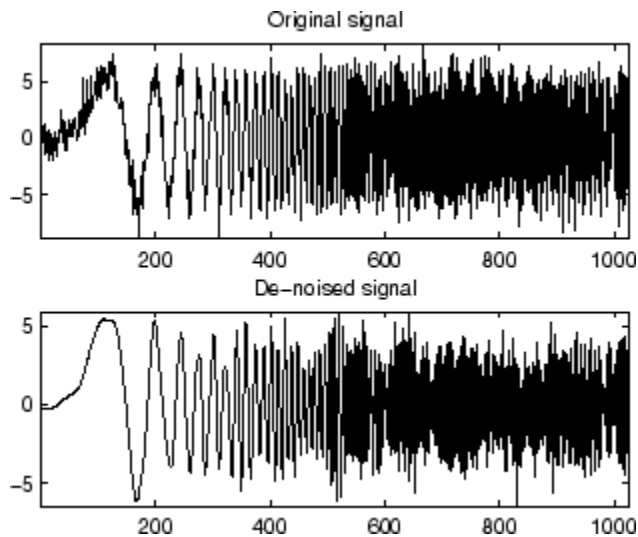
thr =

    4.5740

% Use wpdencmp for denoising the signal using the above
% threshold with soft thresholding and keeping the
% approximation.
keepapp = 1;
xd = wpdencmp(tree,'s','nobest',thr,keepapp);

% Plot original and denoised signals.
figure(1)
subplot(211), plot(x),
title('Original signal')
```

```
subplot(212), plot(xd)
title('De-noised signal')
```



```
% Example 2: Image denoising.
% Load original image.
load noiswom;
nbc = size(map,1);

% Perform a wavelet packet decomposition of the image
% at level 3 using coif2.
wname = 'coif2'; lev = 3;
tree = wpdec2(X,lev,wname);

% Estimate the noise standard deviation from the
% detail coefficients at level 1.
det1 = [wpcoef(tree,2) wpcoef(tree,3) wpcoef(tree,4)];
sigma = median(abs(det1(:)))/0.6745;

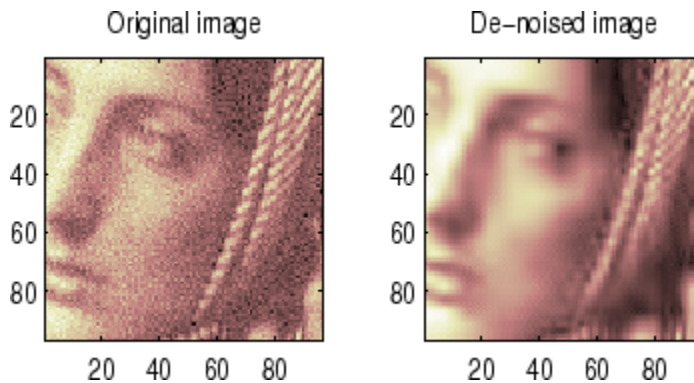
% Use wpbmpen for selecting global threshold
% for image denoising.
alpha = 1.1;
thr = wpbmpen(tree,sigma,alpha)

thr =
```

38.5125

```
% Use wpdencmp for denoising the image using the above
% thresholds with soft thresholding and keeping the
% approximation.
keepapp = 1;
xd = wpdencmp(tree,'s','nobest',thr,keepapp);

% Plot original and denoised images.
figure(2)
colormap(pink(nbc));
subplot(221), image(wcodemat(X,nbc))
title('Original image')
subplot(222), image(wcodemat(xd,nbc))
title('De-noised image')
```



See Also

wbmpen | wden | wdencomp | wdenoise | wpdencmp

Introduced before R2006a

wpcoef

Wavelet packet coefficients

Syntax

```
X = wpcoef(T,N)  
X = wpcoef(T)
```

Description

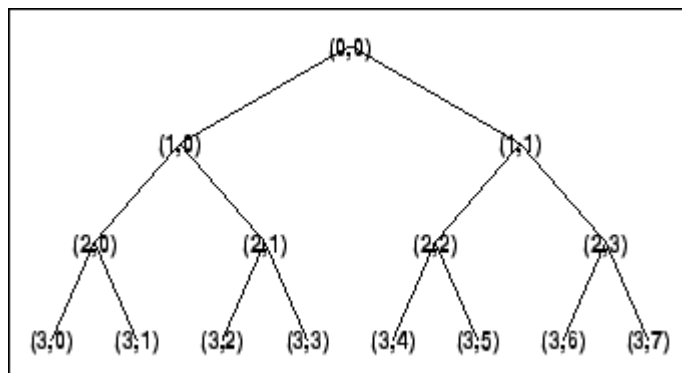
`wpcoef` is a one- or two-dimensional wavelet packet analysis function.

`X = wpcoef(T,N)` returns the coefficients associated with the node `N` of the wavelet packet tree `T`. If `N` doesn't exist, `X = []`;

`X = wpcoef(T)` is equivalent to `X = wpcoef(T,0)`.

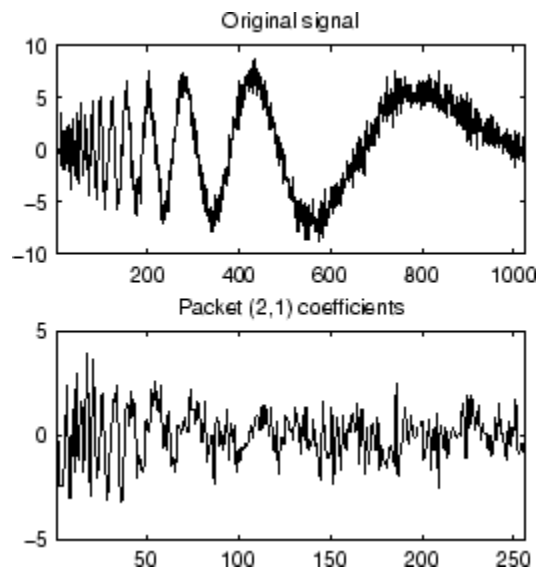
Examples

```
% The current extension mode is zero-padding (see dwtmode).  
  
% Load signal.  
load noisdopp; x = noisdopp;  
  
figure(1); subplot(211);  
plot(x); title('Original signal');  
  
% Decompose x at depth 3 with db1 wavelet packets  
% using Shannon entropy.  
wpt = wpdec(x,3,'db1');  
  
% Plot wavelet packet tree wpt.  
plot(wpt)
```



```
% Read packet (2,1) coefficients.
cfs = wpcoef(wpt,[2 1]);
```

```
figure(1); subplot(212);
plot(cfs); title('Packet (2,1) coefficients');
```



See Also

wpcoef | wpdec | wpdec2 | wprcoef

Topics

“Reconstructing a Signal Approximation from a Node”

Introduced before R2006a

wpcutree

Cut wavelet packet tree

Syntax

```
T = wpcutree(T,L)
T
[T,RN] = wpcutree(T,L)
```

Description

wpcutree is a one- or two-dimensional wavelet packet analysis function.

T = wpcutree(T,L) cuts the tree T at level L.

[T,RN] = wpcutree(T,L) returns the same arguments as above and, in addition, the vector RN contains the indices of the reconstructed nodes.

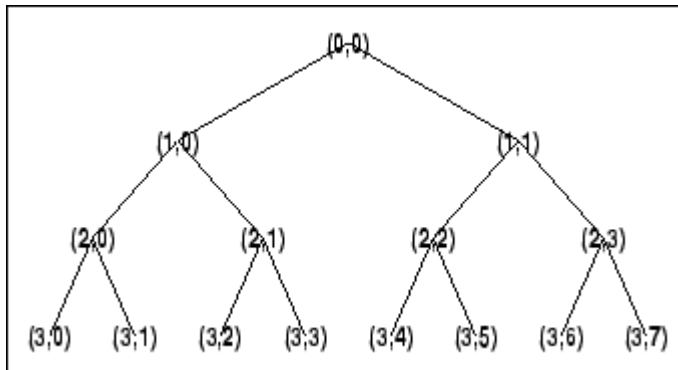
Examples

```
% The current extension mode is zero-padding (see dwtmode).

% Load signal.
load noisdopp; x = noisdopp;

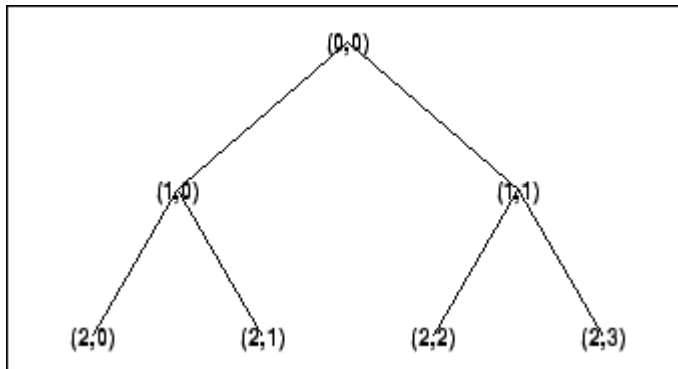
% Decompose x at depth 3 with db1 wavelet packets
% using Shannon entropy.
wpt = wpdec(x,3,'db1');

% Plot wavelet packet tree wpt.
plot(wpt)
```



```
% Cut wavelet packet tree at level 2.  
nwpt = wpcuttree(wpt,2);
```

```
% Plot new wavelet packet tree nwpt.  
plot(nwpt)
```



See Also

wpdec | wpdec2

Introduced before R2006a

wpdec

Wavelet packet decomposition 1-D

Syntax

```
T = wpdec(X,N,wname,E,P)
T = wpdec(X,N,wname)
T = wpdec(X,N,wname,'shannon')
```

Description

wpdec is a one-dimensional wavelet packet analysis function.

$T = \text{wpdec}(X, N, \text{wname}, E, P)$ returns a wavelet packet tree T corresponding to the wavelet packet decomposition of the vector X at level N , using the wavelet specified by wname (see `wfilters` for more information).

$T = \text{wpdec}(X, N, \text{wname})$ is equivalent to $T = \text{wpdec}(X, N, \text{wname}, 'shannon')$.

E is a character vector or string scalar containing the type of entropy and P is an optional parameter depending on the value of T (see `wentropy` for more information).

Entropy Type Name (E)	Parameter (P)	Comments
'shannon'		P is not used.
'log energy'		P is not used.
'threshold'	$0 \leq P$	P is the threshold.
'sure'	$0 \leq P$	P is the threshold.
'norm'	$1 \leq P$	P is the power.
'user'	character vector or string scalar	P is a character vector or string scalar containing the file name of your own entropy function, with a single input X .

Entropy Type Name (E)	Parameter (P)	Comments
FunName	No constraints on P	FunName is any other character vector or string scalar except those used for the previous Entropy Type Names listed above. FunName contains the file name of your own entropy function, with X as input and P as additional parameter to your entropy function.

Note The 'user' option is historical and still kept for compatibility, but it is obsoleted by the last option described in the table above. The FunName option do the same as the 'user' option and in addition gives the possibility to pass a parameter to your own entropy function.

The wavelet packet method is a generalization of wavelet decomposition that offers a richer signal analysis. Wavelet packet atoms are waveforms indexed by three naturally interpreted parameters: position and scale as in wavelet decomposition, and frequency.

For a given orthogonal wavelet function, a library of wavelet packets bases is generated. Each of these bases offers a particular way of coding signals, preserving global energy and reconstructing exact features. The wavelet packets can then be used for numerous expansions of a given signal.

Simple and efficient algorithms exist for both wavelet packets decomposition and optimal decomposition selection. Adaptive filtering algorithms with direct applications in optimal signal coding and data compression can then be produced.

In the orthogonal wavelet decomposition procedure, the generic step splits the approximation coefficients into two parts. After splitting we obtain a vector of approximation coefficients and a vector of detail coefficients, both at a coarser scale. The information lost between two successive approximations is captured in the detail coefficients. The next step consists in splitting the new approximation coefficient vector; successive details are never re-analyzed.

In the corresponding wavelet packets situation, each detail coefficient vector is also decomposed into two parts using the same approach as in approximation vector splitting.

This offers the richest analysis: the complete binary tree is produced in the one-dimensional case or a quaternary tree in the two-dimensional case.

Examples

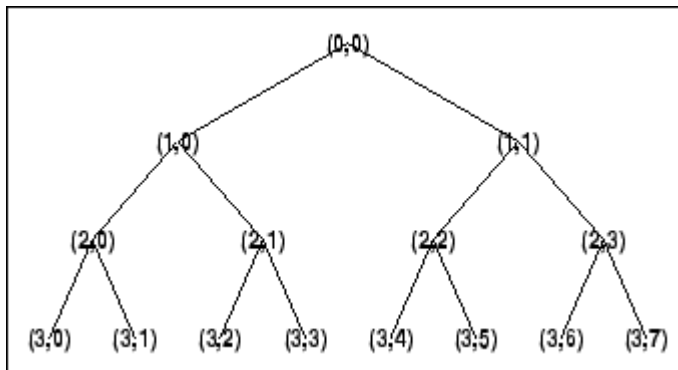
```
% The current extension mode is zero-padding (see dwtmode).

% Load signal.
load noisdopp; x = noisdopp;

% Decompose x at depth 3 with db1 wavelet packets
% using Shannon entropy.
wpt = wpdec(x,3,'db1','shannon');

% The result is the wavelet packet tree wpt.

% Plot wavelet packet tree (binary tree, or tree of order 2).
plot(wpt)
```



Algorithms

The algorithm used for the wavelet packets decomposition follows the same line as the wavelet decomposition process (see `dwt` and `wavedec` for more information).

References

Coifman, R.R.; M.V. Wickerhauser, (1992), "Entropy-based Algorithms for best basis selection," *IEEE Trans. on Inf. Theory*, vol. 38, 2, pp. 713-718.

Meyer, Y. (1993), *Les ondelettes. Algorithmes et applications*, Colin Ed., Paris, 2nd edition. (English translation: *Wavelets: Algorithms and Applications*, SIAM).

Wickerhauser, M.V. (1991), "INRIA lectures on wavelet packet algorithms," *Proceedings ondelettes et paquets d'ondes*, 17-21 June, Rocquencourt, France, pp. 31-99.

Wickerhauser, M.V. (1994), *Adapted wavelet analysis from theory to software algorithms*, A.K. Peters.

See Also

wavedec | waveinfo | wenergy | wpdec | wprec

Topics

"Build Wavelet Tree Objects"

"Examples Using Wavelet Packet Tree Objects"

"Objects in the Wavelet Toolbox Software"

Introduced before R2006a

wpdec2

Wavelet packet decomposition 2-D

Syntax

```
T = wpdec2(X,N,wname,E,P)
T = wpdec2(X,N,wname)
T = wpdec2(X,N,wname,'shannon')
```

Description

wpdec2 is a two-dimensional wavelet packet analysis function.

$T = \text{wpdec2}(X, N, \text{wname}, E, P)$ returns a wavelet packet tree T corresponding to the wavelet packet decomposition of the matrix X , at level N , with the specified wavelet wname (see `wfilters` for more information).

$T = \text{wpdec2}(X, N, \text{wname})$ is equivalent to $T = \text{wpdec2}(X, N, \text{wname}, 'shannon')$.

E is a character vector or string scalar containing the type of entropy and P is an optional parameter depending on the value of T (see `wentropy` for more information).

Entropy Type Name (E)	Parameter (P)	Comments
'shannon'		P is not used.
'log energy'		P is not used.
'threshold'	$0 \leq P$	P is the threshold.
'sure'	$0 \leq P$	P is the threshold.
'norm'	$1 \leq P$	P is the power.
'user'	Character vector or string scalar	P is a character vector or string scalar containing the file name of your own entropy function, with a single input X .

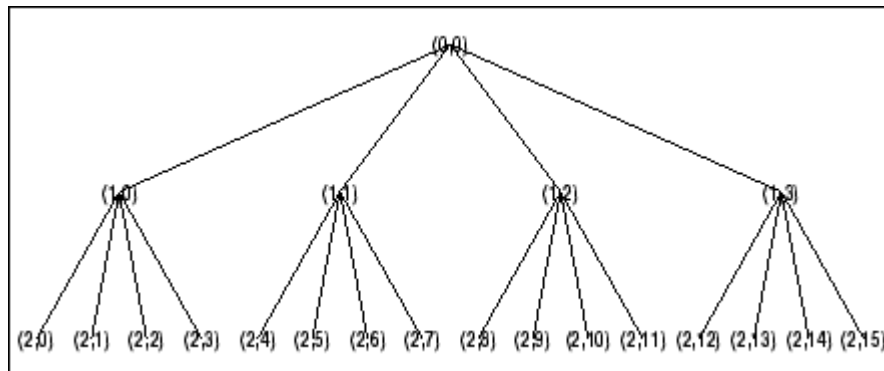
Entropy Type Name (E)	Parameter (P)	Comments
FunName	No constraints on P	FunName is any other character vector or string scalar except those used for the previous Entropy Type Names listed above. FunName contains the file name of your own entropy function, with X as input and P as additional parameter to your entropy function.

Note The 'user' option is historical and still kept for compatibility, but it is obsoleted by the last option described in the preceding table. The FunName option does the same as the 'user' option and in addition, allows you to pass a parameter to your own entropy function.

See `wpdec` for a more complete description of the wavelet packet decomposition.

Examples

```
% The current extension mode is zero-padding (see dwtmode).  
  
% Load image.  
load tire  
% X contains the loaded image.  
  
% For an image the decomposition is performed using:  
t = wpdec2(X,2,'db1');  
% The default entropy is shannon.  
  
% Plot wavelet packet tree  
% (quarternary tree, or tree of order 4).  
plot(t)
```

Tips

When X represents an indexed image, X is an m -by- n matrix. When X represents a truecolor image, it is an m -by- n -by-3 array, where each m -by- n matrix represents a red, green, or blue color plane concatenated along the third dimension.

For more information on image formats, see the `image` and `imfinfo` reference pages.

Algorithms

The algorithm used for the wavelet packets decomposition follows the same line as the wavelet decomposition process (see `dwt2` and `wavedec2` for more information).

References

Coifman, R.R.; M.V. Wickerhauser (1992), "Entropy-based algorithms for best basis selection," *IEEE Trans. on Inf. Theory*, vol. 38, 2, pp. 713-718.

Meyer, Y. (1993), *Les ondelettes. Algorithmes et applications*, Colin Ed., Paris, 2nd edition. (English translation: *Wavelets: Algorithms and Applications*, SIAM).

Wickerhauser, M.V. (1991), "INRIA lectures on wavelet packet algorithms," *Proceedings ondelettes et paquets d'ondes*, 17-21 June, Rocquencourt, France, pp. 31-99.

Wickerhauser, M.V. (1994), *Adapted wavelet analysis from theory to software Algorithms*, A.K. Peters.

See Also

wavedec2 | waveinfo | wenergy | wpdec | wprec2

Topics

“Build Wavelet Tree Objects”

“Examples Using Wavelet Packet Tree Objects”

“Objects in the Wavelet Toolbox Software”

Introduced before R2006a

wpdencmp

Denoising or compression using wavelet packets

`wpdencmp` performs a denoising or compression process of a signal or image using wavelet packets. The ideas and procedures for denoising and compression using either wavelet or wavelet packet decompositions are the same. See `wdenoise` or `wdencomp` for more information.

Syntax

```
[xd,treed,perf0,perfl2] = wpdencmp(x,sorh,n,wname,crit,par,keepapp)
[ ___ ] = wpdencmp(tree,sorh,crit,par,keepapp)
```

Description

`[xd,treed,perf0,perfl2] = wpdencmp(x,sorh,n,wname,crit,par,keepapp)` returns a denoised or compressed version `xd` of the input data `x` obtained by wavelet packet coefficient thresholding. `wpdencmp` also returns the wavelet packet best tree decomposition `treed` of `xd` (see `besttree` for more information), and the L^2 energy recovery and compression scores in percentages as `perfl2` and `perf0`, respectively.

`[___] = wpdencmp(tree,sorh,crit,par,keepapp)` uses the wavelet packet decomposition `tree` of the data to be denoised or compressed.

Examples

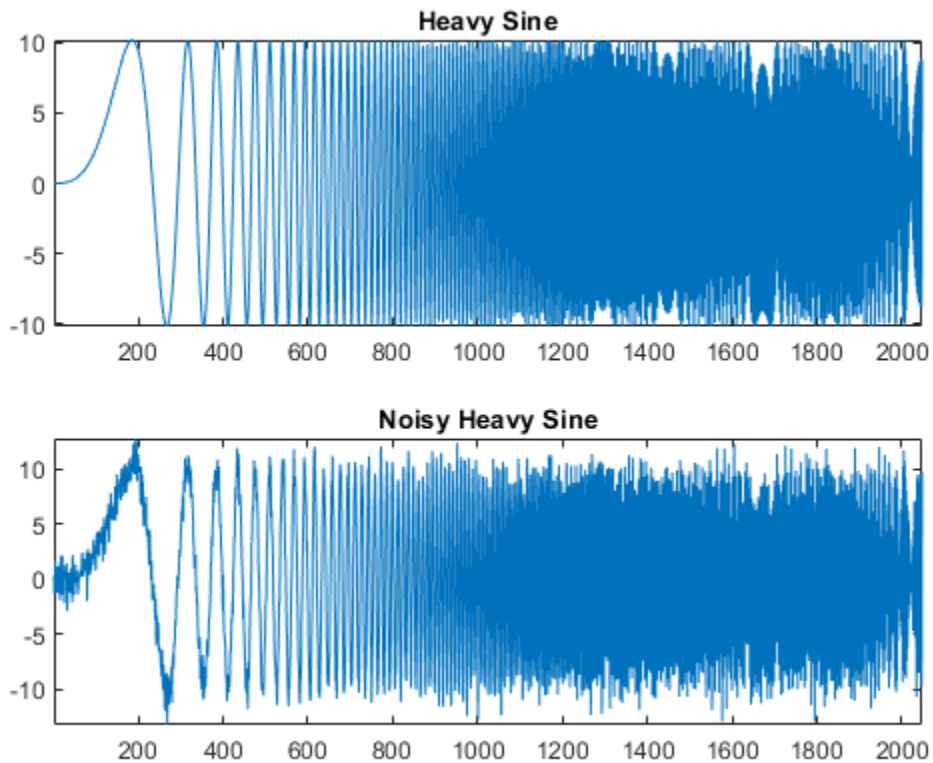
1-D Denoising Using Wavelet Packets

This example shows how to denoise using wavelet packets.

Use `wnoise` to generate the heavy sine signal and a noisy version.

```
init = 1000;
[xref,x] = wnoise(5,11,7,init);
```

```
figure
subplot(2,1,1)
plot(xref)
axis tight
title('Heavy Sine')
subplot(2,1,2)
plot(x)
axis tight
title('Noisy Heavy Sine')
```

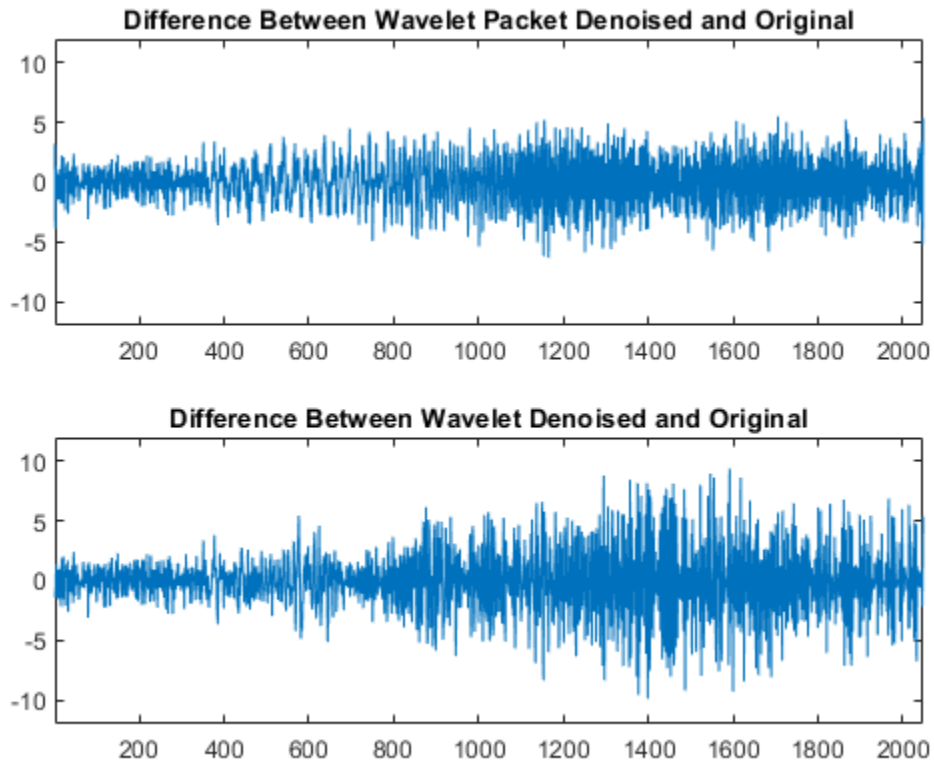


Denoise the noisy signal using a four-level wavelet packet decomposition. Use the order 4 Daubechies least asymmetric wavelet.

```
n = length(x);
thr = sqrt(2*log(n*log(n)/log(2)));
xwpc = wpdencomp(x, 's', 4, 'sym4', 'sure', thr, 1);
```

Compare with a wavelet-based denoising result. Use `wdenoise` with comparable input arguments. Plot the differences between the two denoised signals and original signal.

```
xwd = wdenoise(x, 4, 'Wavelet', 'sym4', 'DenoisingMethod', 'UniversalThreshold', 'ThresholdR
figure
subplot(2,1,1)
plot(x-xwpc)
axis tight
ylim([-12 12])
title('Difference Between Wavelet Packet Denoised and Original')
subplot(2,1,2)
plot(x-xwd)
axis tight
ylim([-12 12])
title('Difference Between Wavelet Denoised and Original')
```

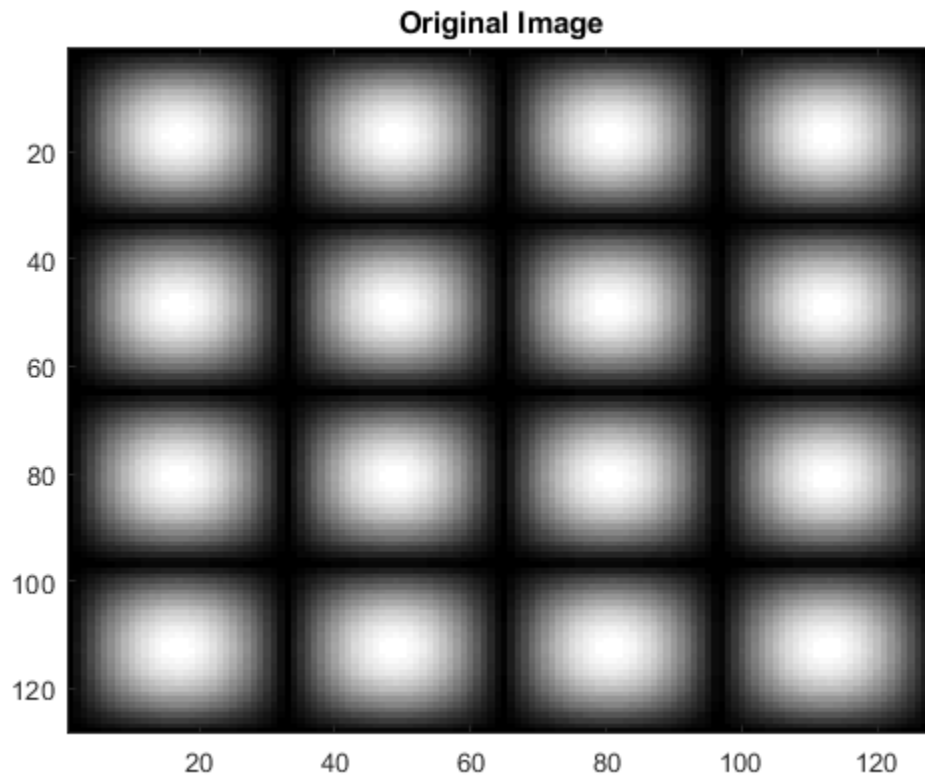


2-D Denoising Using Wavelet Packets

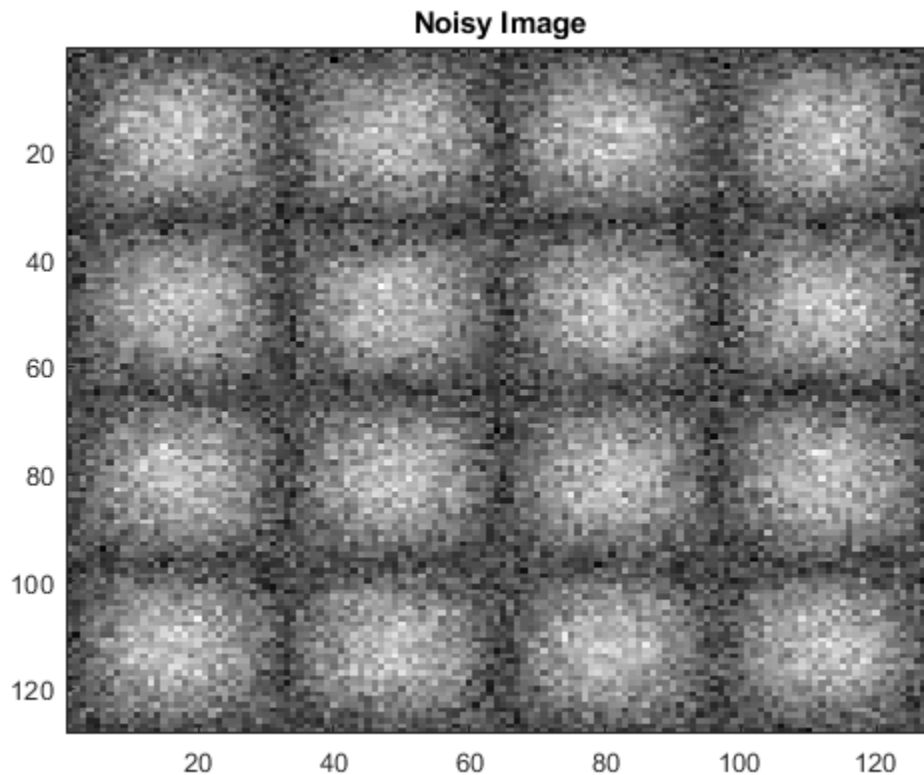
This example shows how to denoise an image using wavelet packets.

Load an image and generate a noisy copy. For reproducibility set the random seed.

```
rng default
load sinsin
x = X/18 + randn(size(X));
imagesc(X)
colormap(gray)
title('Original Image')
```

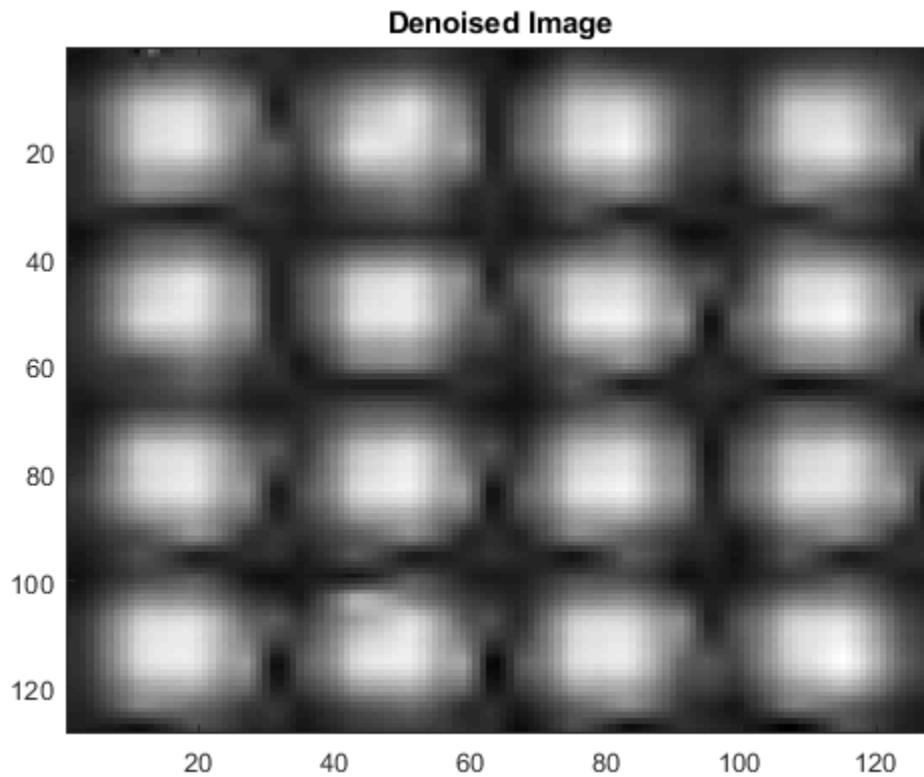


```
figure
imagesc(x)
colormap(gray)
title('Noisy Image')
```



Denoise the noisy image using wavelet packet decomposition. Use `ddencmp` to determine denoising parameters. Do a three-level decomposition with the order 4 Daubechies least asymmetric wavelet.

```
[thr,sorh,keepapp,crit] = ddencmp('den','wp',x);  
xd = wpdencmp(x,sorh,3,'sym4',crit,thr,keepapp);  
figure  
imagesc(xd)  
colormap(gray)  
title('Denoised Image')
```

1-D Compression Using Wavelet Packets

This example shows how to compress a 1-D signal using wavelet packets.

Load a signal. Use `ddencmp` to determine compression values for that signal.

```
load sumlichr
x = sumlichr;
[thr,sorh,keepapp,crit] = ddencmp('cmp','wp',x)

thr = 0.5193
```

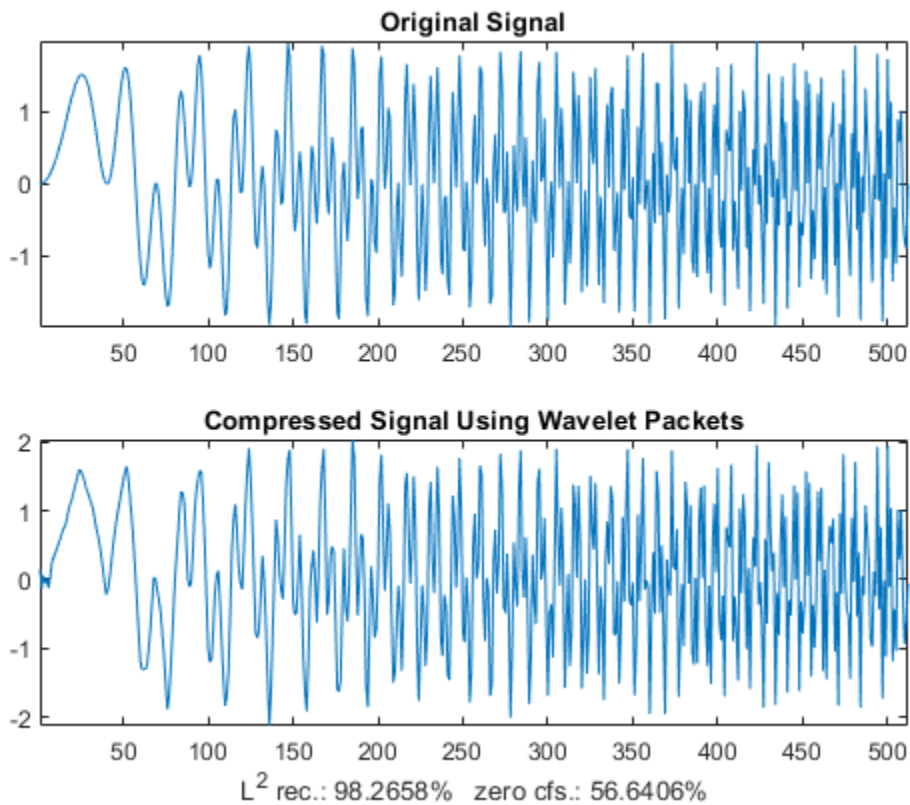
```
sorh =  
'h'  
  
keepapp = 1  
  
crit =  
'threshold'
```

Compress the signal using global thresholding with threshold best basis. Use the order 4 Daubechies least asymmetric wavelet and do a three-level wavelet packet decomposition.

```
[xc,wpt,perf0,perfl2] = wpdencmp(x,sorh,3,'sym4',crit,thr,keepapp);
```

Compare the original signal with the compressed version.

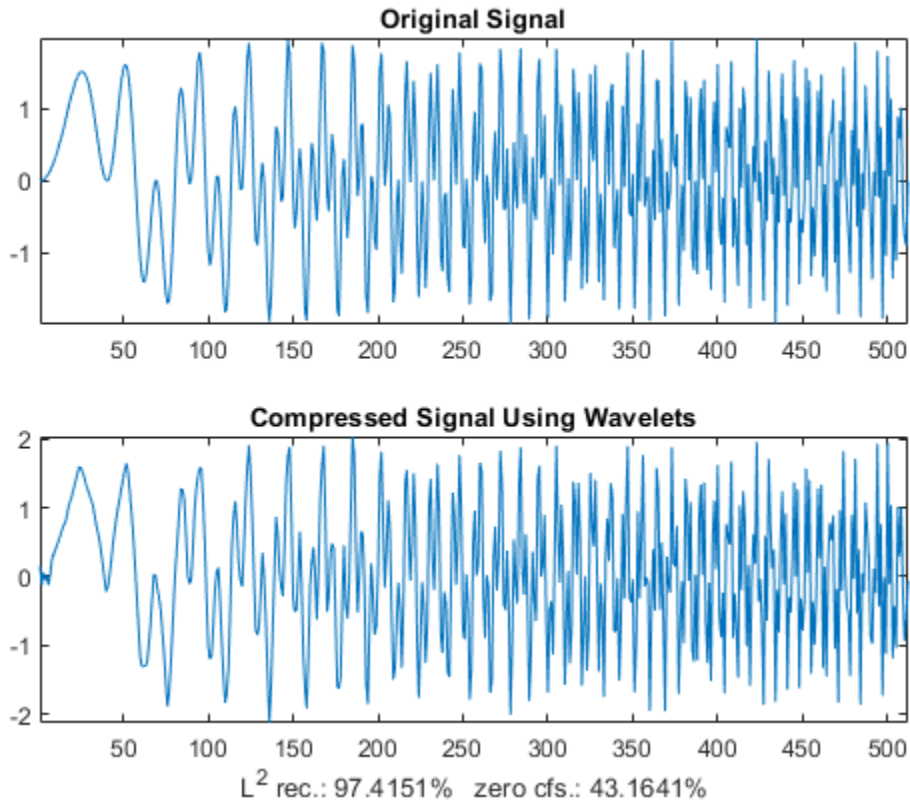
```
subplot(2,1,1)  
plot(x)  
title('Original Signal')  
axis tight  
subplot(2,1,2)  
plot(xc)  
xlabel(['L^2 rec.: ',num2str(perfl2),'% zero cfs.: ',num2str(perf0),'%'])  
title('Compressed Signal Using Wavelet Packets')  
axis tight
```



Compress the signal again, but this do a three-level wavelet decomposition. Keep all the other parameters the same.

```
[thr,sorh,keepapp] = ddencmp('cmp','wv',x);
[xcwv,~,~,perf0wv,perf12wv] = wdencmp('gbl',x,'sym4',3,thr,sorh,keepapp);
figure
subplot(2,1,1)
plot(x)
title('Original Signal')
axis tight
subplot(2,1,2)
plot(xc)
xlabel(['L^2 rec.: ',num2str(perf12wv),'% zero cfs.: ',num2str(perf0wv),'%'])
```

```
title('Compressed Signal Using Wavelets')  
axis tight
```



A larger fraction of coefficients are set equal to 0 when compressing using a wavelet packet decomposition.

Input Arguments

x — Input data

real-valued vector or matrix

Input data to denoise or compress, specified by a real-valued vector or matrix.

Data Types: double

tree — Wavelet packet decomposition

wavelet packet decomposition

Wavelet packet decomposition of the data to be denoised or compressed, specified as a wavelet packet tree. See `wpdec` and `wpdec2` for more information.

sorh — Type of thresholding

's' | 'h'

Type of thresholding to perform:

- 's' — Soft thresholding
- 'h' — Hard thresholding

See `wthresh` for more information.

n — Wavelet packet decomposition level

positive integer

Wavelet packet decomposition level, specified as a positive integer.

wname — Name of wavelet

character vector | string scalar

Name of wavelet, specified as a character vector or string scalar, to use for denoising. See `wavemngr` for more information.

crit — Entropy type

'shannon' | 'log energy' | 'threshold' | 'sure' | 'norm' | 'user' | ...

Entropy type, specified as one of the following:

Entropy Type (crit)	Threshold Parameter (par)	Comments
'shannon'		par is not used.
'log energy'		par is not used.
'threshold'	$0 \leq \text{par}$	par is the threshold.
'sure'	$0 \leq \text{par}$	par is the threshold.

Entropy Type (<i>crit</i>)	Threshold Parameter (<i>par</i>)	Comments
'norm'	$1 \leq \text{par}$	<i>par</i> is the power.
'user'	Character vector	<i>par</i> is a character vector containing the file name of your own entropy function, with a single input <i>x</i> .
' <i>FunName</i> '	No constraints on <i>par</i>	<i>FunName</i> is any character vector other than the previous entropy types listed. <i>FunName</i> contains the file name of your own entropy function, with <i>x</i> as input and <i>par</i> as an additional parameter to your entropy function.

crit and threshold parameter *par* together define the entropy criterion used to determine the best decomposition. See `wentropy` for more information.

If *crit* = 'nobest', no optimization is done, and the current decomposition is thresholded.

par — Threshold parameter

real number | character vector | string scalar

Threshold parameter, specified by a real number, character vector, or string scalar. *par* and the entropy type *crit* together define the entropy criterion used to determine the best decomposition. See `wentropy` for more information.

Data Types: double

keepapp — Threshold approximation setting

0 | 1

Threshold approximation setting, specified as either 0 or 1. If *keepapp* = 1, the approximation coefficients cannot be thresholded. If *keepapp* = 0, the approximation coefficients can be thresholded.

Data Types: double

Output Arguments

xd — Denoised or compressed data

real-valued vector or matrix

Denoised or compressed data, returned as a real-valued vector or matrix. `xd` and `x` have the same dimensions.

treed — Wavelet packet best tree decomposition

wavelet packet tree

Wavelet packet best tree decomposition of `xd`, returned as a wavelet packet tree.

perf0 — Compression score

real number

Compression score, returned as a real number. `perf0` is the percentage of thresholded coefficients that are equal to 0.

perf12 — L^2 energy recovery

real number

L^2 energy recovery, returned as a real number. `perf12` is equal to

$100 \times \left(\frac{\text{vector-norm of wavelet packet coefficients of } xd}{\text{vector-norm of wavelet packet coefficients of } x} \right)^2$. If `x` is a one-dimensional signal

$$\frac{100 \|xd\|^2}{\|x\|^2}.$$

and `wname` an orthogonal wavelet, `perf12` simplifies to

References

- [1] Antoniadis, A., and G. Oppenheim, eds. *Wavelets and Statistics*. Lecture Notes in Statistics. New York: Springer Verlag, 1995.
- [2] Coifman, R. R., and M. V. Wickerhauser. "Entropy-Based Algorithms for Best Basis Selection." *IEEE Transactions on Information Theory*. Vol. 38, Number 2, 1992, pp. 713-718.
- [3] DeVore, R. A., B. Jawerth, and B. J. Lucier. "Image Compression Through Wavelet Transform Coding." *IEEE Transactions on Information Theory*. Vol. 38, Number 2, 1992, pp. 719-746.

- [4] Donoho, D. L. "Progress in Wavelet Analysis and WVD: A Ten Minute Tour." *Progress in Wavelet Analysis and Applications* (Y. Meyer, and S. Roques, eds.). Gif-sur-Yvette: Editions Frontières, 1993.
- [5] Donoho, D. L., and I. M. Johnstone. "Ideal Spatial Adaptation by Wavelet Shrinkage." *Biometrika*. Vol. 81, 1994, pp. 425-455.
- [6] Donoho, D. L., I. M. Johnstone, G. Kerkyacharian, and D. Picard. "Wavelet Shrinkage: Asymptopia?" *Journal of the Royal Statistical Society, series B*. Vol. 57, Number 2, 1995, pp. 301-369.

See Also

Functions

besttree | ddencomp | wden | wdencomp | wdenoise | wenergy | wentropy | wpbopen | wpdec | wpdec2 | wthresh

Apps

Wavelet Signal Denoiser

Introduced before R2006a

wpfun

Wavelet packet functions

Syntax

```
[WPWS,X] = wpfun('wname',NUM,PREC)
[WPWS,X] = wpfun('wname',NUM)
[WPWS,X] = wpfun('wname',NUM,7)
```

Description

wpfun is a wavelet packet analysis function.

`[WPWS,X] = wpfun('wname',NUM,PREC)` computes the wavelet packets for a wavelet '*wname*' (see `wfilters` for more information), on dyadic intervals of length $2^{-\text{PREC}}$.

PREC must be a positive integer. Output matrix WPWS contains the W functions of index from 0 to NUM, stored row-wise as $[W_0; W_1; \dots; W_{\text{NUM}}]$. Output vector X is the corresponding common X-grid vector.

```
[WPWS,X] = wpfun('wname',NUM) is equivalent to
[WPWS,X] = wpfun('wname',NUM,7).
```

The computation scheme for wavelet packets generation is easy when using an orthogonal wavelet. We start with the two filters of length $2N$, denoted $h(n)$ and $g(n)$, corresponding to the wavelet.

Now by induction let us define the following sequence of functions $(W_n(x), n = 0,1,2,\dots)$ by

$$W_{2n}(x) = \sqrt{2} \sum_{k=0, \dots, 2N-1} h(k)W_n(2x-k)$$

$$W_{2n+1}(x) = \sqrt{2} \sum_{k=0, \dots, 2N-1} g(k)W_n(2x-k)$$

where $W_0(x) = \phi(x)$ is the scaling function and $W_1(x) = \psi(x)$ is the wavelet function.

For example for the Haar wavelet we have

$$N = 1, h(0) = h(1) = \frac{1}{\sqrt{2}}$$

and

$$g(0) = -g(1) = \frac{1}{\sqrt{2}}$$

The equations become

$$W_{2n}(x) = W_n(2x) + W_n(2x - 1)$$

and

$$(W_{2n+1}(x) = W_n(2x) - W_n(2x - 1))$$

$W_0(x) = \phi(x)$ is the haar scaling function and $W_1(x) = \psi(x)$ is the haar wavelet, both supported in $[0,1]$.

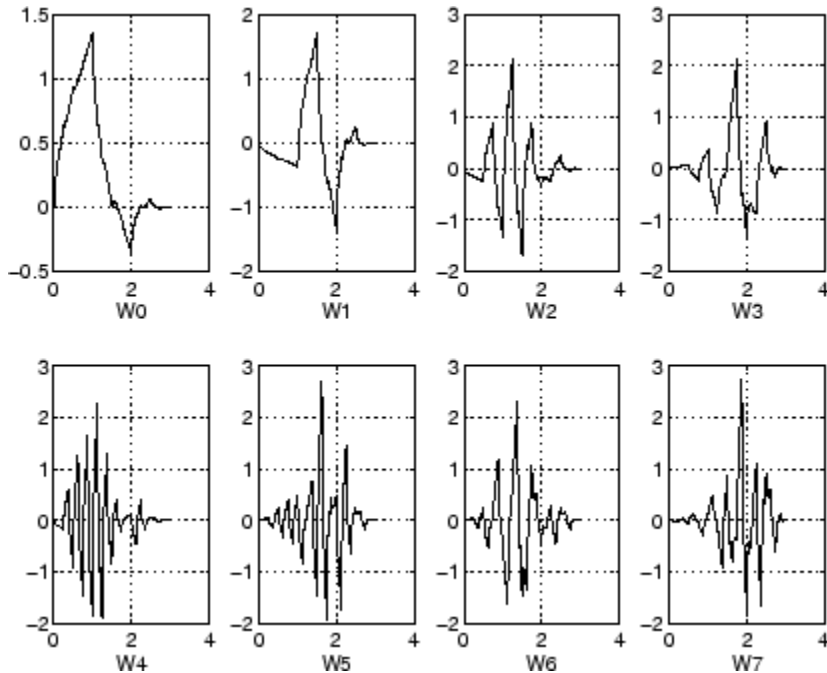
Then we can obtain W_{2^n} by adding two $1/2$ -scaled versions of W_n with distinct supports $[0,1/2]$ and $[1/2,1]$, and obtain $W_{2^{n+1}}$ by subtracting the same versions of W_n .

Starting from more regular original wavelets, using a similar construction, we obtain smoothed versions of this system of W -functions, all with support in the interval $[0, 2N-1]$.

Examples

```
% Compute the db2 Wn functions for n = 0 to 7, generating
% the db2 wavelet packets.
[wp,x] = wfun('db2',7);

% Using some plotting commands,
% the following figure is generated.
```



References

Coifman, R.R.; M.V. Wickerhauser (1992), "Entropy-based Algorithms for best basis selection," *IEEE Trans. on Inf. Theory*, vol. 38, 2, pp. 713-718.

Meyer, Y. (1993), *Les ondelettes. Algorithmes et applications*, Colin Ed., Paris, 2nd edition. (English translation: *Wavelets: Algorithms and applications*, SIAM).

Wickerhauser, M.V. (1991), "INRIA lectures on wavelet packet algorithms," *Proceedings ondelettes et paquets d'ondes*, 17-21 June, Rocquencourt, France, pp. 31-99.

Wickerhauser, M.V. (1994), *Adapted wavelet analysis from theory to software algorithms*, A.K. Peters.

See Also

wavefun | waveinfo

Introduced before R2006a

wpjoin

Recompose wavelet packet

Syntax

```
T = wpjoin(T,N)
[T,X] = wpjoin(T,N)
T = wpjoin(T)
T = wpjoin(T,0)
[T,X] = wpjoin(T)
[T,X] = wpjoin(T,0)
```

Description

wpjoin is a one- or two-dimensional wavelet packet analysis function.

wpjoin updates the wavelet packet tree after the recomposition of a node.

The nodes are numbered from left to right and from top to bottom. The root index is 0.

$T = \text{wpjoin}(T,N)$ returns the modified wavelet packet tree T corresponding to a recomposition of the node N .

$[T,X] = \text{wpjoin}(T,N)$ also returns the coefficients of the node.

$T = \text{wpjoin}(T)$ is equivalent to $T = \text{wpjoin}(T,0)$.

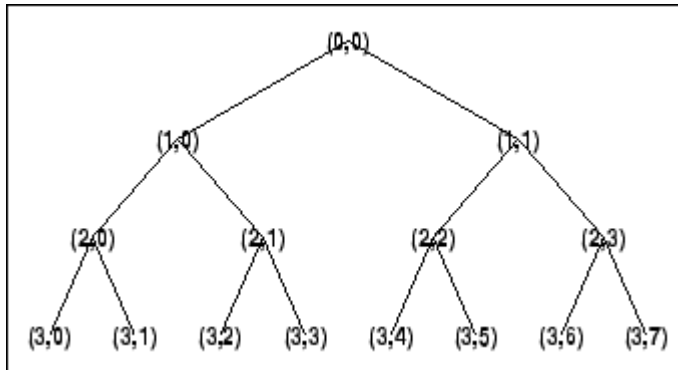
$[T,X] = \text{wpjoin}(T)$ is equivalent to $[T,X] = \text{wpjoin}(T,0)$.

Examples

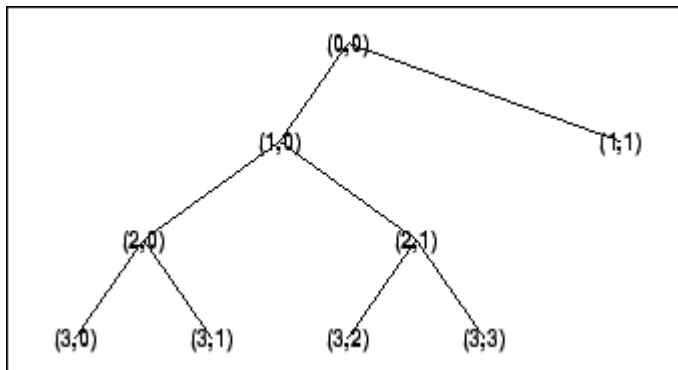
```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load signal.
load noisdopp; x = noisdopp;
```

```
% Decompose x at depth 3 with db1 wavelet packets.  
wpt = wpdec(x,3,'db1');  
  
% Plot wavelet packet tree wpt.  
plot(wpt)
```



```
% Recompose packet (1,1) or 2  
wpt = wpjoin(wpt,[1 1]);  
  
% Plot wavelet packet tree wpt.  
plot(wpt)
```



See Also

[wpdec](#) | [wpdec2](#) | [wpsplt](#)

Introduced before R2006a

wprcoef

Reconstruct wavelet packet coefficients

Syntax

```
X = wprcoef(T,N)
X = wprcoef(T)
X = wprcoef(T,θ)
```

Description

wprcoef is a one- or two-dimensional wavelet packet analysis function.

$X = \text{wprcoef}(T,N)$ computes reconstructed coefficients of the node N of the wavelet packet tree T .

$X = \text{wprcoef}(T)$ is equivalent to $X = \text{wprcoef}(T,\theta)$.

Examples

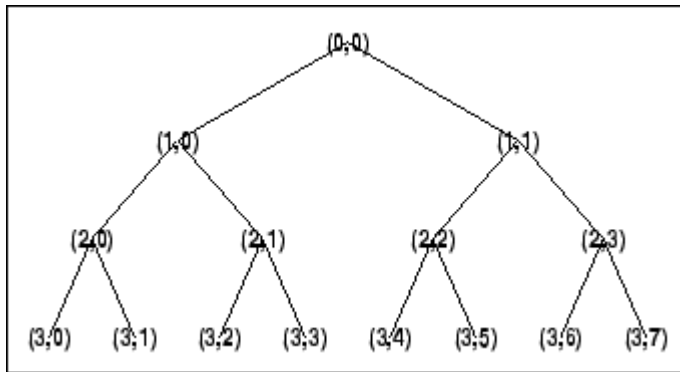
```
% The current extension mode is zero-padding (see dwtmode)

% Load signal.
load noisdopp; x = noisdopp;

figure(1); subplot(211);
plot(x); title('Original signal');

% Decompose x at depth 3 with db1 wavelet packets
% using Shannon entropy.
t = wptdec(x,3,'db1','shannon');

% Plot wavelet packet tree.
plot(t)
```

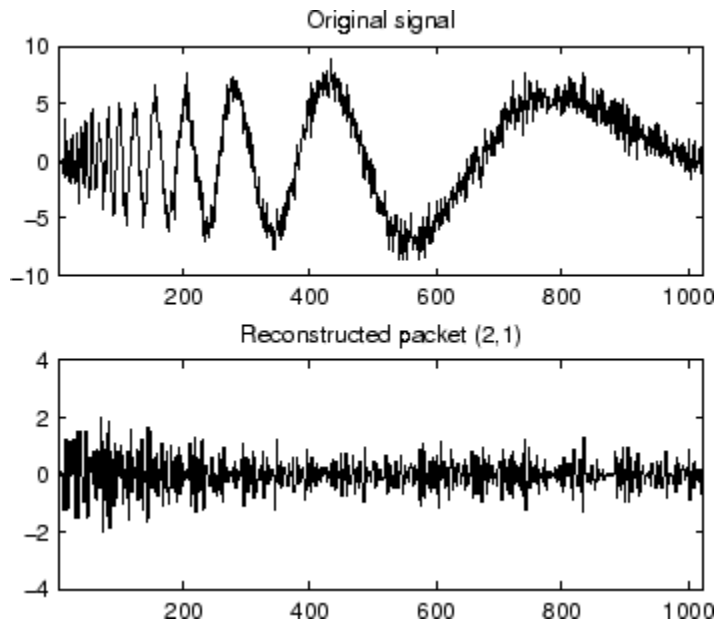



```

% Reconstruct packet (2,1).
rcfs = wprcoef(t,[2 1]);

figure(1); subplot(212);
plot(rcfs); title('Reconstructed packet (2,1)');

```



See Also

wpdec | wpdec2 | wprec | wprec2

Topics

“Reconstructing a Signal Approximation from a Node”

Introduced before R2006a

wprec

Wavelet packet reconstruction 1-D

Syntax

```
X = wprec(T)
wprec(wpdec(X, 'wname'))
```

Description

wprec is a one-dimensional wavelet packet analysis function.

$X = \text{wprec}(T)$ returns the reconstructed vector X corresponding to a wavelet packet tree T .

wprec is the inverse function of wpdec in the sense that the abstract statement $\text{wprec}(\text{wpdec}(X, \text{'wname'}))$ would give back X .

See Also

wpdec | wpdec2 | wpjoin | wprec2 | wpsplt

Introduced before R2006a

wprec2

Wavelet packet reconstruction 2-D

Syntax

```
X = wprec2(T)
wprec2(wpdec2(X, 'wname'))
```

Description

wprec2 is a two-dimensional wavelet packet analysis function.

$X = \text{wprec2}(T)$ returns the reconstructed matrix X corresponding to a wavelet packet tree T .

wprec2 is the inverse function of wpdec2 in the sense that the abstract statement $\text{wprec2}(\text{wpdec2}(X, \text{'wname'}))$ would give back X .

Tips

If T is obtained from an indexed image analysis or a truecolor image analysis, X is an m -by- n matrix or an m -by- n -by-3 array, respectively.

For more information on image formats, see the `image` and `imfinfo` reference pages.

See Also

`wprec` | `wprec2` | `wpjoin` | `wprec` | `wpsplt`

Introduced before R2006a

wpspectrum

Wavelet packet spectrum

Syntax

```
[SPEC, TIMES, FREQ] = wpspectrum(WPT, Fs)
[...] = wpspectrum(WPT, Fs, 'plot')
[... , TNF0] = wpspectrum(...)
```

Description

[SPEC, TIMES, FREQ] = wpspectrum(WPT, Fs) returns a matrix of wavelet packet spectrum estimates, SPEC, for the binary wavelet packet tree object, WPT. Fs is the sampling frequency in Hertz. SPEC is a 2^J -by- N matrix where J is the level of the wavelet packet transform and N is the length of the time series. TIMES is a 1-by- N vector of times and FREQ is a 1-by- 2^J vector of frequencies.

[...] = wpspectrum(WPT, Fs, 'plot') displays the wavelet packet spectrum.

[... , TNF0] = wpspectrum(...) returns the terminal nodes of the wavelet packet tree in frequency order.

Input Arguments

WPT

WPT is a binary wavelet packet tree of class `wptree`.

Fs

Sampling frequency in Hertz as a scalar of class `double`.

Default: 1

plot

The character vector 'plot' displays the wavelet packet spectrum. Enter 'plot' after *F_s* to produce a plot of the wavelet packet spectrum.

Output Arguments

SPEC

Wavelet packet spectrum. SPEC is a 2^J -by- N matrix where J is the level of the wavelet packet transform and N is the length of node 0 in the wavelet packet tree object.

The frequency spacing between the rows of SPEC is $F_s/2^{J+1}$.

TIMES

Time vector. TIMES is a vector of times in seconds equal in length to node 0 of the wavelet packet tree object. The time spacing between elements is $1/F_s$.

FREQ

Frequency vector. FREQ is a vector of frequencies of length 2^J where J is the level of the wavelet packet tree object. The frequency spacing in FREQ is $F_s/2^{J+1}$.

TNFO

Terminal nodes. TNFO is a vector of the terminal nodes of the wavelet packet tree object in frequency order.

Examples

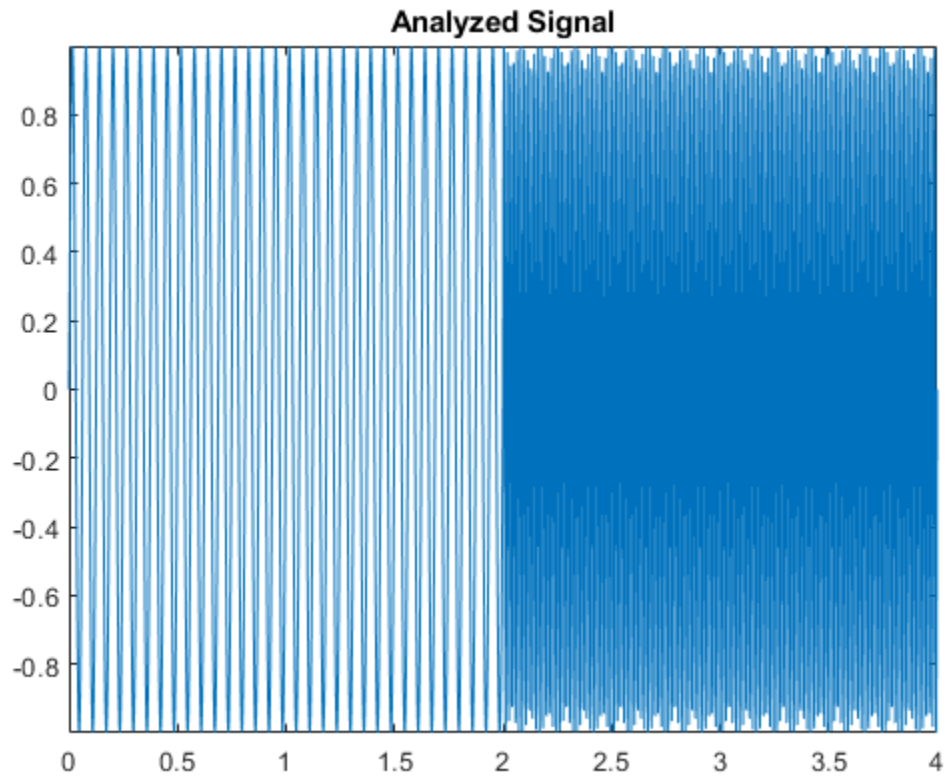
Wavelet Packet Spectrum for Sinusoids

This example shows wavelet packet spectrum for signal consisting of two sinusoids with disjoint support.

Define wavelet.

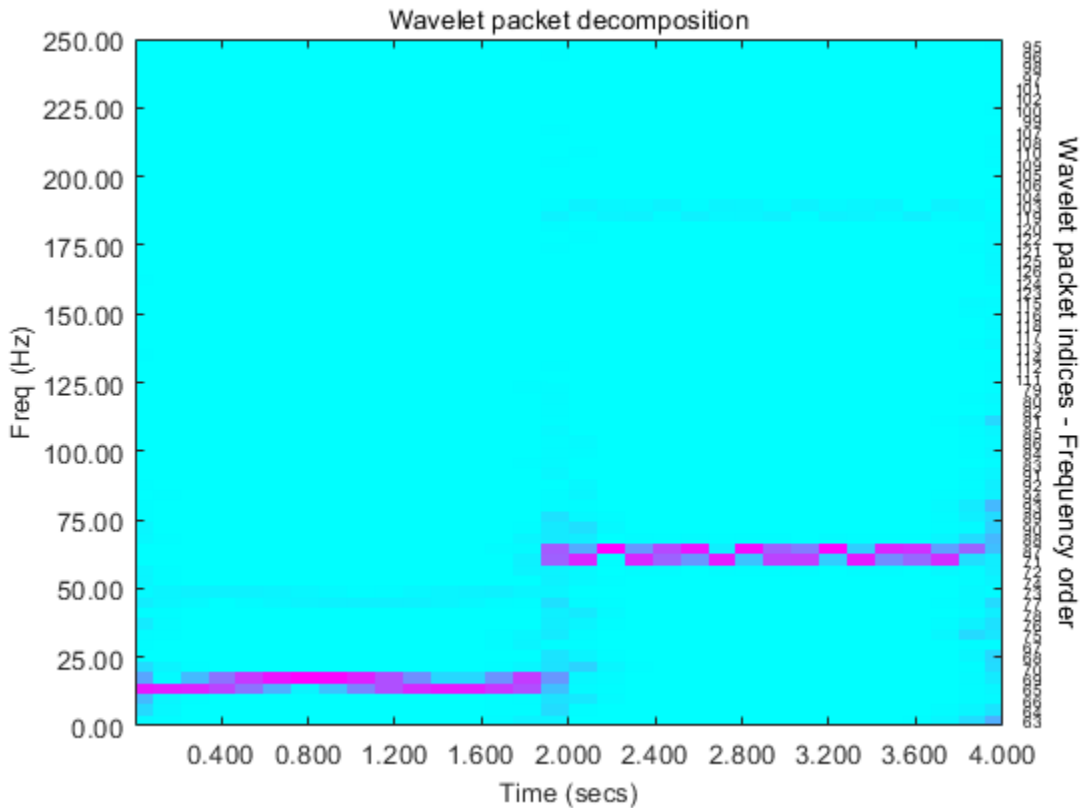
```
fs = 500;  
t = 0:1/fs:4;
```

```
y = sin(32*pi*t).*(t<2) + sin(128*pi*t).*(t>=2);  
plot(t,y);  
axis tight  
title('Analyzed Signal');
```



Define wavelet packet spectrum.

```
level = 6;  
wpt = wpdec(y, level, 'sym6');  
figure;  
[S,T,F] = wpspectrum(wpt, fs, 'plot');
```



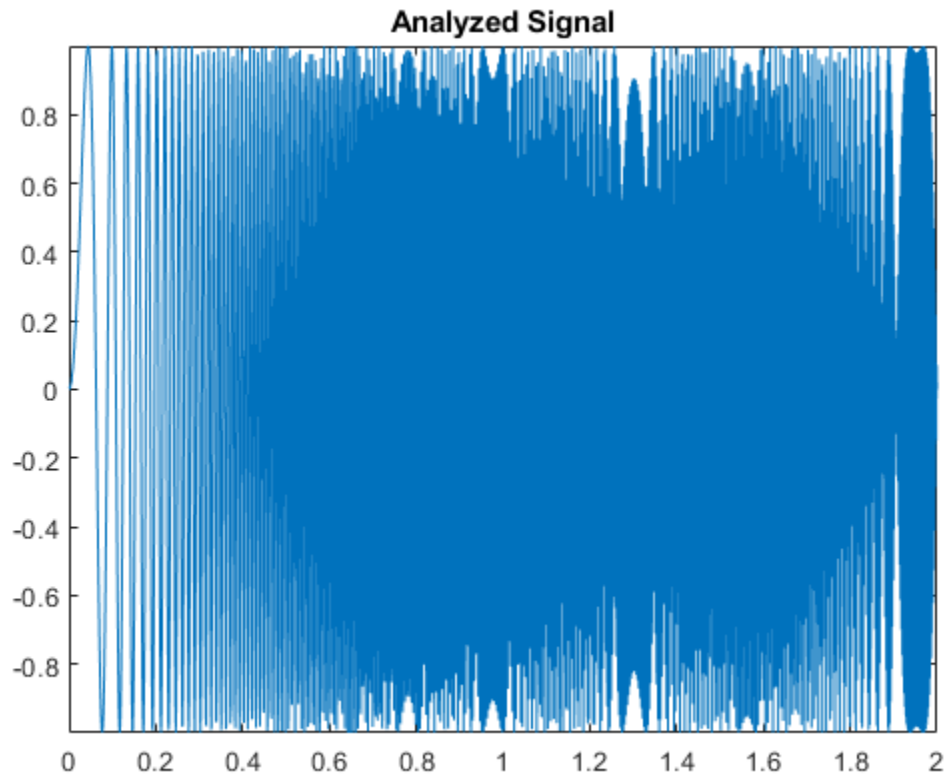
Wavelet Packet Spectrum of Chirp Signal

Create the chirp signal.

```
fs = 1000;  
t = 0:1/fs:2;  
% create chirp signal  
y = sin(256*pi*t.^2);
```

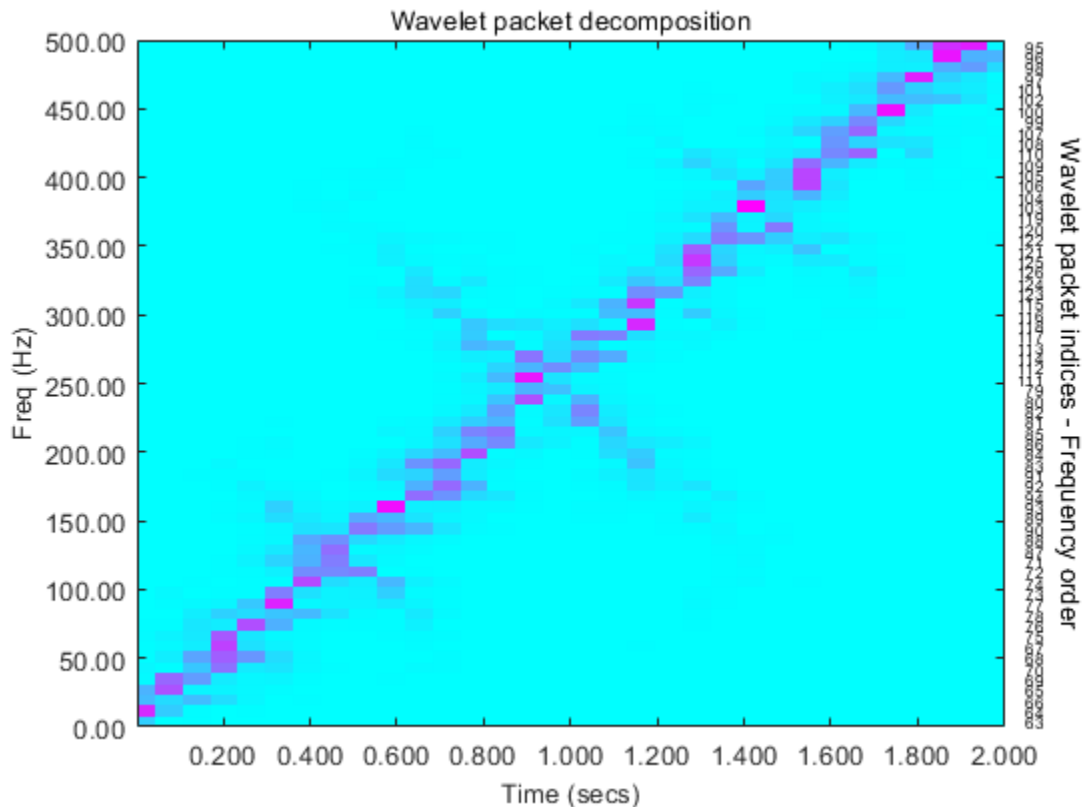
Plot the analyzed signal.


```
plot(t,y);  
axis tight  
title('Analyzed Signal');
```



Get the wavelet packet spectrum estimates.

```
level = 6;  
wpt = wpdec(y,level,'sym8');  
figure;  
[S,T,F] = wpspectrum(wpt,fs,'plot');
```



Definitions

Wavelet Packet Spectrum

The wavelet packet spectrum contains the absolute values of the coefficients from the frequency-ordered terminal nodes of the input binary wavelet packet tree. The terminal nodes provide the finest level of frequency resolution in the wavelet packet transform. If J denotes the level of the wavelet packet transform and F_s is the sampling frequency, the terminal nodes approximate bandpass filters of the form:

$$\left[\frac{nFs}{2^{J+1}}, \frac{(n+1)Fs}{2^{J+1}} \right) \quad n = 0, 1, 2, 3, \dots, 2^J - 1$$

At the terminal level of the wavelet packet tree, the transform divides the interval from 0 to the Nyquist frequency into bands of approximate width $F_s/2^{J+1}$.

Algorithms

wpspectrum computes the wavelet packet spectrum as follows:

- Extract the wavelet packet coefficients corresponding to the terminal nodes. Take the absolute value of the coefficients.
- Order the wavelet packet coefficients by frequency ordering.
- Determine the time extent on the original time axis corresponding to each wavelet packet coefficient. Repeat each wavelet packet coefficient to fill in the time gaps between neighboring wavelet packet coefficients and create a vector equal in length to node 0 of the wavelet packet tree object.

References

Wickerhauser, M.V. *Lectures on Wavelet Packet Algorithms*, Technical Report, Washington University, Department of Mathematics, 1992.

See Also

otnodes | wpdec

Topics

“Wavelet Packet Spectrum”

Introduced in R2010b

wpsplt

Split (decompose) wavelet packet

Syntax

```
T = wpsplt(T,N)
[T,cA,cD] = wpsplt(T,N)
[T,cA,cH,cV,cD] = wpsplt(T,N)
```

Description

wpsplt is a one- or two-dimensional wavelet packet analysis function.

wpsplt updates the wavelet packet tree after the decomposition of a node.

$T = \text{wpsplt}(T,N)$ returns the modified wavelet packet tree T corresponding to the decomposition of the node N .

For a one-dimensional decomposition,

$[T,cA,cD] = \text{wpsplt}(T,N)$ with cA = approximation and cD = detail of node N .

For a two-dimensional decomposition,

$[T,cA,cH,cV,cD] = \text{wpsplt}(T,N)$ with cA = approximation and cH, cV, cD = horizontal, vertical, and diagonal details of node N .

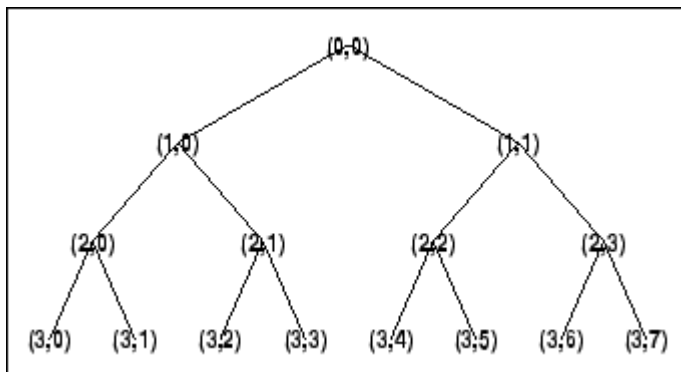
Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load signal.
load noisdopp;
x = noisdopp;
```

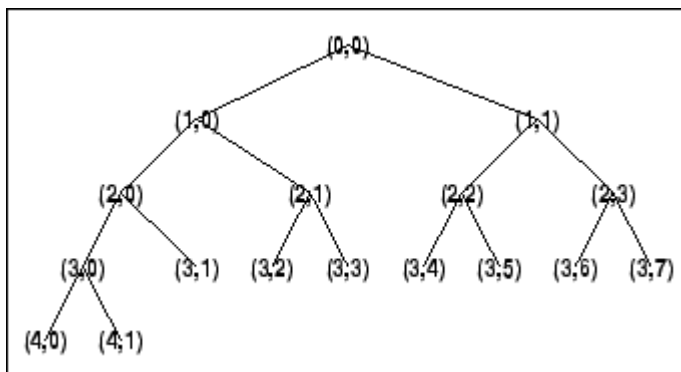
```
% Decompose x at depth 3 with db1 wavelet packets.
wpt = wpdec(x,3,'db1');

% Plot wavelet packet tree wpt.
plot(wpt)
```



```
% Decompose packet (3,0).
wpt = wpsplt(wpt,[3 0]);
% or equivalently wpsplt(wpt,7).

% Plot wavelet packet tree wpt.
plot(wpt)
```



See Also

wavedec | wavedec2 | wpdec | wpdec2 | wpjoin

Introduced before R2006a

wpthcoef

Wavelet packet coefficients thresholding

Syntax

$NT = \text{wpthcoef}(T, \text{KEEPAPP}, \text{SORH}, \text{THR})$

Description

wpthcoef is a one- or two-dimensional de-noising and compression utility.

$NT = \text{wpthcoef}(T, \text{KEEPAPP}, \text{SORH}, \text{THR})$ returns a new wavelet packet tree NT obtained from the wavelet packet tree T by coefficients thresholding.

If $\text{KEEPAPP} = 1$, approximation coefficients are not thresholded; otherwise, they can be thresholded.

If $\text{SORH} = 's'$, soft thresholding is applied; if $\text{SORH} = 'h'$, hard thresholding is applied (see `wthresh` for more information).

THR is the threshold value.

See Also

`wpdec` | `wpdec2` | `wpdencmp` | `wthresh`

Introduced before R2006a

wptree

WPTREE constructor

Syntax

```
T = wptree(ORDER,DEPTH,X,WNAME,ENT_TYPE,PARAMETER)
T = wptree(ORDER,DEPTH,X,WNAME)
T = wptree(ORDER,DEPTH,X,WNAME,'shannon')
T = wptree(ORDER,DEPTH,X,WNAME,ENT_TYPE,ENT_PAR,USERDATA)
```

Description

`T = wptree(ORDER,DEPTH,X,WNAME,ENT_TYPE,PARAMETER)` returns a complete wavelet packet tree `T`.

`ORDER` is an integer representing the order of the tree (the number of “children” of each non terminal node). `ORDER` must be equal to 2 or 4.

If `ORDER = 2`, `T` is a WPTREE object corresponding to a wavelet packet decomposition of the vector (signal) `X`, at level `DEPTH` with a particular wavelet `WNAME`.

If `ORDER = 4`, `T` is a WPTREE object corresponding to a wavelet packet decomposition of the matrix (image) `X`, at level `DEPTH` with a particular wavelet `WNAME`.

`ENT_TYPE` is a character vector or string scalar containing the entropy type and `ENT_PAR` is an optional parameter used for entropy computation (see `wentropy`, `wpdec`, or `wpdec2` for more information).

`T = wptree(ORDER,DEPTH,X,WNAME)` is equivalent to `T = wptree(ORDER,DEPTH,X,WNAME,'shannon')`

With `T = wptree(ORDER,DEPTH,X,WNAME,ENT_TYPE,ENT_PAR,USERDATA)` you may set a `userdata` field.

The function `wptree` returns a WPTREE object.

For more information on object fields, see the `get` function or type


```
help wptree/get
```

Class WPTREE (Parent class: DTREE)

Fields

'dtree'	DTREE parent object
'wavInfo'	Structure (wavelet information)
'entInfo'	Structure (entropy information)

The wavelet information structure, 'wavInfo', contains

'wavName'	Wavelet name
'Lo_D'	Low Decomposition filter
'Hi_D'	High Decomposition filter
'Lo_R'	Low Reconstruction filter
'Hi_R'	High Reconstruction filter

The entropy information structure, 'entInfo', contains

'entName'	Entropy name
'entPar'	Entropy parameter

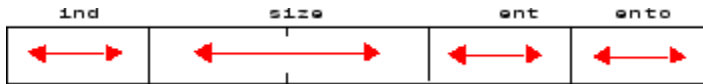
Fields from the DTREE parent object:

'allNI'	All nodes information
---------	-----------------------

'allNI' is an array of size nbnode by 5, which contains

ind	Index
size	Size of data
ent	Entropy
ento	Optimal entropy

Each line is built based on the following scheme:

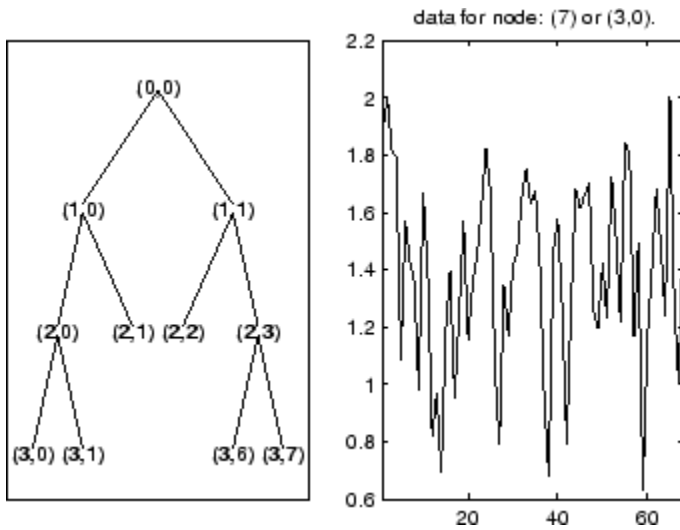


Examples

```
% Create a wavelet packet tree.  
x = rand(1,512);  
t = wptree(2,3,x,'db3');  
t = wpjoin(t,[4;5]);
```

```
% Plot tree t4.  
plot(t);
```

```
% Click the node (3,0), (see the plot function).
```



See Also

[dtree](#) | [ntree](#)

Introduced before R2006a

wpviewcf

Plot wavelet packets colored coefficients

Syntax

```
wpviewcf(T,CMODE)
wpviewcf(T,CMODE,NBCOL)
```

Description

`wpviewcf(T,CMODE)` plots the colored coefficients for the terminal nodes of the tree T .

T is a wavelet packet tree and $CMODE$ is an integer, which represents the color mode. The color modes are listed in the table below.

Color Mode	Description
1	Frequency order - Global coloration - Absolute values
2	Frequency order - By level - Absolute values
3	Frequency order - Global coloration - Values
4	Frequency order - By level coloration - Values
5	Natural order - Global coloration - Absolute values
6	Natural order - By level - Absolute values
7	Natural order - Global coloration - Values
8	Natural order - By level coloration - Values

`wpviewcf(T,CMODE,NBCOL)` uses $NBCOL$ colors.

Examples

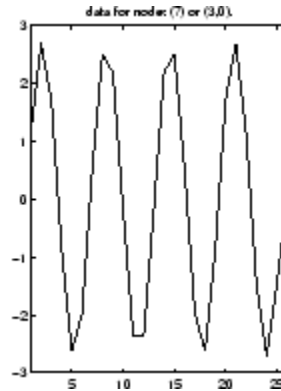
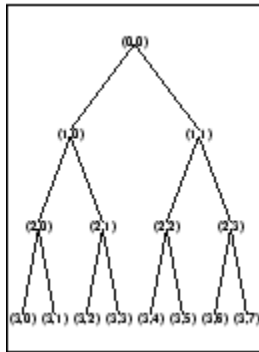
```
% Create a wavelet packet tree.
x = sin(8*pi*[0:0.005:1]);
```

```
t = wpdec(x,3,'db1');
```

```
% Plot tree t.
```

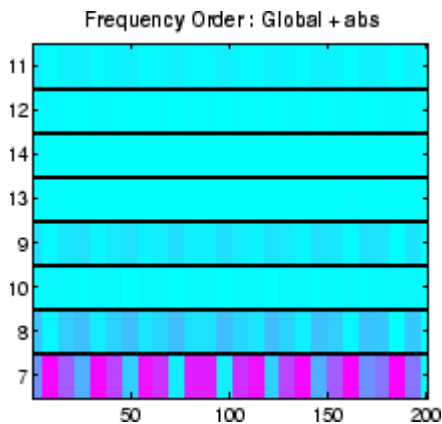
```
% Click the node (3,0), (see the plot function)
```

```
plot(t);
```



```
% Plot the colored wavelet packet coefficients.
```

```
wpviewcf(t,1);
```



See Also

[wpdec](#)

Introduced before R2006a

wrcoef

Reconstruct single branch from 1-D wavelet coefficients

Syntax

```
X = wrcoef('type',C,L,wname,N)
X = wrcoef('type',C,L,Lo_R,Hi_R,N)
X = wrcoef('type',C,L,wname)
X = wrcoef('type',C,L,Lo_R,Hi_R)
```

Description

wrcoef reconstructs the coefficients of a one-dimensional signal, given a wavelet decomposition structure (C and L) and either a specified wavelet (wname, see wfilters for more information) or specified reconstruction filters (Lo_R and Hi_R).

`X = wrcoef('type',C,L,wname,N)` computes the vector of reconstructed coefficients, based on the wavelet decomposition structure [C,L] (see wavedec for more information), at level N. wname is a character vector or string scalar containing the wavelet name.

Argument 'type' determines whether approximation ('type' = 'a') or detail ('type' = 'd') coefficients are reconstructed. When 'type' = 'a', N is allowed to be 0; otherwise, a strictly positive number N is required. Level N must be an integer such that $N \leq \text{length}(L) - 2$.

`X = wrcoef('type',C,L,Lo_R,Hi_R,N)` computes coefficients as above, given the reconstruction filters you specify.

`X = wrcoef('type',C,L,wname)` and `X = wrcoef('type',C,L,Lo_R,Hi_R)` reconstruct coefficients of maximum level $N = \text{length}(L) - 2$.

Examples

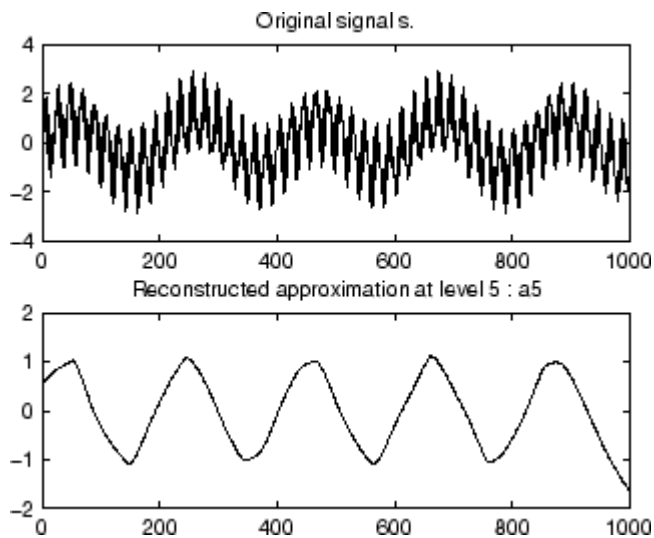
```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load a one-dimensional signal.
load sumsin; s = sumsin;

% Perform decomposition at level 5 of s using sym4.
[c,l] = wavedec(s,5,'sym4');

% Reconstruct approximation at level 5,
% from the wavelet decomposition structure [c,l].
a5 = wrcoef('a',c,l,'sym4',5);

% Using some plotting commands,
% the following figure is generated.
```



See Also

[appcoef](#) | [detcoef](#) | [wavedec](#)

Introduced before R2006a

wrcoef2

Reconstruct single branch from 2-D wavelet coefficients

Syntax

```
X = wrcoef2('type',C,S,wname,N)
X = wrcoef2('type',C,S,Lo_R,Hi_R,N)
X = wrcoef2('type',C,S,wname)
X = wrcoef2('type',C,S,Lo_R,Hi_R)
```

Description

`wrcoef2` is a two-dimensional wavelet analysis function. `wrcoef2` reconstructs the coefficients of an image.

`X = wrcoef2('type',C,S,wname,N)` computes the matrix of reconstructed coefficients of level `N`, based on the wavelet decomposition structure `[C,S]` (see `wavedec2` for more information).

`wname` is a character vector or string scalar containing the name of the wavelet (see `wfilters` for more information). If `'type' = 'a'`, approximation coefficients are reconstructed; otherwise if `'type' = 'h'` (`'v'` or `'d'`, respectively), horizontal (vertical or diagonal, respectively) detail coefficients are reconstructed.

Level `N` must be an integer such that $0 \leq N \leq \text{size}(S,1) - 2$ if `'type' = 'a'` and such that $1 \leq N \leq \text{size}(S,1) - 2$ if `'type' = 'h', 'v', or 'd'`.

Instead of giving the wavelet name, you can give the filters.

For `X = wrcoef2('type',C,S,Lo_R,Hi_R,N)`, `Lo_R` is the reconstruction low-pass filter and `Hi_R` is the reconstruction high-pass filter.

`X = wrcoef2('type',C,S,wname)` or `X = wrcoef2('type',C,S,Lo_R,Hi_R)` reconstruct coefficients of maximum level `N = size(S,1) - 2`.

Examples

```
% The current extension mode is zero-padding (see dwtmode).
```

```
% Load an image.
```

```
load woman;
```

```
% X contains the loaded image.
```

```
% Perform decomposition at level 2  
% of X using sym5.
```

```
[c,s] = wavedec2(X,2,'sym5');
```

```
% Reconstruct approximations at  
% levels 1 and 2, from the wavelet  
% decomposition structure [c,s].
```

```
a1 = wrcoef2('a',c,s,'sym5',1);
```

```
a2 = wrcoef2('a',c,s,'sym5',2);
```

```
% Reconstruct details at level 2,  
% from the wavelet decomposition  
% structure [c,s].
```

```
% 'h' is for horizontal,
```

```
% 'v' is for vertical,
```

```
% 'd' is for diagonal.
```

```
hd2 = wrcoef2('h',c,s,'sym5',2);
```

```
vd2 = wrcoef2('v',c,s,'sym5',2);
```

```
dd2 = wrcoef2('d',c,s,'sym5',2);
```

```
% All these images are of same size sX.
```

```
sX = size(X)
```

```
sX =
```

```
256 256
```

```
sa1 = size(a1)
```

```
sa1 =
```

```
256 256
```

```
shd2 = size(hd2)
```

```
shd2 =
```

```
256 256
```

Tips

If C and S are obtained from an indexed image analysis (respectively a truecolor image analysis) then X is an m -by- n matrix (respectively an m -by- n -by-3 array).

For more information on image formats, see the reference pages of `image` and `imfinfo` functions.

See Also

`appcoef2` | `detcoef2` | `wavedec2`

Introduced before R2006a

wrev

Flip vector

Syntax

```
Y = wrev(X)
```

Description

wrev is a general utility.

`Y = wrev(X)` reverses the vector `X`.

Examples

```
v = [1 2 3];  
wrev(v)  
wrev(v')
```

See Also

`fliplr` | `flipud`

Introduced before R2006a

write

Write values in WPTREE fields

Syntax

```
T = write(T,'cfs',NODE,COEFS)
T = write(T,'cfs',N1,CFS1,'cfs',N2,CFS2, ...)
```

Description

`T = write(T,'cfs',NODE,COEFS)` writes coefficients for the terminal node `NODE`.

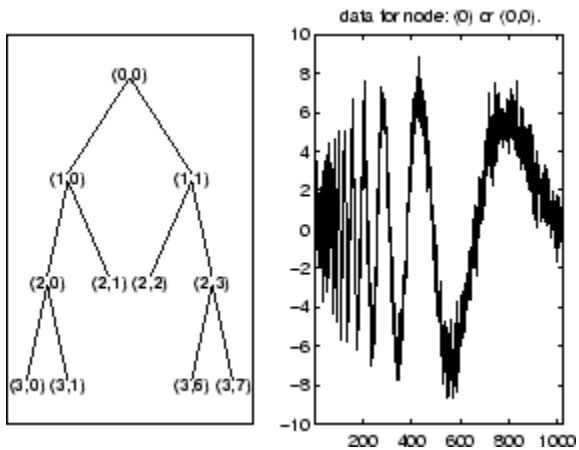
`T = write(T,'cfs',N1,CFS1,'cfs',N2,CFS2, ...)` writes coefficients `CFS1`, `CFS2`, ... for the terminal nodes `N1`, `N2`,

Caution The coefficients values must have the suitable size. You can use `S = read(T,'sizes',NODE)` or `S = read(T,'sizes',[N1;N2; ...])` in order to get those sizes.

Examples

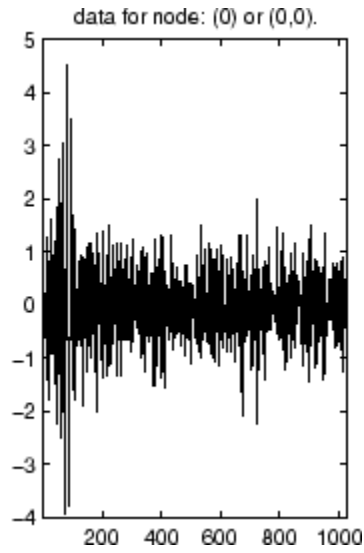
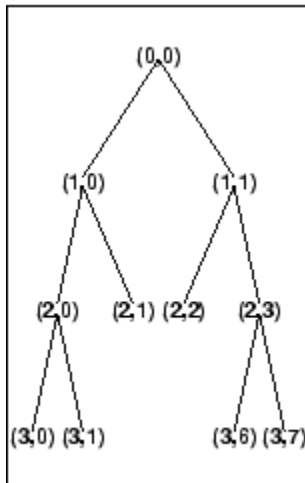
```
% Create a wavelet packet tree.
load noisdopp; x = noisdopp;
t = wpdec(x,3,'db3');
t = wpjoin(t,[4;5]);
```

```
% Plot tree t and click the node (0,0) (see the plot function).
plot(t);
```



```
% Write values.
sNod = read(t,'sizes',[4,5,7]);
cfs4 = zeros(sNod(1,:));
cfs5 = zeros(sNod(2,:));
cfs7 = zeros(sNod(3,:));
t = write(t,'cfs',4,cfs4,'cfs',5,cfs5,'cfs',7,cfs7);

% Plot tree t and click the node (0,0) (see the plot function).
plot(t)
```



See Also

`disp` | `get` | `read` | `set`

Introduced before R2006a

wscalogram

Scalogram for continuous wavelet transform

Note This function is no longer recommended. Use `cwt` instead.

Syntax

```
SC = wscalogram(TYPEPLOT,COEFS)
SC = wscalogram(TYPEPLOT,COEFS,'PropName1',PropVal1,...)
```

Description

`SC = wscalogram(TYPEPLOT,COEFS)` computes the scalogram `SC` which represents the percentage of energy for each coefficient. `COEFS` is the matrix of the continuous wavelet coefficients (see `cwt`).

The scalogram is obtained by computing:

```
S = abs(coefs.*coefs); SC = 100*S./sum(S(:))
```

When `TYPEPLOT` is equal to `'image'`, a scaled image of scalogram is displayed. When `TYPEPLOT` is equal to `'contour'`, a contour representation of scalogram is displayed. Otherwise, the scalogram is returned without plot representation.

`SC = wscalogram(TYPEPLOT,COEFS,'PropName1',PropVal1,...)` allows you to modify some properties. The valid choices for `PropName` are:

<code>'scales'</code>	Scales used for the CWT.
<code>'ydata'</code>	Signal used for the CWT.
<code>'xdata'</code>	x values corresponding to the signal values.
<code>'power'</code>	Positive real value. Default value is zero.

If `power > 0`, coefficients are first normalized

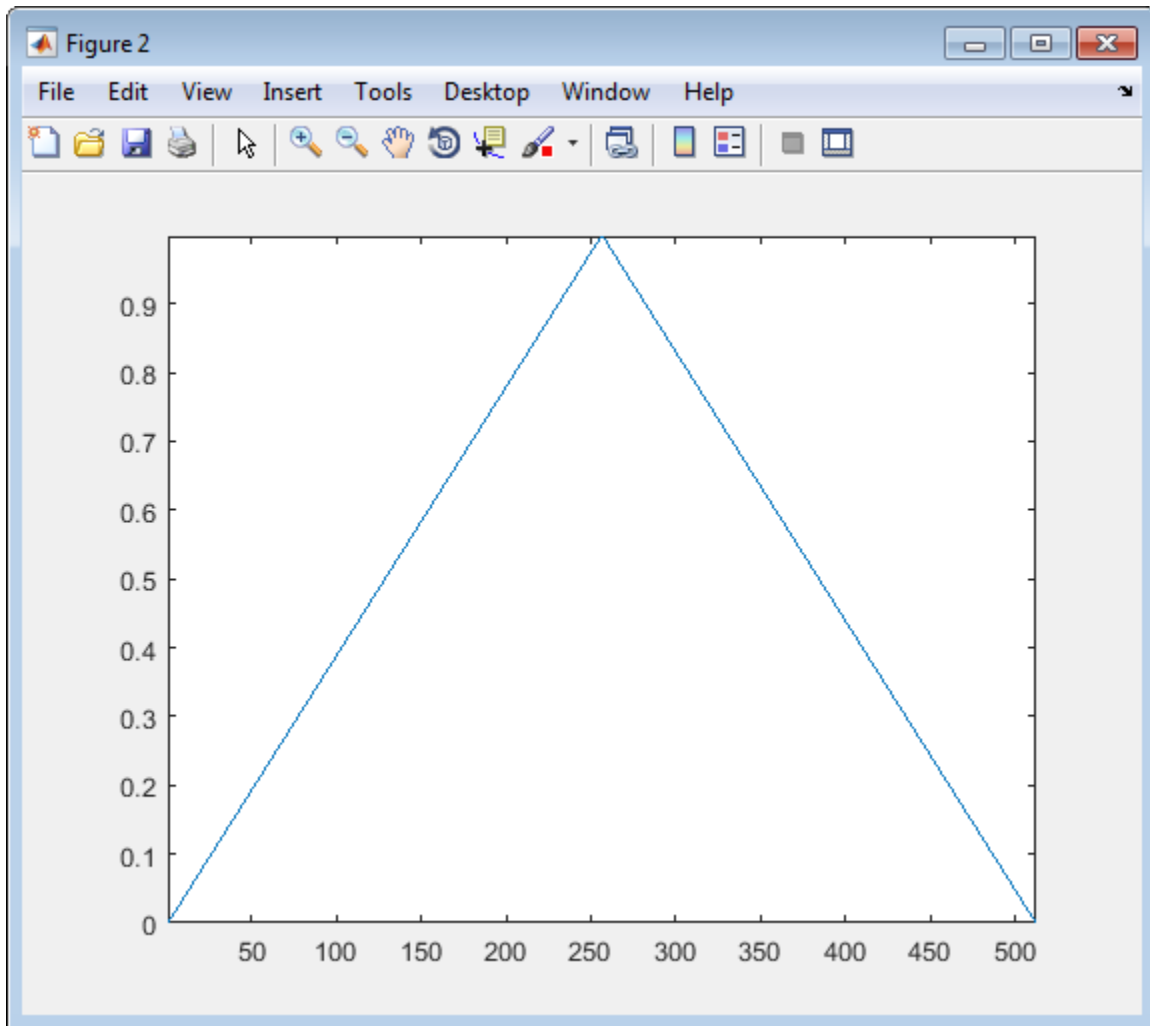
```
coefs(k,:) = coefs(k,)/(scales(k)^power)
```

and then the scalogram is computed as explained above.

Examples

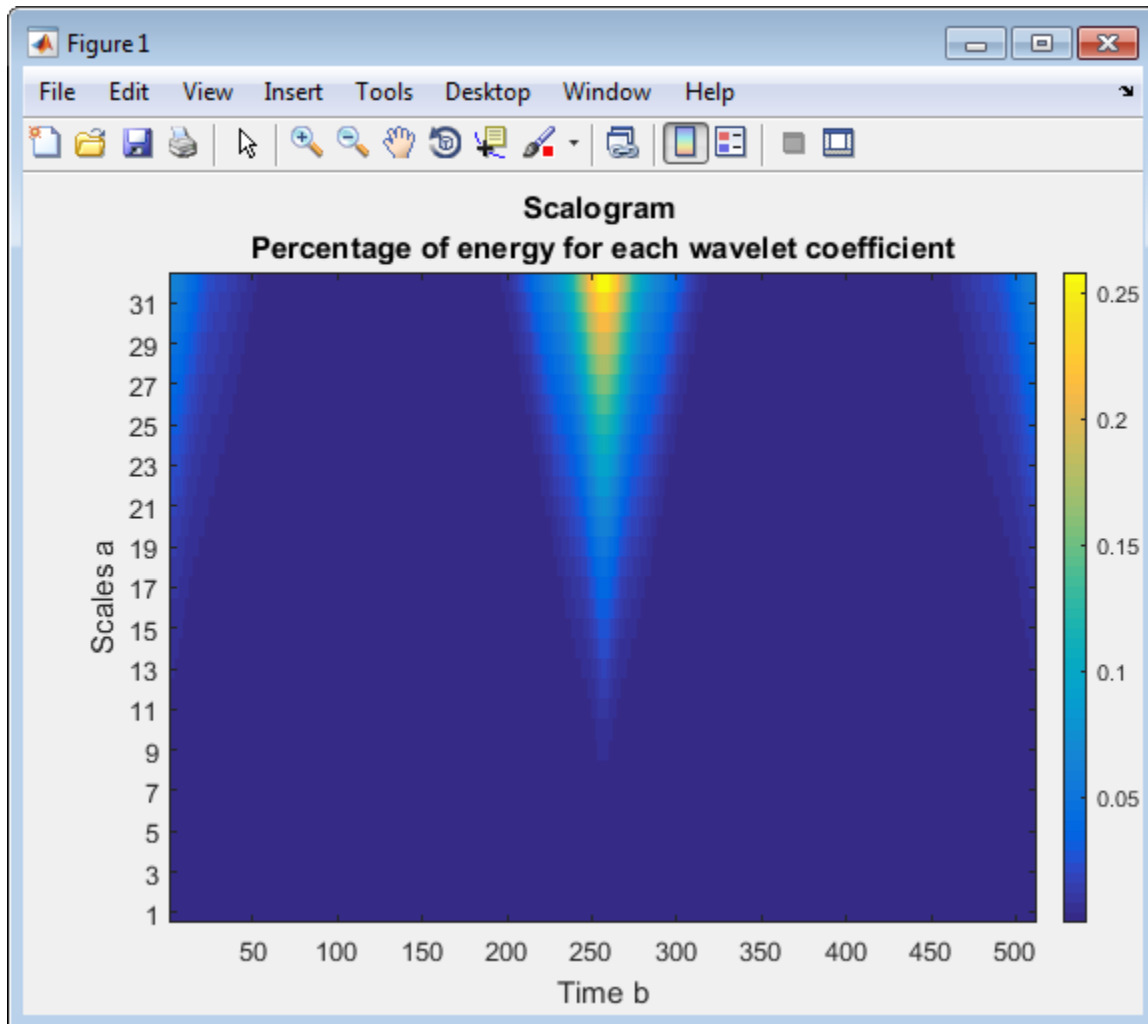
```
% Compute signal s  
t = linspace(-1,1,512);  
s = 1-abs(t);
```

```
% Plot signal s  
figure;  
plot(s), axis tight
```

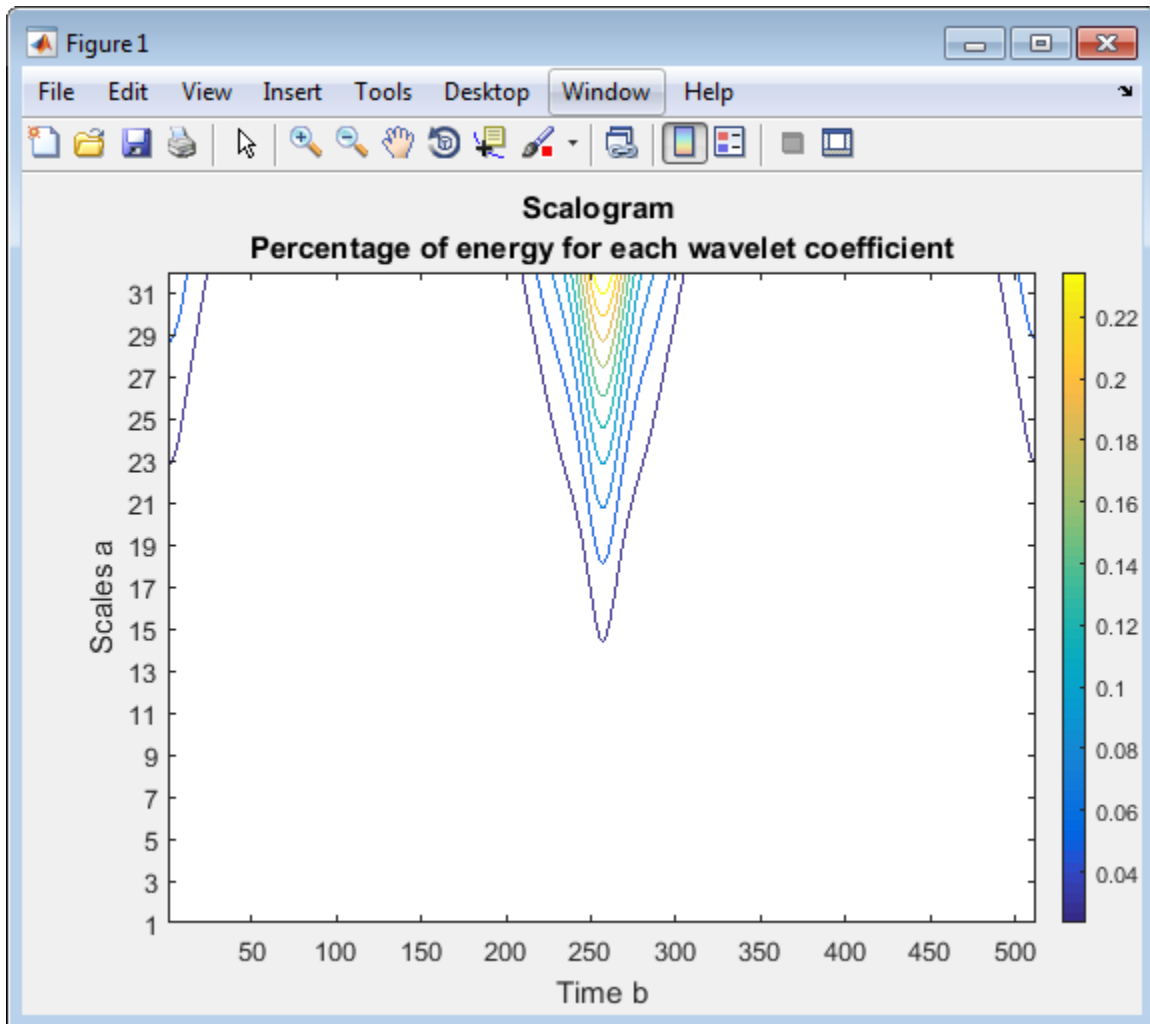



```
% Compute coefficients COEFS using cwt
COEFS = cwt(s,1:32,'cgau4');

% Compute and plot the scalogram (image option)
figure;
SC = wscalogram('image',COEFS);
```



```
% Compute and plot the scalogram (contour option)
figure;
SC = wscalogram('contour',COEFS);
```



See Also

cwt

Introduced in R2008a

wsst

Wavelet synchrosqueezed transform

Syntax

```
sst = wsst(x)
[sst,f] = wsst(x)
[ ___ ] = wsst(x,fs)
[ ___ ] = wsst(x,ts)
[ ___ ] = wsst( ___,wav)
wsst( ___ )
[ ___ ] = wsst( ___,Name,Value)
```

Description

`sst = wsst(x)` returns the wavelet synchrosqueezed transform, `sst`, which you use to examine data in the time-frequency plane. The synchrosqueezed transform has reduced energy smearing when compared to the continuous wavelet transform. The input, `x`, must be a 1-D real-valued signal with at least four samples. `wsst` computes the synchrosqueezed transform using the analytic Morlet wavelet.

`[sst,f] = wsst(x)` returns a vector of frequencies, `f`, in cycles per sample. The frequencies correspond to the rows of `sst`.

`[___] = wsst(x,fs)` computes the synchrosqueezed transform using the specified sampling frequency, `fs`, in Hz, to compute the synchrosqueezed transform. If you specify an `f` output, `wsst` returns the frequencies in Hz. You can use any previous combination of output values.

`[___] = wsst(x,ts)` uses a duration `ts` with a positive, scalar input, as the sampling interval. The duration can be in years, days, hours, minutes, or seconds. If you specify `ts` and the `f` output, `wsst` returns the frequencies in `f` in cycles per unit time, where the time unit is derived from specified duration.

[___] = `wsst(___ ,wav)` uses the analytic wavelet specified by `wav` to compute the synchrosqueezed transform. Valid values are 'amor' and 'bump', which specify the analytic Morlet and bump wavelet, respectively.

`wsst(___)` with no output arguments plots the synchrosqueezed transform as a function of time and frequency. If you do not specify a sampling frequency, `fs`, or interval, `ts`, the synchrosqueezed transform is plotted in cycles per sample. If you specify a sampling frequency, the synchrosqueezed transform is plotted in Hz. If you specify a sampling interval using a duration, the plot is in cycles per unit time. The time units are derived from the duration.

[___] = `wsst(___ ,Name,Value)` returns the synchrosqueezed transform with additional options specified by one or more `Name,Value` pair arguments.

Examples

Synchrosqueezed Transform of Speech Signal

Obtain the wavelet synchrosqueezed transform of a speech sample using default values.

```
load mtlb;  
sst = wsst(mtlb);
```

Synchrosqueezed Transform and Reconstruction of Speech Signal

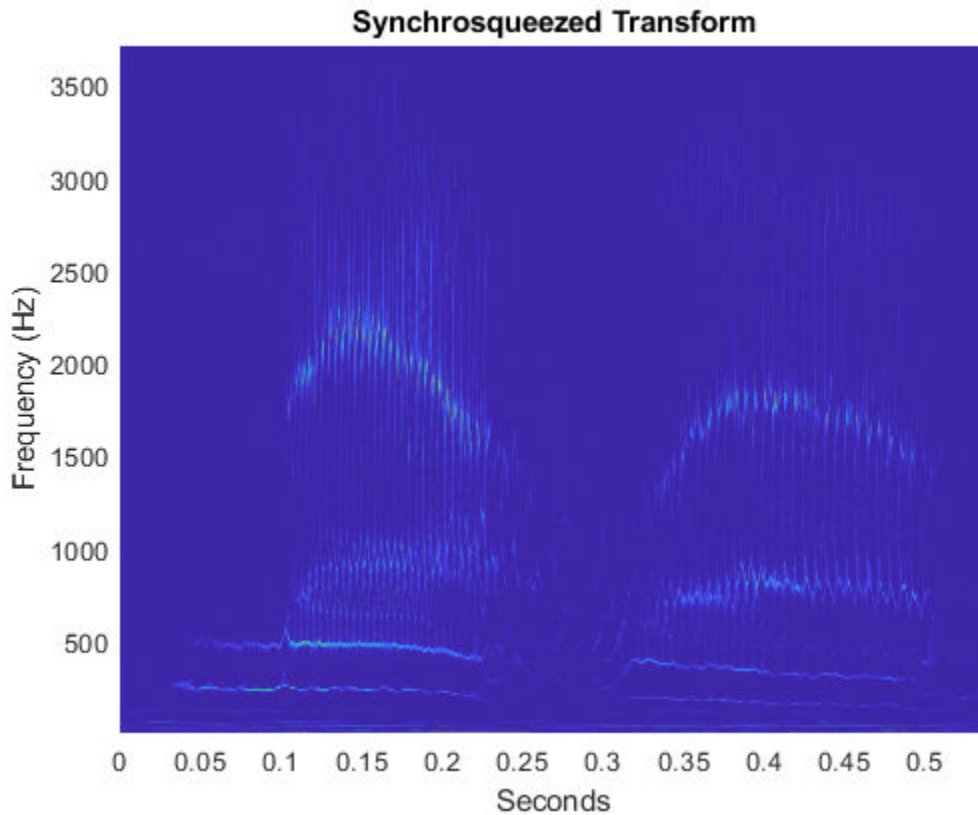
Obtain the wavelet synchrosqueezed transform of a speech signal and compare the original and reconstructed signals.

Load the speech signal and obtain its synchrosqueezed transform.

```
load mtlb  
soundsc(mtlb,Fs)  
dt = 1/Fs;  
t = 0:dt: numel(mtlb)*dt-dt;  
[sst,f] = wsst(mtlb,Fs);
```

Plot the synchrosqueezed transform.

```
pcolor(t,f,abs(sst))  
shading interp  
xlabel('Seconds')  
ylabel('Frequency (Hz)')  
title('Synchrosqueezed Transform')
```



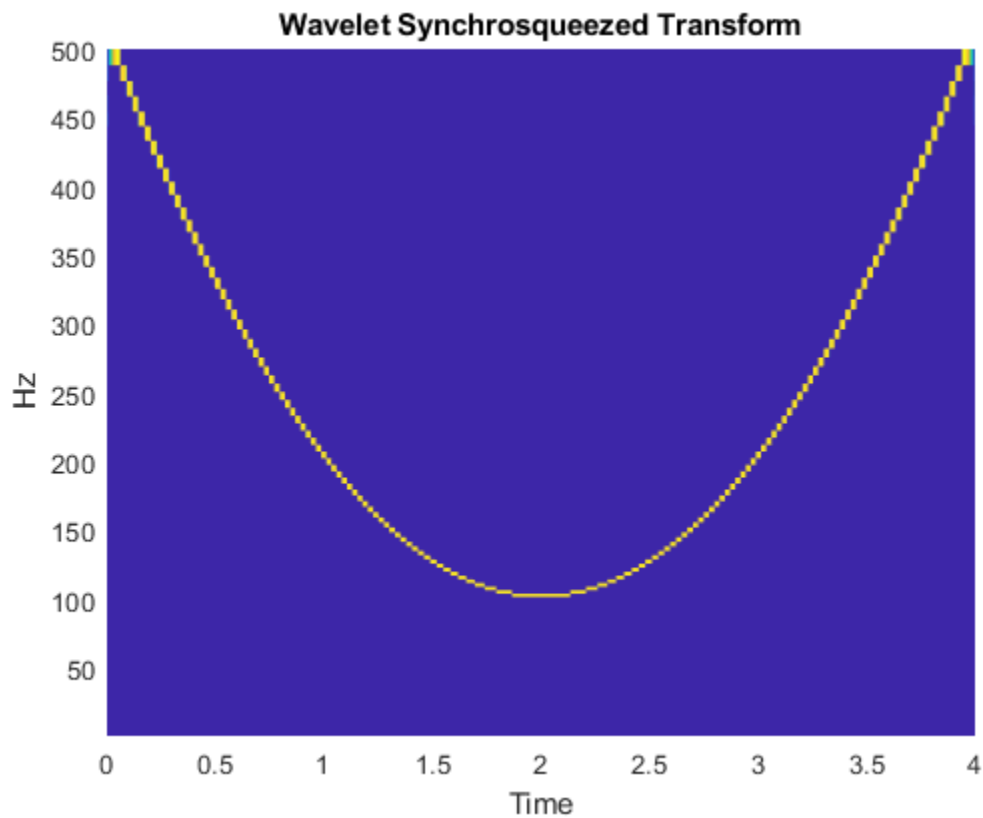
Obtain the inverse synchrosqueezed transform and play the reconstructed speech signal.

```
xrec = iwsst(sst);  
soundsc(xrec,Fs)
```

Synchrosqueezed Transform of Quadratic Chirp

Obtain and plot the wavelet synchrosqueezed transform of a quadratic chirp. The chirp is sampled at 1000 Hz.

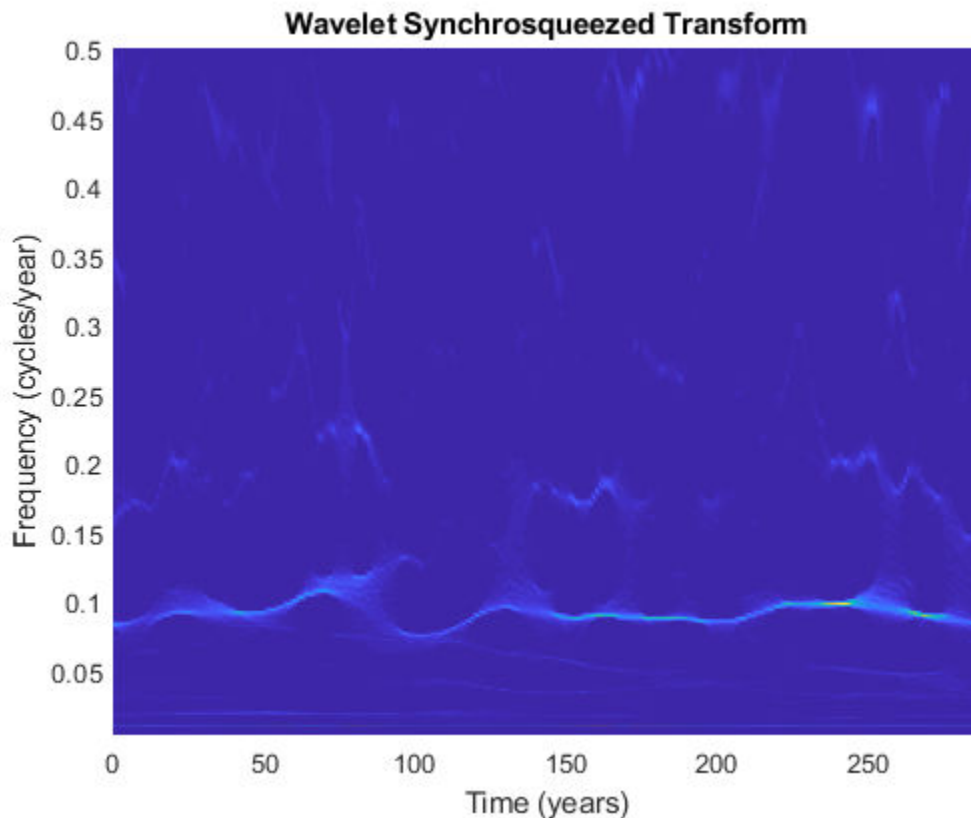
```
load quadchirp;  
[sst,f] = wsst(quadchirp,1000);  
hp = pcolor(tquad,f,abs(sst));  
hp.EdgeColor = 'none';  
title('Wavelet Synchrosqueezed Transform');  
xlabel('Time'); ylabel('Hz');
```



Synchrosqueezed Transform of Sunspot Data

Obtain the wavelet synchrosqueezed transform of sunspot data using the default Morlet wavelet. Specify the sampling interval to be one year.

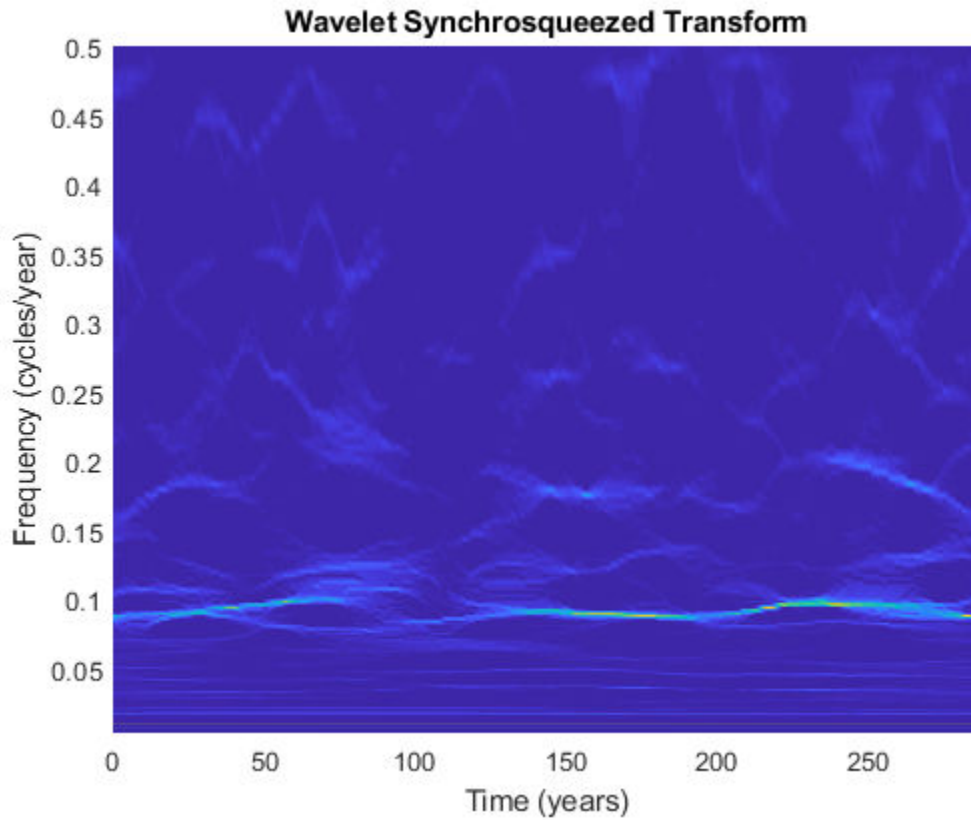
```
load sunspot.dat;  
wsst(sunspot(:,2),years(1))
```



Synchrosqueezed Transform of Sunspot Data Using Bump Wavelet

Obtain and plot the wavelet synchrosqueezed transform of sunspot data using the bump wavelet. Specify the sampling interval to be 1 for one sample per year.


```
load sunspot.dat
wsst(sunspot(:,2),years(1),'bump')
```



Input Arguments

x — Input signal

row or column vector of real values

Input signal, specified as a row or column vector. **x** must be a 1-D, real-valued signal with at least four samples.

fs — Sampling frequency

positive scalar

Sampling frequency, specified as a positive scalar.

ts — Sampling interval

duration with positive scalar input

Sampling interval, also known as the sampling period, specified as a duration with positive scalar input. Valid durations are years, days, hours, seconds, and minutes. You cannot use calendar durations (`caldays`, `calweeks`, `calmonths`, `calquarters`, or `calyears`). You cannot specify both `ts` and `fs`.

Example: `sst = wsst(x, hours(12))`

wav — Analytic wavelet

'amor' (default) | 'bump'

Analytic wavelet used to compute the synchrosqueezed transform, specified as one of the following:

- 'amor' — Analytic Morlet wavelet
- 'bump' — Bump wavelet

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'VoicesPerOctave', 26`

VoicesPerOctave — Number of voices per octave

32 (default) | even integer from 10 to 48

Number of voices per octave to use in the synchrosqueezed transform, specified as the comma-separated pair consisting of `'VoicesPerOctave'` and an even integer from 10 to 48. The product of the number of voices per octave and the number of octaves is the number of scales. The number of octaves depends on the size of the input `x` and is `floor(log2(numel(x))) - 1`.

ExtendSignal — Extend input signal symmetrically

false (default) | true

Option to extend the input signal symmetrically, specified as the comma-separated pair consisting of 'ExtendSignal' and either `false` or `true`. Extending the signal symmetrically can mitigate boundary effects. If you specify `false`, then the signal is not extended. If you specify `true`, then the signal is extended.

Output Arguments

sst — Synchrosqueezed transform

matrix

Synchrosqueezed transform, returned as a matrix. By default, the synchrosqueezed transform uses $\text{floor}(\log_2(\text{numel}(x))) - 1$ octaves, 32 voices per octave, and the analytic Morlet wavelet. `sst` is an N_a -by- N matrix where N_a is the number of scales, and N is the number of samples in x . The default number of scales is $32 * (\text{floor}(\log_2(\text{numel}(x))) - 1)$.

f — Frequencies

vector

Frequencies of the synchrosqueezed transform, returned as a vector. The frequencies correspond to the rows of the `sst`. If you do not specify `fs` or `ts`, the frequencies are in cycles per sample. If you specify `fs`, the frequencies are in Hz. If you specify `ts`, the frequencies are in cycles per unit time. The length of the frequency vector is the same as the number of `sst` rows. If you specify `ts` as the sampling interval, `ts` is used to compute the scale-to-frequency conversion for `f`.

References

- [1] Daubechies, I., J. Lu, and H.-T. Wu. "Synchrosqueezed wavelet transforms: an empirical mode decomposition-like tool." *Applied and Computational Harmonic Analysis*. Vol. 30, Number 2, 2011, pp. 243–261.
- [2] Thakur, G., E. Brevdo, N. S. Fučkar, and H.-T. Wu. "The Synchrosqueezing algorithm for time-varying spectral analysis: robustness properties and new paleoclimate applications." *Signal Processing*. Vol. 93, Number 5, 2013, pp. 1079–1094.

See Also

days | duration | hours | iwsst | minutes | seconds | wsstridge | years

Topics

“Time-Frequency Reassignment and Mode Extraction with Synchrosqueezing”
“Wavelet Synchrosqueezing”

Introduced in R2016a

wsstridge

Time-frequency ridges from wavelet synchrosqueezing

Syntax

```
fridge = wsstridge(sst)
[fridge,iridge] = wsstridge(sst)
[ ___ ] = wsstridge(sst,penalty)
[ ___ ] = wsstridge( ___,f)
[ ___ ]= wsstridge( ___,Name,Value)
```

Description

`fridge = wsstridge(sst)` extracts the maximum energy time-frequency ridge in cycles per sample from the wavelet synchrosqueezed transform, `sst`. The `sst` input is the output of `wsst`. Each ridge is a separate signal mode.

`[fridge,iridge] = wsstridge(sst)` returns in `iridge` the row indices of `sst`. The row indices are the maximum time-frequency ridge at each sample. Use `iridge` to reconstruct the signal mode along a time-frequency ridge using `iwsst`.

`[___] = wsstridge(sst,penalty)` multiplies the squared distance between frequency bins by the `penalty` value. You can include any of the output arguments from previous syntaxes.

`[___] = wsstridge(___,f)` returns the maximum energy time-frequency ridge in cycles per unit time based on the `f` input frequency vector. `f` is the frequency output of `wsst`. The `f` input and `fridge` output have the same units.

`[___]= wsstridge(___,Name,Value)` returns the time-frequency ridge with additional options specified by one or more `Name,Value` pair arguments.

Examples

Extract Time-Frequency Ridge from Chirp Signal

Obtain the wavelet synchrosqueezed transform of a quadratic chirp and extract the maximum time-frequency ridge, in `fridge`, and the associated row indices, in `iridge`.

Load the chirp signal and obtain its synchrosqueezed transform.

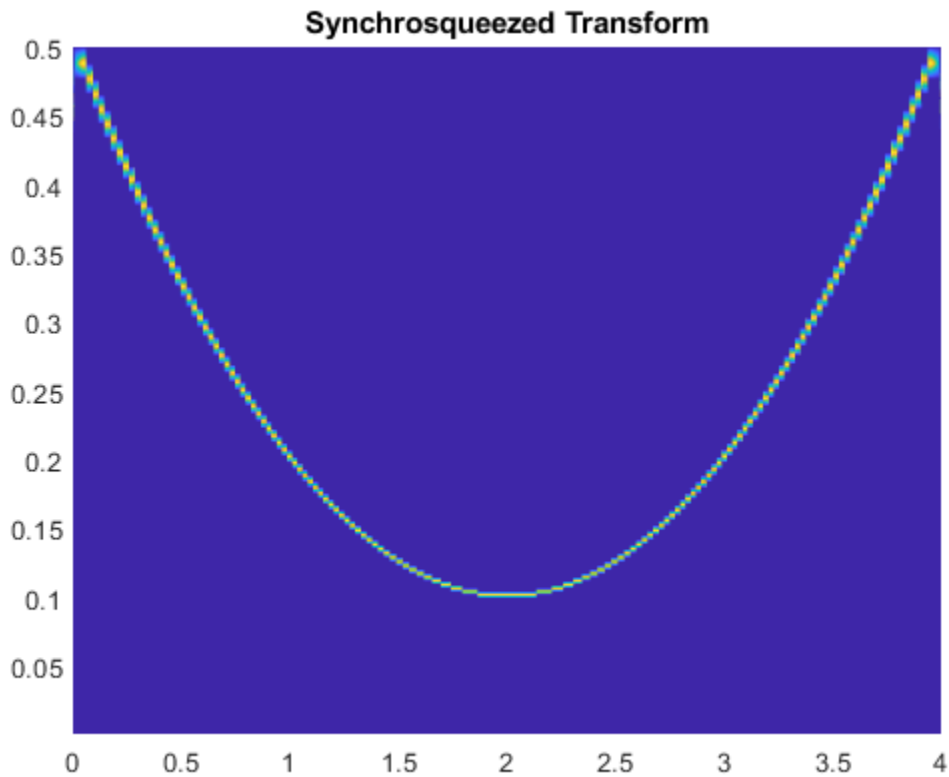
```
load quadchirp;  
[sst,f] = wssst(quadchirp);
```

Extract the maximum time-frequency ridge.

```
[fridge,iridge] = wsstridge(sst);
```

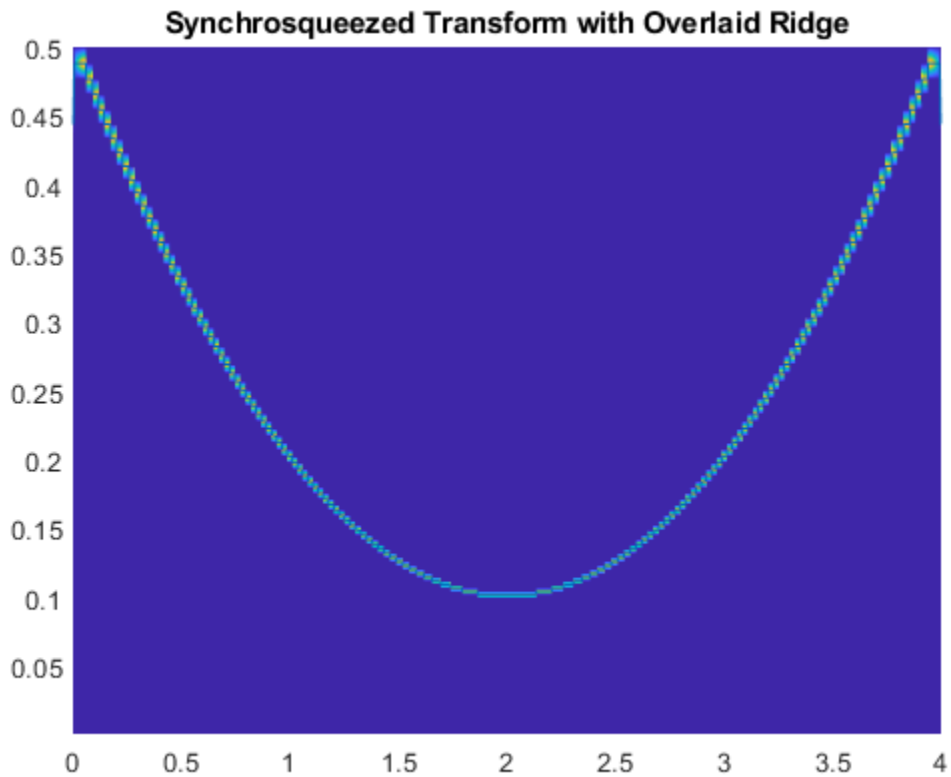
Plot the synchrosqueezed transform.

```
pcolor(tquad,f,abs(sst))  
shading interp  
title('Synchrosqueezed Transform')
```



Overlay the plot of the maximum energy frequency ridge.

```
hold on
plot(tquad,fridge)
title('Synchrosqueezed Transform with Overlaid Ridge')
```



Extract Time-Frequency Ridge from Multicomponent Signal

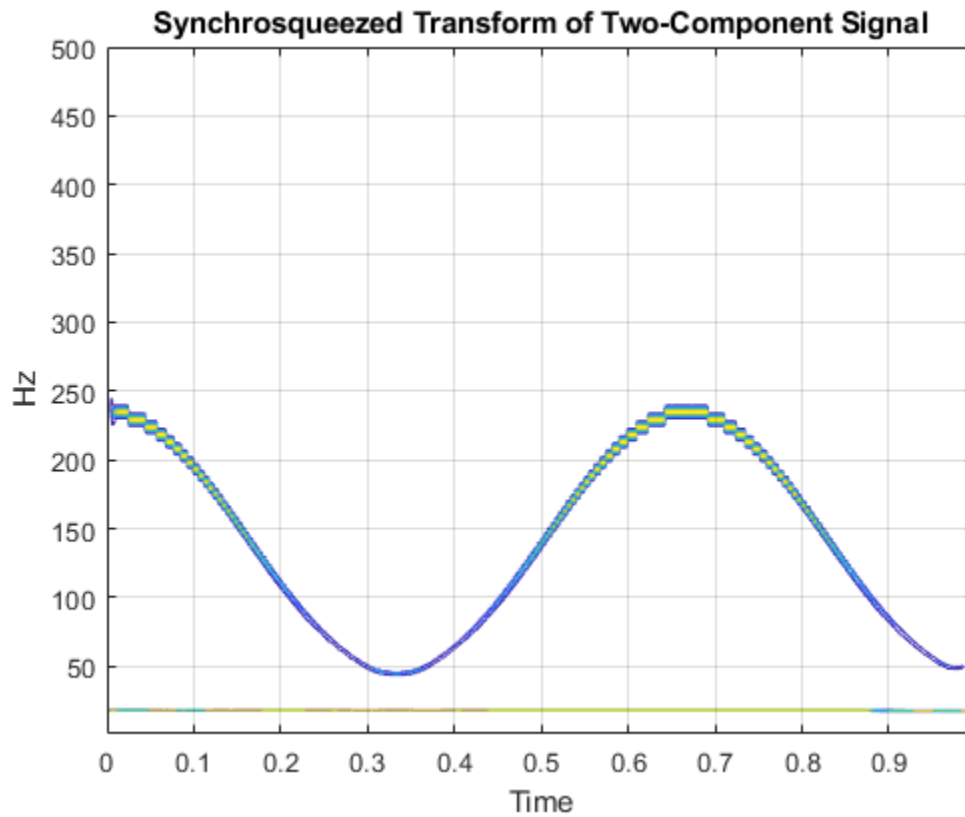
Extract the two highest energy modes from a multicomponent signal.

Obtain and plot the wavelet synchrosqueezed transform.

```
load multicompsig;  
sig = sig1+sig2;  
[sst,F] = wsst(sig,sampfreq);  
contour(t,F,abs(sst));  
xlabel('Time'); ylabel('Hz');
```

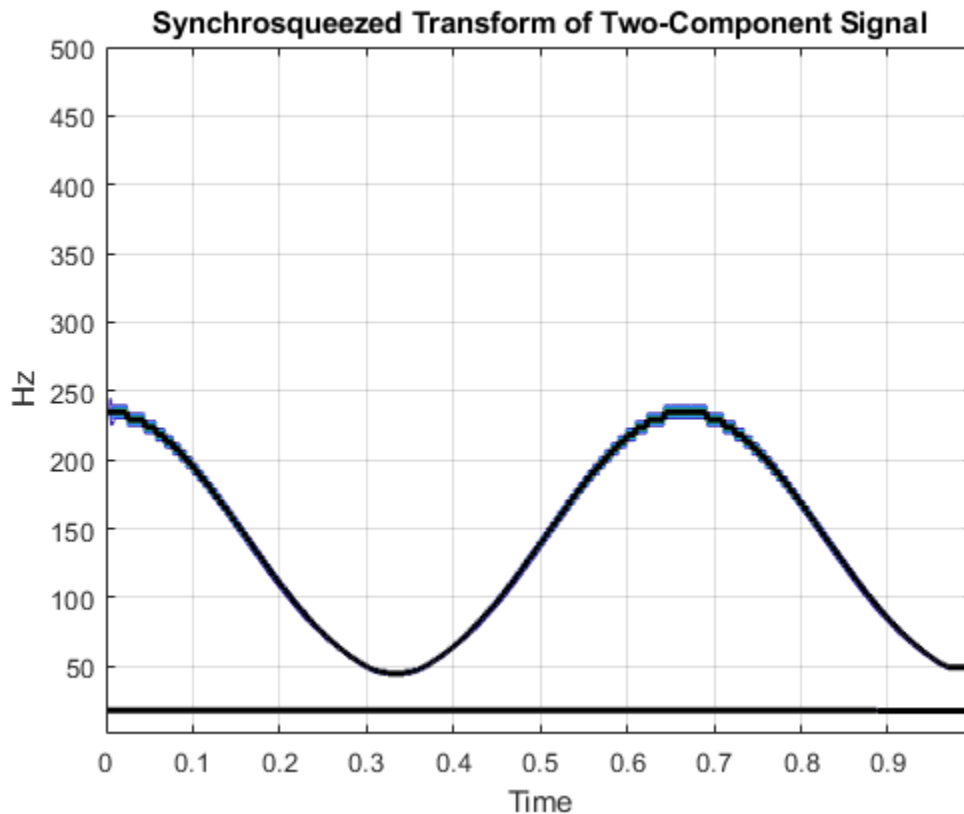


```
grid on;  
title('Synchrosqueezed Transform of Two-Component Signal');
```



Using a penalty of 10, extract the two highest energy modes and plot the result.

```
[fridge,iridge] = wsstridge(sst,10,F,'NumRidges',2);  
hold on;  
plot(t,fridge,'k','linewidth',2);
```



Input Arguments

sst — Synchrosqueezed transform

matrix

Synchrosqueezed transform, specified as a matrix. `sst` is a time-frequency matrix and is the output of `wsst`.

penalty — Frequency bins scaling penalty

0 (default) | nonnegative scalar

Frequency bins scaling penalty, specified as a nonnegative scalar. This input penalizes changes in frequency by multiplying the penalty value by the squared distance between frequency bins. Use a penalty term when you extract multiple ridges, or when you have a single modulated component in additive noise. The penalty term prevents jumps in frequency that occur when the region of highest energy in the time-frequency plane changes abruptly.

f – **Synchrosqueezed transform frequencies**

vector

Synchrosqueezed transform frequencies corresponding to the rows of the synchrosqueezed transform, which is the vector output of `wsst`. The number of elements in the frequency vector is equal to the number of rows in the `sst` input.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'NumRidges', 3`

NumRidges – **Number of highest energy time-frequency ridges**

1 (default) | positive integer

Number of highest energy time-frequency ridges to extract, specified as the comma-separated pair consisting of `'NumRidges'` and a positive integer. If this integer is greater than 1, `wsstridge` iteratively determines the maximum energy time-frequency ridge by removing the previously computed ridges and the default or specified `'NumFrequencyBins'` on either side of each ridge bin.

NumFrequencyBins – **Number of frequency bins to remove**

4 (default) | positive integer

Number of frequency bins to remove from synchrosqueezed transform `sst` when extracting multiple ridges, specified as the comma-separated pair consisting of `'NumFrequencyBins'` and a positive integer. This integer must be less than or equal to `round(size(sst,1)/4)`. You can specify the number of frequency bins to remove only if you extract more than one ridge. After extracting the highest energy time-frequency ridge, `wsstridge` removes the `sst` values corresponding to the `iridge` indices at each

time step. The energy is removed along the time-frequency ridge extended on both sides of the `iridge` index by the specified number of frequency bins. If the index of the extended time-frequency ridge exceeds the number of frequency bins at any time step, `wsstridge` truncates the removal region at the first or last frequency bin. To specify 'NumFrequencyBins', you must specify 'NumRidges'.

Output Arguments

fidge — Time-frequency ridge frequencies

vector or matrix

Time-frequency ridge frequencies, returned as a vector or matrix. The frequencies correspond to the time-frequency ridge at each time step. `fidge` is an N -by-`nr` matrix where N is the number of time samples (columns) in `sst` and `nr` is the number of ridges. The first column of the matrix contains the frequencies for the maximum energy time-frequency ridge in `sst`. Subsequent columns contain the frequencies for the time-frequency ridges in decreasing energy order. By default, `fidge` contains frequencies in cycles per sample.

iridge — Time-frequency ridge indices

vector or matrix

Time-frequency ridge row indices of `sst`, returned as a vector or matrix. The row indices in `iridge` correspond to the row index of the maximum time-frequency ridge for each `sst` column. `iridge` is an N -by-`nr` matrix where N is the number of time samples (columns) in `sst`, and `nr` is the number of ridges. The first column of the matrix contains the indices for the maximum energy time-frequency ridge in `sst`. Subsequent columns contain the indices for the time-frequency ridges in decreasing energy order.

Algorithms

The function uses a penalized forward-backward greedy algorithm to extract the maximum-energy ridges from a time-frequency matrix. The algorithm finds the maximum time-frequency ridge by minimizing $-\ln A$ at each time point, where A is the absolute value of the matrix. Minimizing $-\ln A$ is equivalent to maximizing the value of A . The algorithm optionally constrains jumps in frequency with a penalty that is proportional to the distance between frequency bins.

The following example illustrates the time-frequency ridge algorithm using a penalty that is two times the distance between frequency bins. Specifically, the distance between the elements (j, k) and (m, n) is defined as $(j - m)^2$. The time-frequency matrix has three frequency bins and three time steps. The matrix columns correspond to time steps, and the matrix rows correspond to frequency bins. The values in the second row represent a sine wave.

1 Suppose you have the matrix:

1	4	4
2	2	2
5	5	4

2 Update the value for the (1,2) element as follows.

- a** Leave the values at the first time point unaltered. Begin the algorithm with the (1,2) element of the matrix, which presents the first frequency bin at the second time point. The bin value is 4. Penalize the values in the first column based on their distance from the (1,2) element. Applying the penalty to the first column produces

original value + penalty \times distance

1	+ 2 \times 0	=	1
2	+ 2 \times 1	=	4
5	+ 2 \times 4	=	13

1	4
4	2
13	5

The minimum value of the first column is 1, which is in bin 1.

- b** Add the minimum value in column 1 to the current bin value, 4. The updated value for (1,2) becomes 5, which came from bin 1.
- 3** Update the values for the remaining elements in column 2 as follows.

Recompute the original column 1 values with the penalty factor using the same process as in Step 2a. Obtain the remaining second column values using the same process as in Step 2b. For example, when updating the (2,2) element, which has bin value 2, applying the penalty to the column yields

original value + penalty \times distance

$$\begin{aligned}1 + 2 \times 1 &= 3 \\2 + 2 \times 0 &= 2 \\5 + 2 \times 1 &= 7\end{aligned}$$

Add the minimum value, 2, to the current bin value. The updated value for (2,2) becomes 4. After updating the (3,2) element, the matrix is

$$\begin{array}{l}1 \quad 5_{(1)} \quad 4 \\2 \quad 4_{(2)} \quad 2 \\5 \quad 9_{(2)} \quad 4\end{array}$$

Only the second column has been updated. The subscripts indicate the index of the bin in the previous column from which a value came.

- 4 Repeat Step 2 for the third column. But now the penalty is applied to the updated second column. For example, when updating the (1,3) element, the penalty is

$$\begin{aligned}5 + 2 \times 0 &= 5 \\4 + 2 \times 1 &= 6 \\9 + 2 \times 4 &= 17\end{aligned}$$

The minimum value, 5, which is in the first bin, is added to the (1,3) bin value. After updating all the values in the third column, the final matrix is

$$\begin{array}{l}1 \quad 5_{(1)} \quad 9_{(1)} \\2 \quad 4_{(2)} \quad 6_{(2)} \\5 \quad 9_{(2)} \quad 10_{(2)}\end{array}$$

- 5 Starting at the last column of the matrix, find the minimum value. Walk back in time through the matrix by going from the current bin to the origin of that bin at the previous time point. Keep track of the bin indices, which form the path composing the ridge. The algorithm smooths the transition by using the origin bin instead of the bin with the minimum value. For this example, the ridge indices are 2, 2, 2, which matches the energy path of the sine wave in row 2 of the matrix shown in Step 1.

If you are extracting multiple ridges, the algorithm removes the first ridge from the time-frequency matrix and repeats the process.

References

- [1] Daubechies, I., J. Lu, and H.-T. Wu. "Synchrosqueezed wavelet transforms: an empirical mode decomposition-like tool." *Applied and Computational Harmonic Analysis*. Vol. 30, Number 2, 2011, pp. 243-261.

[2] Thakur, G., E. Brevdo, N. S. Fučkar, and H.-T. Wu. "The Synchrosqueezing algorithm for time-varying spectral analysis: Robustness properties and new paleoclimate applications." *Signal Processing*. Vol. 93, Number 4, 2013, pp. 1079-1094.

See Also

iwssst | wssst

Topics

"Time-Frequency Reassignment and Mode Extraction with Synchrosqueezing"
"Wavelet Synchrosqueezing"

Introduced in R2016a

wt

Continuous wavelet transform with a filter bank

Syntax

```
cfs = wt(fb,x)
[cfs,f] = wt(fb,x)
[cfs,f,coi] = wt(fb,x)
[cfs,f,coi,scalcfs] = wt(fb,x)
[cfs,p] = wt(fb,x)
[cfs,p,coi] = wt(fb,x)
[cfs,p,coi,scalcfs] = wt(fb,x)
```

Description

`cfs = wt(fb,x)` returns the continuous wavelet transform (CWT) coefficients of the signal `x`, using the CWT filter bank `fb`. `x` is a double-precision real- or complex-valued vector. `x` must have at least 4 samples. If `x` is real-valued, `cfs` is a 2-D matrix where each row corresponds to one scale. The column size of `cfs` is equal to the length of `x`. If `x` is complex-valued, `cfs` is a 3-D matrix, where the first page is the CWT for the positive scales (analytic part or counterclockwise component) and the second page is the cwt for the negative scales (anti-analytic part or clockwise component).

`[cfs,f] = wt(fb,x)` returns the frequencies, `f`, corresponding to the scales (rows) of `cfs` if the `SamplingPeriod` property is not specified in the CWT filter bank `fb`. If you do not specify a sampling frequency, `f` is in cycles/sample.

`[cfs,f,coi] = wt(fb,x)` returns the cone of influence, `coi` for the CWT. `coi` is in the same units as `f`. If the input `x` is complex, the `coi` applies to both pages of `cfs`.

`[cfs,f,coi,scalcfs] = wt(fb,x)` returns the scaling coefficients, `scalcfs` for the wavelet transform if the analyzing wavelet is 'Morse' or 'amor'. Scaling coefficients are not supported for the bump wavelet.

`[cfs,p] = wt(fb,x)` returns the periods, `p`, corresponding to the scales (rows) of `cfs` if you specify a sampling period in the CWT filter bank, `fb`. `p` has the same units and format as the duration scalar sampling period.

`[cfs,p,coi] = wt(fb,x)` returns the cone of influence `coi` in periods for the CWT. `coi` is an array of durations with the same format property as the sampling period. If the input `x` is complex, the `coi` applies to both pages of `cfs`.

`[cfs,p,coi,scalcfs] = wt(fb,x)` returns the scaling coefficients, `scalcfs`, for the wavelet transform if the analyzing wavelet is 'Morse' or 'amor'. Scaling coefficients are not supported for the bump wavelet.

Examples

Continuous Wavelet Transform Using a Filter Bank

Load a real-valued signal. Create a CWT filter bank that can be applied to the signal.

```
load noisdopp.mat
sig = noisdopp;
fb = cwtfilterbank('SignalLength',numel(sig));
```

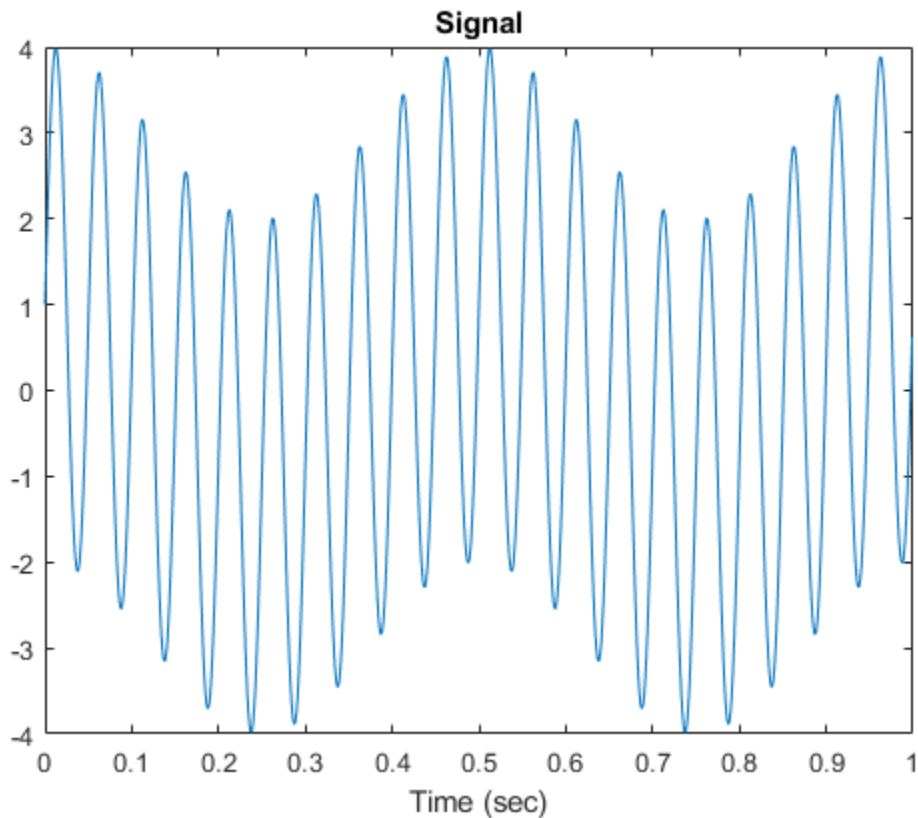
Use the filter bank to obtain the continuous wavelet transform of the signal.

```
cfs = wt(fb,sig);
```

Inverse Continuous Wavelet Transform Using Scaling Coefficients

Create and plot a signal sampled at 1000 Hz. Create a CWT filter bank that can be used on the signal. Since the signal is periodic, set the boundary extension property of the filter bank to 'periodic'.

```
Fs = 1000;
t = 0:1/Fs:1-1/Fs;
sig = 3*sin(2*pi*20*t)+cos(2*pi*2*t);
fb = cwtfilterbank('SignalLength',length(sig),'SamplingFrequency',Fs,'Boundary','periodic');
plot(t,sig)
xlabel('Time (sec)')
title('Signal')
```



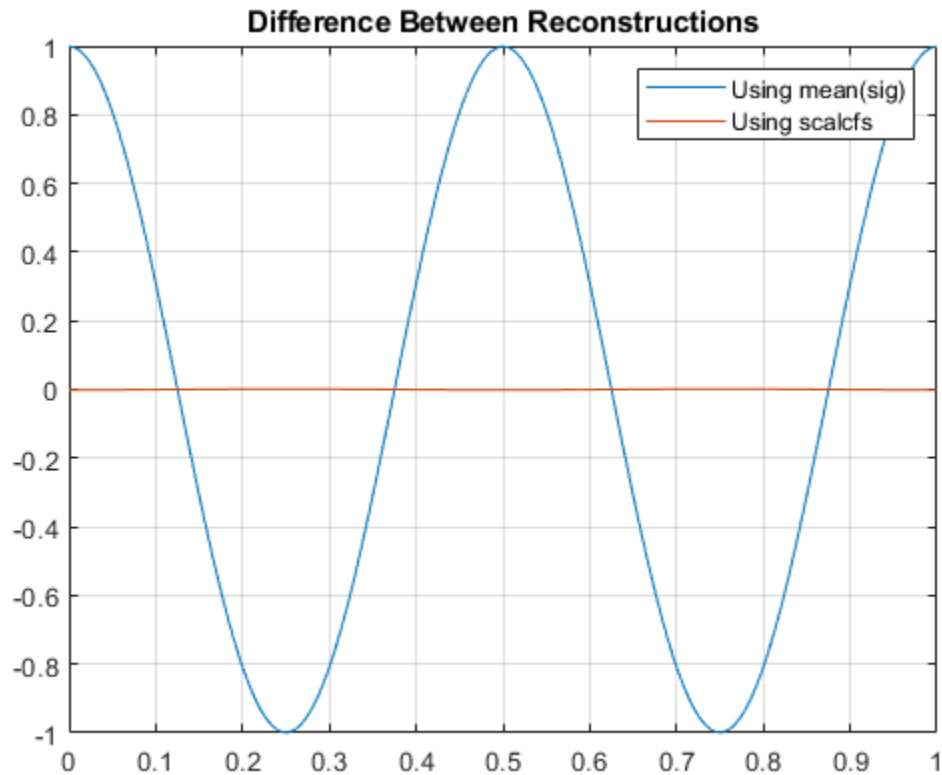
Take the CWT of the signal. Return the wavelet and scaling coefficients.

```
[cfs,~,~,scalcfs] = wt(fb,sig);
```

Reconstruct the signal two ways. First use the mean of the signal, then use the scaling coefficients. Plot the difference between the original signal and both reconstructions.

```
xrec0 = icwt(cfs,'SignalMean',mean(sig));  
xrec1 = icwt(cfs,'ScalingCoefficients',scalcfs);  
plot(t,sig-xrec0)  
hold on  
plot(t,sig-xrec1)  
grid on
```

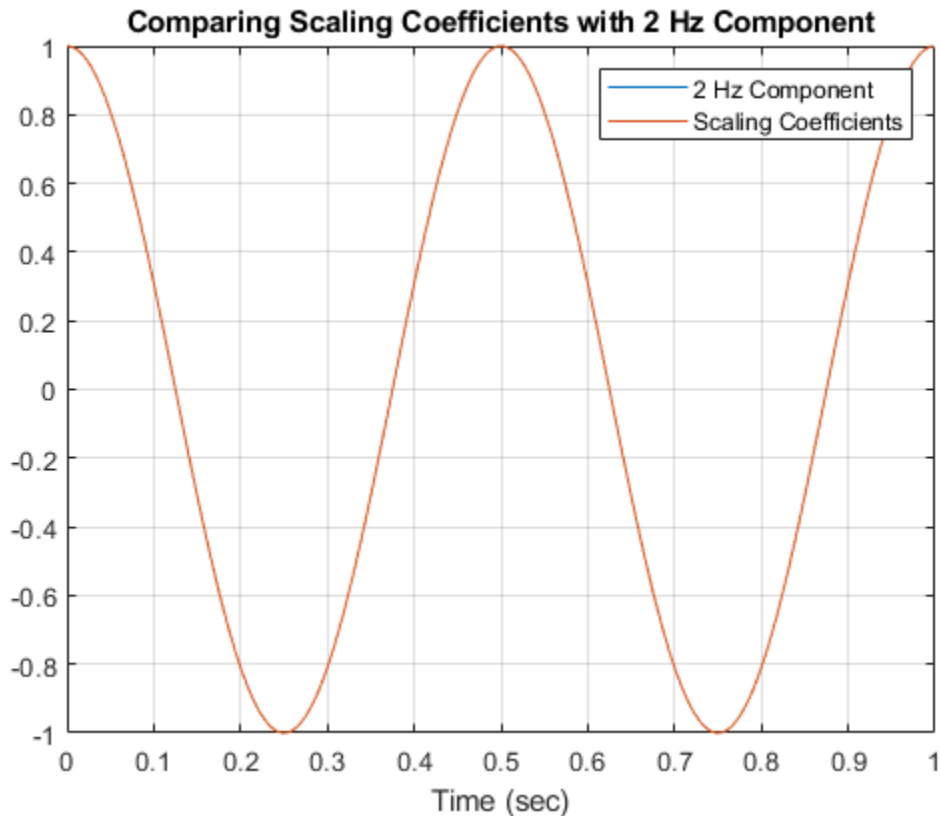
```
legend('Using mean(sig)', 'Using scalcfs')  
title('Difference Between Reconstructions')
```



Observe that using the scaling coefficients results in a significantly more accurate reconstruction. To investigate the source of the dramatic improvement, create a second signal consisting of the 2 Hz component of the original signal. Compare the scaling coefficients with the 2 Hz signal.

```
figure  
sig2hz = cos(2*pi*2*t);  
plot(t,sig2hz)  
hold on  
plot(t,scalcfs)  
grid on
```

```
title('Comparing Scaling Coefficients with 2 Hz Component')  
xlabel('Time (sec)')  
legend('2 Hz Component', 'Scaling Coefficients')
```



Note that the scaling coefficients and 2 Hz signal are virtually identical. Using the scaling coefficients helps with the reconstruction because the 2 Hz component is not representable by a wavelet with this sampling frequency and length.

Input Arguments

fb — Continuous wavelet transform filter bank
object

Continuous wavelet transform (CWT) filter bank, specified as a `cwtfilterbank` object.

x — Input signal

real- or complex-valued vector

Input signal, specified as a double-precision real- or complex-valued vector. `x` must have at least four samples.

Output Arguments

cfs — Continuous wavelet transform

matrix

Continuous wavelet transform, returned as a matrix of complex values. If `x` is real-valued, `cfs` is a 2-D matrix where each row corresponds to one scale. The column size of `cfs` is equal to the length of `x`. If `x` is complex-valued, `cfs` is a 3-D matrix, where the first page is the CWT for the positive scales (analytic part or counterclockwise component), and the second page is the CWT for the negative scales (anti-analytic part or clockwise component).

f — Frequencies

vector

Frequencies, returned as a vector, corresponding to the scales (rows) of `cfs` if the `'SamplingPeriod'` is not specified in `fb`. If you specify a sampling frequency, `f` is in hertz. If you do not specify a frequency, `f` is in cycles/sample.

Data Types: `double`

p — Periods

array

Periods, returned as an array of durations, corresponding to the scales (rows) of `cfs` if you specify a sampling period in `fb`. `p` has the same units and format as the duration scalar sampling period.

Data Types: `duration`

coi — Cone of influence

array of doubles | array of durations

Cone of influence for the CWT, returned as either an array of doubles or array of durations. The cone of influence indicates where edge effects occur in the CWT. If you specify a sampling frequency, `coi` is an array of doubles in the same units as `f`. If you specify a sampling period, `coi` is an array of durations with the same format property as the sampling period. Due to the edge effects, give less credence to areas that are outside or overlap the cone of influence.

For additional information, see “Boundary Effects and the Cone of Influence”.

Data Types: `double` | `duration`

`scalcf`s — Scaling coefficients

real- or complex-valued vector

Scaling coefficients for the wavelet transform, returned as a vector with length equal to the length of `x`. If `x` is real-valued, `scalcf`s is real-valued. If `x` is complex-valued, `scalcf`s is complex-valued.

Data Types: `double`

See Also

`cwt` | `cwtfilterbank` | `icwt`

Topics

“Boundary Effects and the Cone of Influence”

Introduced in R2018a

wtbo

WTBO constructor

Syntax

OBJ = wtbo

OBJ = wtbo(USERDATA)

Description

OBJ = wtbo returns a WTBO object. Any object in the Wavelet Toolbox software is parented by a WTBO object.

With OBJ = wtbo(USERDATA) you can set a userdata field.

Class WTBO (Parent class: none)

Fields

wtboInfo	Object information (not used in the current version of the toolbox)
ud	Userdata field

Introduced before R2006a

wtbxmngr

Wavelet Toolbox manager

Syntax

```
wtbxmngr(OPTION)  
V = wtbxmngr('version')
```

Description

wtbxmngr or wtbxmngr('version') displays the current version of Wavelet Toolbox software.

wtbxmngr(OPTION) sets a toolbox option. Available options are

Option	Description
'LargeFonts'	Sets the size of future-created figures to use large fonts.
'DefaultSize'	Restores the default figure size for future- created figures.
'FigRatio'	Returns the current figure ratio value.
'FigRatio',ratio	Changes the size of future-created figures by multiplying the default size by the specified ratio, where ratio must be between 0.75 and 1.25.

V = wtbxmngr('version') saves the current version of the toolbox to variable V.

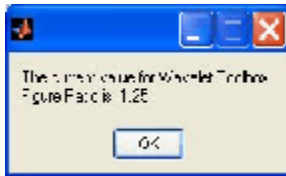
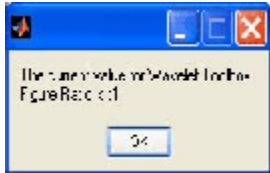
Examples

```
wtbxmngr('version')
```

```
*****  
** Wavelet Toolbox Version: V3.1 **  
*****
```



```
wtbxmng('FigRatio') % Display the current figure ratio
wtbxmng('FigRatio',1.25) % Set the figure ratio to 1.25
wtbxmng('FigRatio') % Display the current figure ratio
wtbxmng('DefaultSize') % Return to the default figure ratio
```



Introduced before R2006a

wthcoef

1-D wavelet coefficient thresholding

Syntax

```
NC = wthcoef('d',C,L,N,P)
NC = wthcoef('d',C,L,N)
NC = wthcoef('a',C,L)
NC = wthcoef('t',C,L,N,T,SORH)
```

Description

`wthcoef` thresholds wavelet coefficients for the denoising or compression of a 1-D signal.

`NC = wthcoef('d',C,L,N,P)` returns coefficients obtained from the wavelet decomposition structure `[C,L]` (see `wavedec` for more information), by rate compression defined in vectors `N` and `P`. `N` contains the detail levels to be compressed and `P` the corresponding percentages of lower coefficients to be set to zero. `N` and `P` must be of same length. Vector `N` must be such that $1 \leq N(i) \leq \text{length}(L) - 2$.

`NC = wthcoef('d',C,L,N)` returns coefficients obtained from `[C,L]` by setting all the coefficients of detail levels defined in `N` to zero.

`NC = wthcoef('a',C,L)` returns coefficients obtained by setting approximation coefficients to zero.

`NC = wthcoef('t',C,L,N,T,SORH)` returns coefficients obtained from the wavelet decomposition structure `[C,L]` by soft (if `SORH = 's'`) or hard (if `SORH = 'h'`) thresholding (see `wthresh` for more information) defined in vectors `N` and `T`. `N` contains the detail levels to be thresholded and `T` the corresponding thresholds. `N` and `T` must be of the same length.

`[NC,L]` is the modified wavelet decomposition structure.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

wavedec | wthresh

Introduced before R2006a

wthcoef2

Wavelet coefficient thresholding 2-D

Syntax

```
NC = wthcoef2('type',C,S,N,T,SORH)
NC = wthcoef2('type',C,S,N)
NC = wthcoef2('a',C,S)
NC = wthcoef2('t',C,S,N,T,SORH)
```

Description

wthcoef2 is a two-dimensional de-noising and compression oriented function.

For `'type' = 'h'` (`'v'` or `'d'`), `NC = wthcoef2('type',C,S,N,T,SORH)` returns the horizontal (vertical or diagonal, respectively) coefficients obtained from the wavelet decomposition structure `[C,S]` (see `wavedec2` for more information), by soft (if `SORH = 's'`) or hard (if `SORH = 'h'`) thresholding defined in vectors `N` and `T`. `N` contains the detail levels to be thresholded and `T` the corresponding thresholds. `N` and `T` must be of the same length. The vector `N` must be such that $1 \leq N(i) \leq \text{size}(S,1) - 2$.

For `'type' = 'h'` (`'v'` or `'d'`), `NC = wthcoef2('type',C,S,N)` returns the horizontal (vertical or diagonal, respectively) coefficients obtained from `[C,S]` by setting all the coefficients of detail levels defined in `N` to zero.

`NC = wthcoef2('a',C,S)` returns the coefficients obtained by setting approximation coefficients to zero.

`NC = wthcoef2('t',C,S,N,T,SORH)` returns the detail coefficients obtained from the wavelet decomposition structure `[C,S]` by soft (if `SORH = 's'`) or hard (if `SORH = 'h'`) thresholding (see `wthresh` for more information) defined in vectors `N` and `T`. `N` contains the detail levels to be thresholded and `T` the corresponding thresholds which are applied in the three detail orientations. `N` and `T` must be of the same length.

`[NC,S]` is the modified wavelet decomposition structure.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

wavedec2 | wthresh

Introduced before R2006a

wthresh

Soft or hard thresholding

Syntax

```
Y = wthresh(X, sorh, T)
```

Description

`Y = wthresh(X, sorh, T)` returns the soft or hard thresholding, indicated by `sorh`, of the vector or matrix `X`. `T` is the threshold value.

Examples

Hard and Soft Thresholding

Generate a signal and set a threshold.

```
y = linspace(-1,1,100);  
thr = 0.4;
```

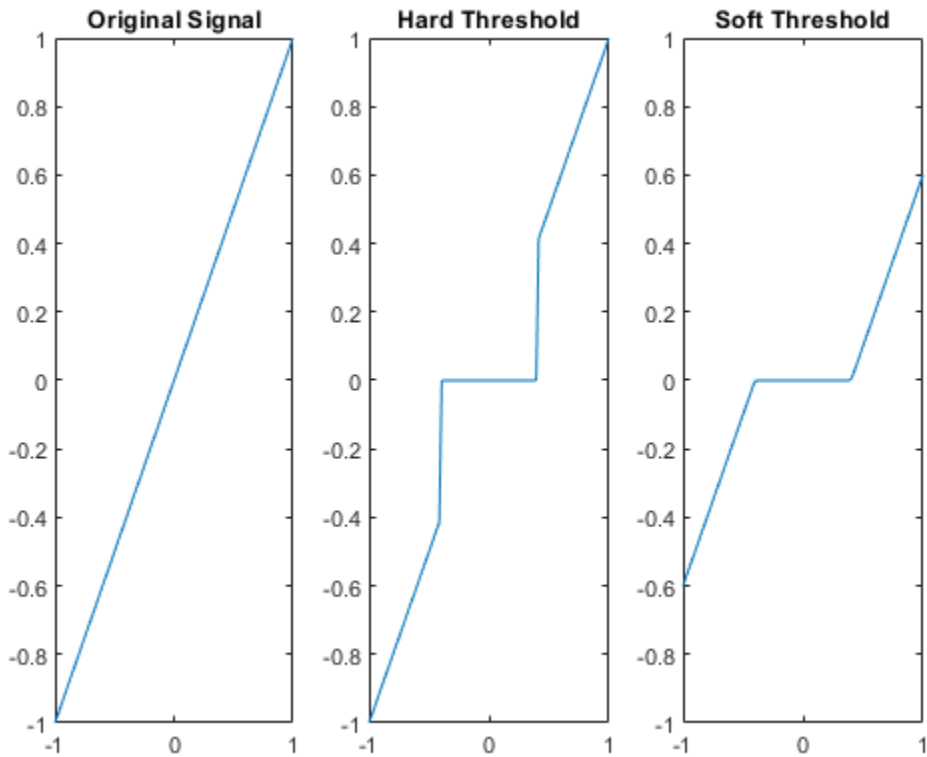
Perform hard and soft thresholding.

```
ythard = wthresh(y, 'h', thr);  
ytsoft = wthresh(y, 's', thr);
```

Plot the results and compare with the original signal.

```
subplot(1,3,1)  
plot(y,y)  
ylim([-1 1])  
title('Original Signal')  
subplot(1,3,2)  
plot(y,ythard)  
ylim([-1 1])
```

```
title('Hard Threshold')
subplot(1,3,3)
plot(y,ytsoft)
ylim([-1 1])
title('Soft Threshold')
```



Input Arguments

X — Input data

real-valued vector or matrix

Input data to threshold, specified as a real-valued vector or matrix.

Data Types: double

sorh — Type of thresholding

's' | 'h'

Type of thresholding to perform:

- 's' — Soft thresholding
- 'h' — Hard thresholding

T — Threshold value

positive real number

Threshold value, specified as a positive real number.

Output Arguments

Y — Thresholded data

real-valued vector or matrix

Thresholded data, returned as a real-valued vector or matrix. Y has the same dimensions as X.

Algorithms

If sorh is 's', Y is the soft thresholding of X: $Y = \text{sign}(X) \cdot (|X| - T)_+$ where

$$(x)_+ = \begin{cases} x & \text{if } x \geq 0 \\ 0 & \text{otherwise} \end{cases}$$

Soft thresholding is wavelet shrinkage.

If sorh is 'h', Y is the hard thresholding of X: $Y = X \cdot \mathbf{1}_{(|X| > T)}$ where

$$\mathbf{1}_{(|X| > T)} = \begin{cases} 1 & \text{if } |X| > T \\ 0 & \text{otherwise} \end{cases}$$

Hard thresholding is cruder than soft thresholding.

Extended Capabilities

C/C++ Code Generation

Generate C and C++ code using MATLAB® Coder™.

See Also

Functions

`wden` | `wdencmp` | `wdenoise` | `wpdencmp`

Apps

Wavelet Signal Denoiser

Introduced before R2006a

wthrmngr

Threshold settings manager

`wthrmngr` returns a global threshold or level-dependent thresholds for wavelet-based denoising and compression. The function derives thresholds from the wavelet coefficients in a wavelet decomposition.

The thresholds are used by Wavelet Toolbox denoising and compression tools, such as command-line functions and the **Wavelet Analyzer** app.

Syntax

```
thr = wthrmngr(opt,method,C,L)
thr = wthrmngr(opt,method,C,L,alpha)
thr = wthrmngr(opt,method,C,L,scale)

thr = wthrmngr(opt,method,swtdec,alpha)
thr = wthrmngr(opt,method,swtdec,scale)

thr = wthrmngr(opt,method,wpt)
thr = wthrmngr(opt,'rem_n0',X)
```

Description

`thr = wthrmngr(opt,method,C,L)` returns the threshold for the $[C,L]$ wavelet decomposition of the signal or image to compress or denoise. For signals, $[C,L]$ is the output of `wavedec`. For images, $[C,L]$ is the output of `wavedec2`.

`thr = wthrmngr(opt,method,C,L,alpha)` returns the $[C,L]$ wavelet decomposition threshold using the sparsity parameter `alpha`. For signals, $[C,L]$ is the output of `wavedec`. For images, $[C,L]$ is the output of `wavedec2`.

To learn more about `alpha`, see `wdcbm` or `wdcbm2` for compression, and `wbmpen` for denoising.

`thr = wthrmngr(opt,method,C,L,scale)` returns the $[C,L]$ wavelet decomposition threshold using the type of multiplicative threshold rescaling specified in `scale`. For signals, $[C,L]$ is the output of `wavedec`. For images, $[C,L]$ is the output of `wavedec2`.

The `'rigrsure'`, `'heursure'`, and `'minimaxi'` denoising methods are only applicable to signals.

To learn more about multiplicative threshold rescaling, see `wden`.

`thr = wthrmngr(opt,method,swtdec,alpha)` returns the level-dependent threshold for the stationary wavelet decomposition, `swtdec`, of the signal or image to denoise. `alpha` specifies the sparsity parameter (see `wbmpen`). For signals, `swtdec` is the output of `swt`. For images, `swtdec` is the output of `swt2`.

Thresholds are derived from a subset of the coefficients in the stationary wavelet decomposition. For more information, see "Coefficient Selection" on page 1-1653.

`thr = wthrmngr(opt,method,swtdec,scale)` returns the level-dependent threshold for the stationary wavelet decomposition using the type of multiplicative threshold rescaling specified in `scale`. For signals, `swtdec` is the output of `swt`. For images, `swtdec` is the output of `swt2`.

Thresholds are derived from a subset of the coefficients in the stationary wavelet decomposition. For more information, see "Coefficient Selection" on page 1-1653.

The `'rigrsure'`, `'heursure'`, and `'minimaxi'` denoising methods apply only to signals.

To learn more about multiplicative threshold rescaling, see `wden`.

`thr = wthrmngr(opt,method,wpt)` returns the global threshold for the wavelet packet decomposition, `wpt`, of the signal or image to compress or denoise.

`thr = wthrmngr(opt,'rem_n0',X)` returns the global threshold to compress the signal or image, `X`, using the specified wavelet option and method `'rem_n0'`.

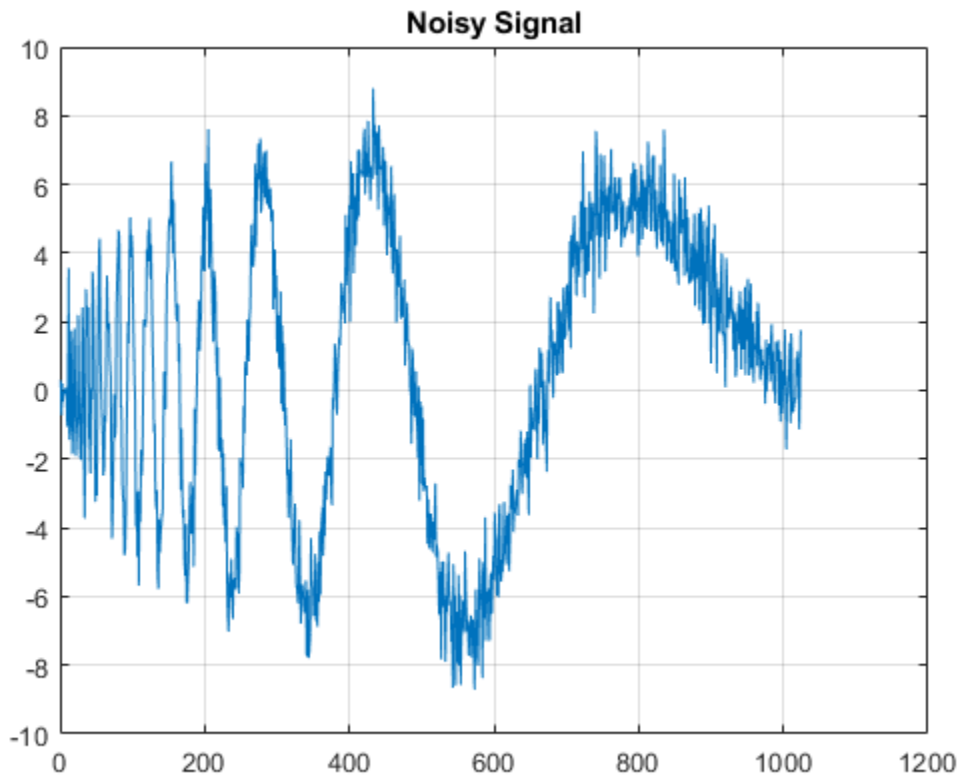
If `opt` is `'dw1dcompGBL'` or `'dw2dcompGBL'`, thresholds are based on the finest-scale wavelet coefficients obtained using the Haar wavelet. If `opt` is `'wp1dcompGBL'` or `'wp2dcompGBL'`, thresholds are based on the finest-scale wavelet packet coefficients obtained using the Haar wavelet.

Examples

Global Threshold — Discrete Wavelet Decomposition

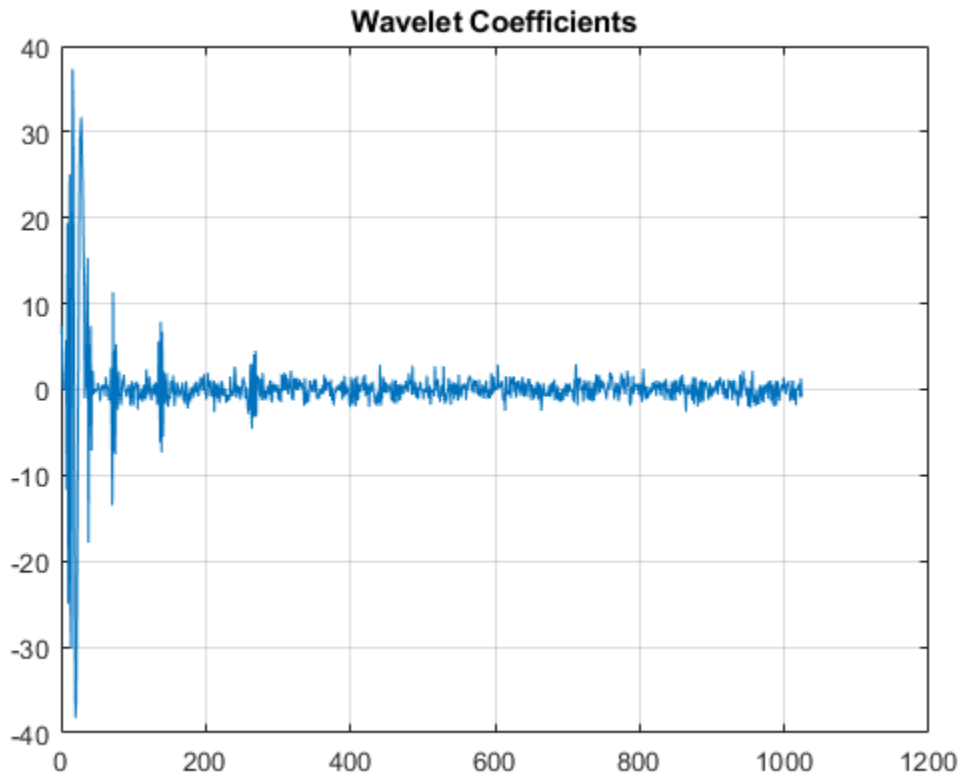
Load and plot a noisy signal.

```
load noisdopp  
plot(noisdopp)  
grid on  
title('Noisy Signal')
```



Generate a level 5 wavelet decomposition of the noisy signal using the order 4 Daubechies wavelet. Plot the coefficients.

```
[c,l] = wavedec(noisdopp,5,'db4');
plot(c)
grid on
title('Wavelet Coefficients')
```



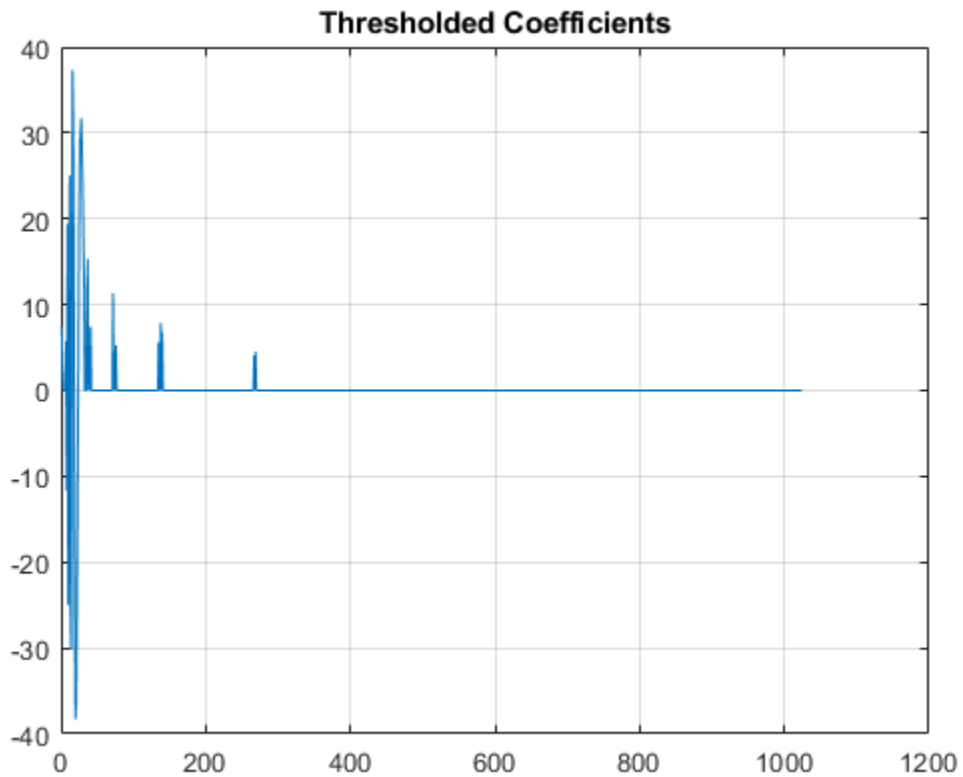
Determine a global threshold for compressing the signal.

```
thr = wthrmngr('dw1dcompGBL','bal_sn',c,l);
```

The index of the first wavelet detail coefficient in c is $l(1)+1$. Apply the threshold to all the detail coefficients. Plot the thresholded coefficients. Observe that most of the coefficients have been set to 0.

```
c(l(1)+1:end) = c(l(1)+1:end).*(c(l(1)+1:end)>thr);
plot(c)
```

```
grid on  
title('Thresholded Coefficients')
```



Reconstruct the signal from the thresholded coefficients. Plot the reconstruction.

```
xrec = waverec(c,l,'db4');  
plot(xrec)  
grid on  
title('Compressed Signal')
```

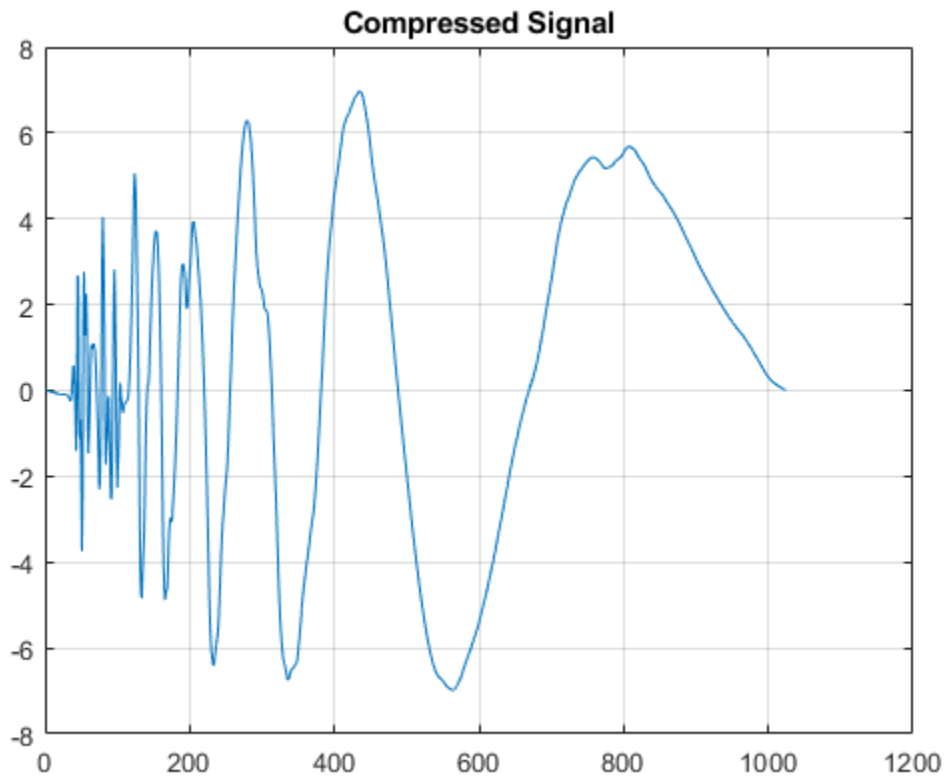


Image Compression — Birgé-Massart Thresholds

Compress an image using the Birgé-Massart strategy.

Load an image and add white Gaussian noise. For purposes of reproducibility, set the random seed to the default value.

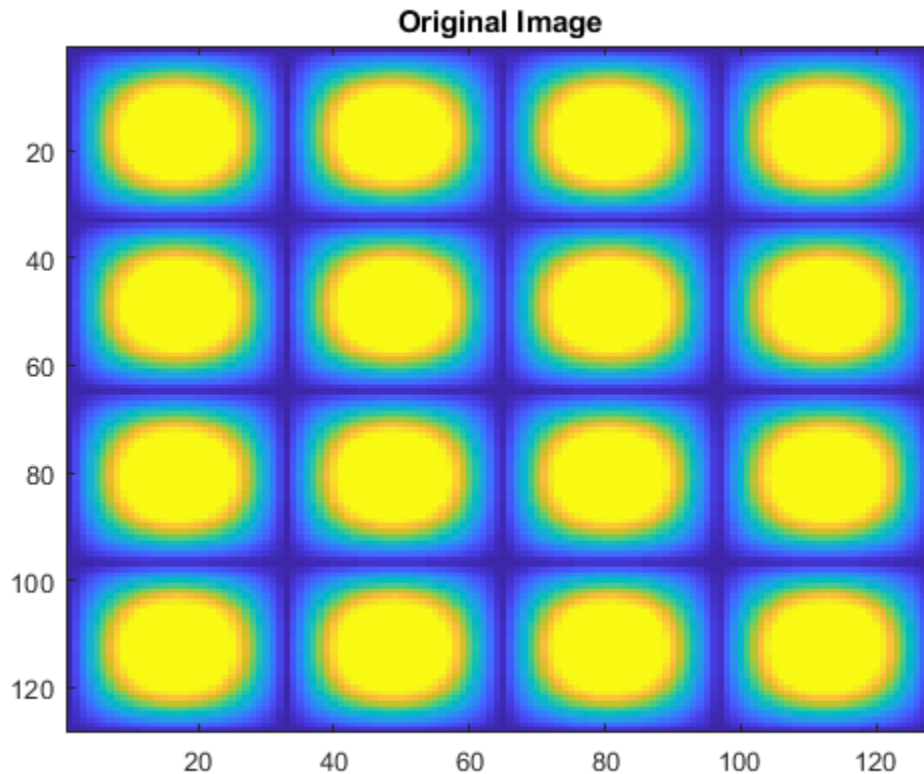
```
rng default
load sinsin
x = X+18*randn(size(X));
```

Obtain the 2-D discrete wavelet transform down to level 3 using the Daubechies least-asymmetric wavelet with 4 vanishing moments. Obtain the compression thresholds using the Birgé-Massart strategy with sparsity parameter, α , equal to 2.

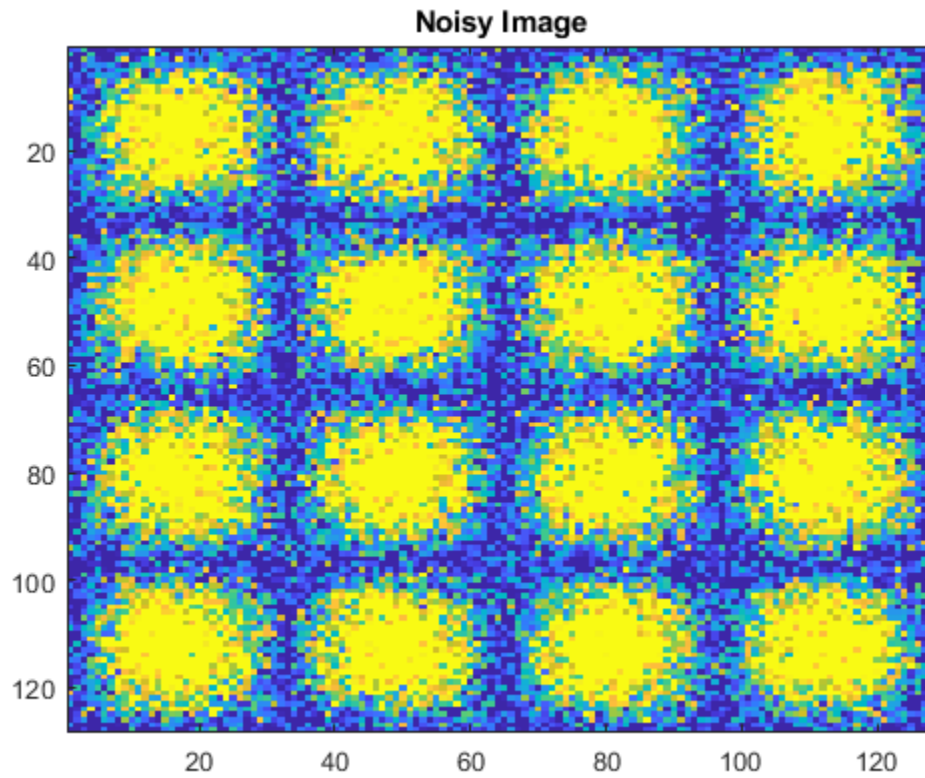
```
[C,L] = wavedec2(x,3,'sym4');  
alpha = 2;  
THR = wthrmngr('dw2dcompLVL','scarcehi',C,L,alpha);
```

Compress the image and display the result.

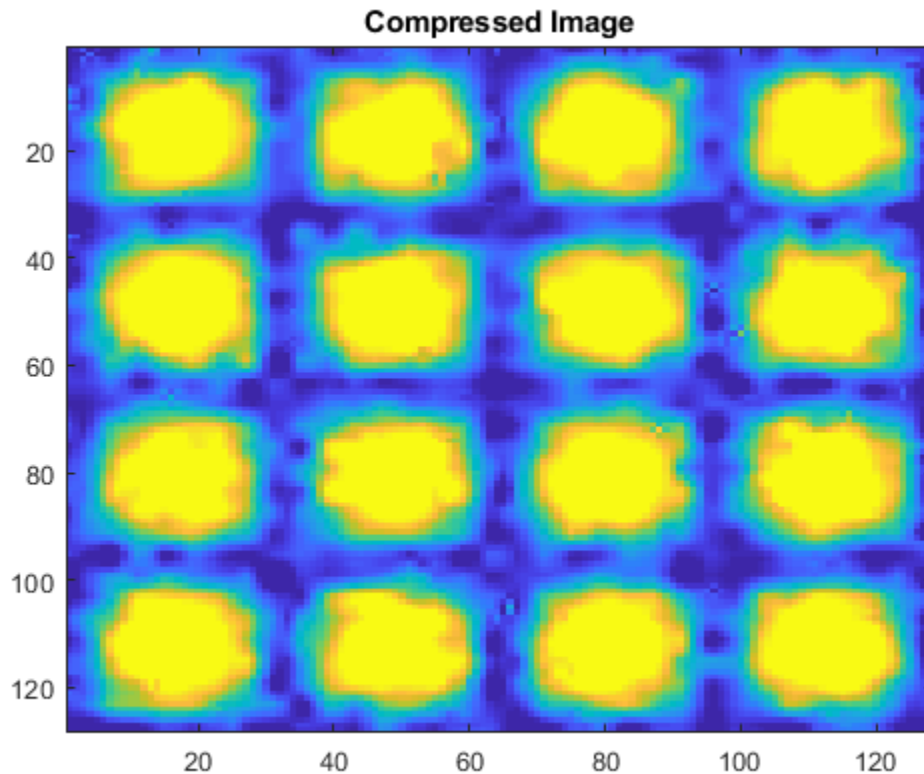
```
xd = wdencmp('lvd',x,'sym4',3,THR,'s');  
image(X)  
title('Original Image')
```




```
figure
image(x)
title('Noisy Image')
```



```
figure
image(xd)
title('Compressed Image')
```



Level-Dependent Threshold – Stationary Wavelet Transform

This example uses a level-dependent threshold derived from the wavelet coefficients at each scale to implement hard thresholding with the stationary wavelet transform.

Load the noisy blocks signal. Obtain the stationary wavelet transform down to level 5 by using the Haar wavelet.

```
load noisbloc
L = 5;
swc = swt(noisbloc,L,'haar');
```

Make a copy of the wavelet transform coefficients. Determine the Donoho-Johnstone universal threshold based on the detail coefficients for each scale. Using the 'mln' option, `wthrmngr` returns a 1-by-L vector, with every element equal to the universal threshold for the corresponding scale.

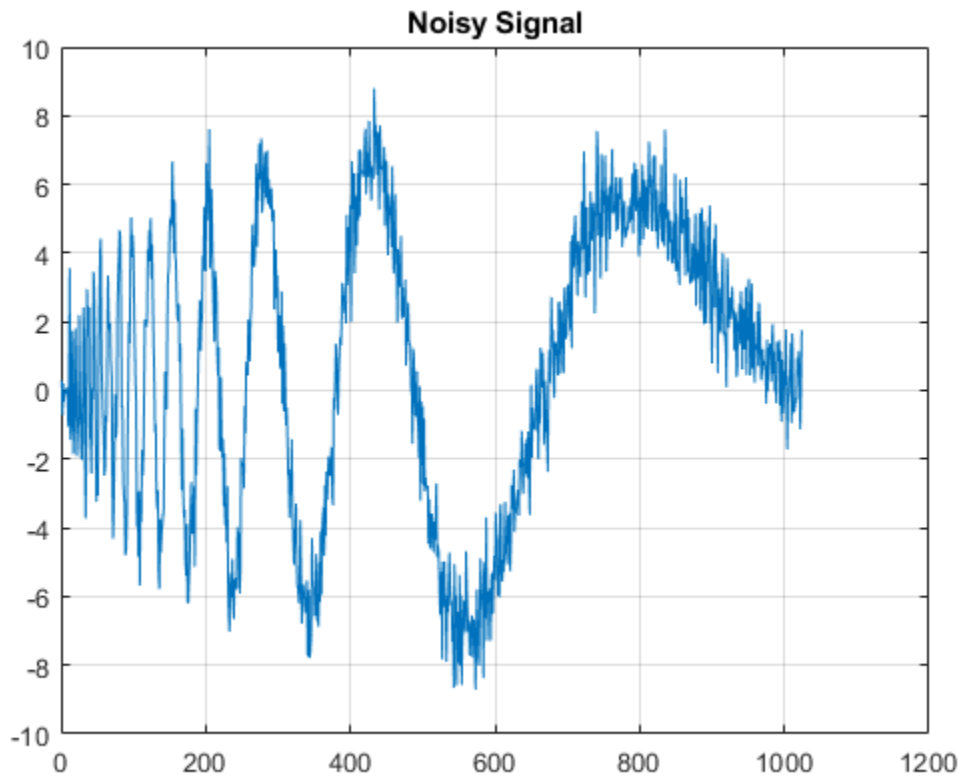
```
swcnew = swc;  
ThreshML = wthrmngr('swlddenoLVL','sqrtwolog',swc,'mln');
```

Use the universal thresholds to implement hard thresholding. The thresholds are applied in a scale-dependent manner.

```
for jj = 1:L  
    swcnew(jj,:) = wthresh(swc(jj,:), 'h', ThreshML(jj));  
end
```

Invert the stationary wavelet transform on the thresholded coefficients, `swcnew`. Plot the original signal and the denoised signal for comparison.

```
noisbloc_denoised = iswt(swcnew, 'haar');  
plot(noisbloc)  
hold on  
plot(noisbloc_denoised, 'r', 'linewidth', 2)  
legend('Original', 'Denoised')
```

Generate a level 3 wavelet packet decomposition of the noisy signal using the order 4 Daubechies wavelet.

```
T = wpdec(noisdopp,3,'db4');
```

Determine a global threshold for denoising the signal.

```
thr = wthrmngr('wp1ddenoGBL','sqrtwologuwn',T);
```

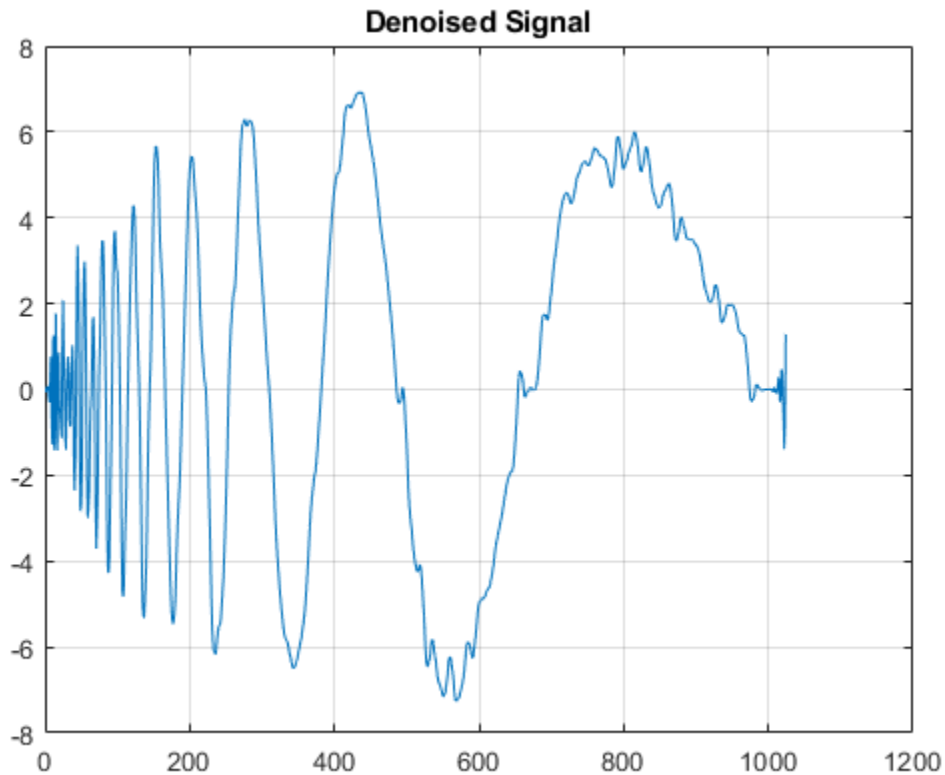
Obtain the leaves from the wavelet packet decomposition tree T and apply the threshold to the leaves. Use hard thresholding.

```
T1 = T;  
sorh = 'h';
```

```
cfs = read(T,'data');  
cfs = wthresh(cfs,sorh,thr);  
T1 = write(T1,'data',cfs);
```

Reconstruct the denoised signal from the thresholded coefficients. Plot the reconstruction.

```
xrec = wprec(T1);  
plot(xrec)  
grid on  
title('Denoised Signal')
```



Level-Independent Threshold — Stationary Wavelet Transform

This example uses a level-independent threshold based on the finest-scale wavelet coefficients to implement hard thresholding with the stationary wavelet transform.

Load the noisy blocks signal. Obtain the stationary wavelet transform down to level 5 by using the Haar wavelet.

```
load noisbloc
L = 5;
swc = swt(noisbloc,L,'haar');
```

Make a copy of the wavelet transform coefficients. Determine the Donoho-Johnstone universal threshold based on the first-level detail coefficients. Using the 'sln' option, wthrmngr returns a 1-by-L vector, with every element equal to the same value. Take the mean of the vector to obtain a scalar threshold.

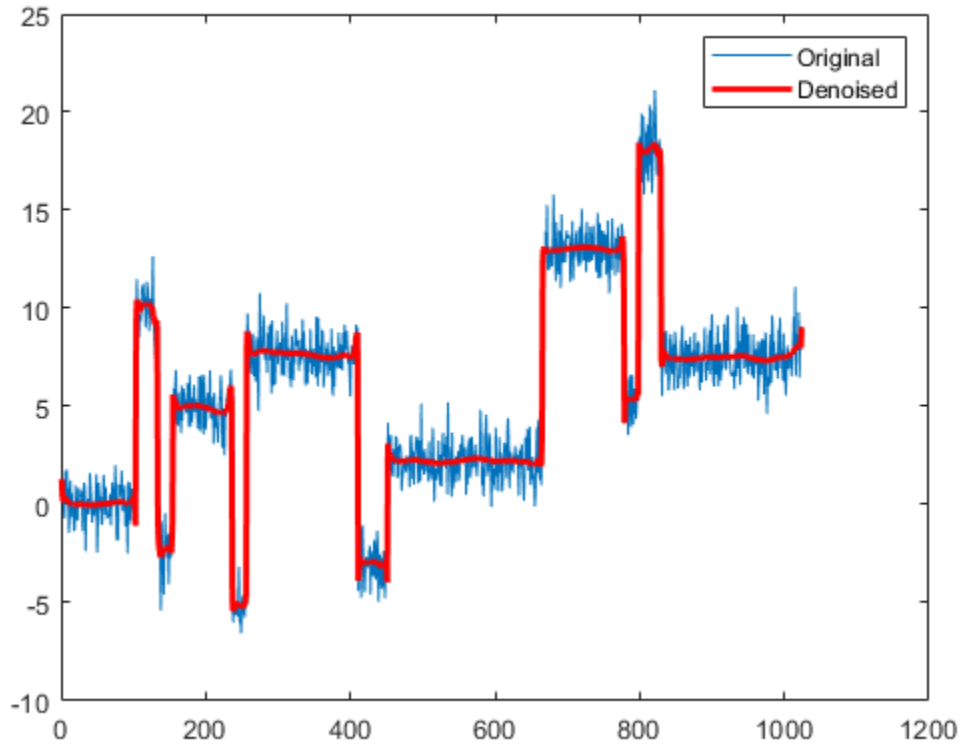
```
swcnew = swc;
ThreshSL = mean(wthrmngr('swlddenoLVL','sqrtwolog',swc,'sln'));
```

Use the universal threshold to implement hard thresholding. The same threshold is applied to the wavelet coefficients at every level.

```
for jj = 1:L
    swcnew(jj,:) = wthresh(swc(jj,:), 'h',ThreshSL);
end
```

Invert the stationary wavelet transform on the thresholded coefficients, swcnew. Plot the original signal and the denoised signal for comparison.

```
noisbloc_denoised = iswt(swcnew,'haar');
plot(noisbloc)
hold on
plot(noisbloc_denoised,'r','linewidth',2)
legend('Original','Denoised')
```



Input Arguments

opt — Type and dimension of compression or denoising

'dw1dcompGBL' | 'dw1dcompLVL' | 'dw1ddenoLVL' | 'sw1ddenoLVL' |
'dw2dcompGBL' | 'dw2dcompLVL' | ...

Type and dimension of compression or denoising, specified as one of the values listed in the tables that follow. `wthrmngr` returns thresholds appropriate for the option you specify.

With a discrete wavelet or wavelet packet decomposition of the data, you can compress or denoise that data. With a stationary wavelet decomposition of the data, you can only denoise the data.

For an explanation of which coefficients are used to determine the thresholds, see “Coefficient Selection” on page 1-1653.

1-D Discrete Wavelet Decomposition Options

In these options, X is the signal, the wavelet coefficients are in the vector C , and the lengths of the coefficient vectors are in L . The argument α is the sparsity parameter, and $scale$ defines the multiplicative threshold rescaling.

For additional information regarding the wavelet decomposition, see `wavedec`. To learn more about α and $scale$, see `wdcbm` and `wden` respectively.

opt	Description	Valid Syntaxes
'dw1dcompG BL'	1-D compression using a global threshold	<ul style="list-style-type: none"> thr = wthrmngr('dw1dcompGBL','rem_n0',X) thr = wthrmngr('dw1dcompGBL','bal_sn',C,L)
'dw1dcompL VL'	1-D compression using level-dependent thresholds	<ul style="list-style-type: none"> thr = wthrmngr('dw1dcompLVL','scarcehi',C,L,alpha), where $2.5 < \alpha < 10$ thr = wthrmngr('dw1dcompLVL','scarceme',C,L,alpha), where $1.5 < \alpha < 2.5$ thr = wthrmngr('dw1dcompLVL','scarcelo',C,L,alpha), where $1 < \alpha < 2$

opt	Description	Valid Syntaxes
'dw1ddenoLVL'	1-D denoising using level-dependent thresholds	<ul style="list-style-type: none"> • thr = wthrmngr('dw1ddenoLVL','sqrtwolog',C,L,scale) • thr = wthrmngr('dw1ddenoLVL','rigrsure',C,L,scale) • thr = wthrmngr('dw1ddenoLVL','heursure',C,L,scale) • thr = wthrmngr('dw1ddenoLVL','minimaxi',C,L,scale) • thr = wthrmngr('dw1ddenoLVL','penalhi',C,L,alpha), where $2.5 < \alpha < 10$ • thr = wthrmngr('dw1ddenoLVL','penalme',C,L,alpha), where $1.5 < \alpha < 2.5$ • thr = wthrmngr('dw1ddenoLVL','penallo',C,L,alpha), where $1 < \alpha < 2$

2-D Discrete Wavelet Decomposition Options

In these options, *X* is the data, the wavelet coefficients are in the vector *C*, and the size of the coefficient matrices are in *L*. The argument *alpha* is the sparsity parameter, and *scale* defines the multiplicative threshold rescaling.

For additional information regarding the wavelet decomposition, see `wavedec2`. To learn more about *alpha* and *scale*, see `wdcbm2` and `wden` respectively.

opt	Description	Valid Syntaxes
'dw2dcompGBL'	2-D compression using a global threshold	<ul style="list-style-type: none"> • thr = wthrmngr('dw2dcompGBL', 'rem_n0', X) • thr = wthrmngr('dw2dcompGBL', 'bal_sn', C, L) • thr = wthrmngr('dw2dcompGBL', 'sqrtbal_sn', C, L)
'dw2dcompLVL'	2-D compression using level-dependent thresholds	<ul style="list-style-type: none"> • thr = wthrmngr('dw2dcompLVL', 'scarcehi', C, L, alpha), where $2.5 < \alpha < 10$ • thr = wthrmngr('dw2dcompLVL', 'scarce', C, L, alpha), where $1.5 < \alpha < 2.5$ • thr = wthrmngr('dw2dcompLVL', 'scarcelo', C, L, alpha), where $1 < \alpha < 2$
'dw2ddenoLVL'	2-D denoising using level-dependent thresholds	<ul style="list-style-type: none"> • thr = wthrmngr('dw2ddenoLVL', 'sqrtbal_sn', C, L) • thr = wthrmngr('dw2ddenoLVL', 'penalhi', C, L, alpha), where $2.5 < \alpha < 10$ • thr = wthrmngr('dw2ddenoLVL', 'penalme', C, L, alpha), where $1.5 < \alpha < 2.5$ • thr = wthrmngr('dw2ddenoLVL', 'penallo', C, L, alpha), where $1 < \alpha < 2$ • thr = wthrmngr('dw2ddenoLVL', 'sqtwo', C, L, scale)

1-D Wavelet Packet Decomposition Options

In these options, *X* is the signal and *wpt* is the wavelet packet decomposition structure of the signal.

For additional information regarding the wavelet packet decomposition, see *wpdec*.

opt	Description	Valid Syntaxes
'wp1dcompGBL'	1-D compression using a global threshold	<ul style="list-style-type: none"> thr = wthrmngr('wp1dcompGBL','rem_n0',X) thr = wthrmngr('wp1dcompGBL','bal_sn',wpt)
'wp1ddenoGBL'	1-D denoising using a global threshold	<ul style="list-style-type: none"> thr = wthrmngr('wp1ddenoGBL','sqrtwologuwn',wpt) thr = wthrmngr('wp1ddenoGBL','sqrtwologswn',wpt) thr = wthrmngr('wp1ddenoGBL','bal_sn',wpt) thr = wthrmngr('wp1ddenoGBL','penalhi',wpt) <p>The <i>wpbmpen</i> function is used with the tuning parameter ALPHA = 6.25.</p> <ul style="list-style-type: none"> thr = wthrmngr('wp1ddenoGBL','penalme',wpt) <p>The <i>wpbmpen</i> function is used with the tuning parameter ALPHA = 2.</p> <ul style="list-style-type: none"> thr = wthrmngr('wp1ddenoGBL','penallo',wpt) <p>The <i>wpbmpen</i> function is used with the tuning parameter ALPHA = 1.5.</p>

2-D Wavelet Packet Decomposition Options

In these options, X is the data and wpt is the wavelet packet decomposition structure of the data.

For additional information regarding the wavelet packet decomposition, see `wpdec2`.

opt	Description	Valid Syntaxes
'wp2dcompG BL'	2-D compression using a global threshold	<ul style="list-style-type: none"> • thr = wthrmngr('wp2dcompGBL', 'rem_n0', X) • thr = wthrmngr('wp2dcompGBL', 'bal_sn', wpt) • thr = wthrmngr('wp2dcompGBL', 'sqrtbal_sn', wpt)

opt	Description	Valid Syntaxes
'wp2ddenoGBL'	2-D denoising using a global threshold	<ul style="list-style-type: none"> • thr = wthrmngr('wp2ddenoGBL','sqrtwologuwn',wpt) • thr = wthrmngr('wp2ddenoGBL','sqrtwologswn',wpt) • thr = wthrmngr('wp2ddenoGBL','sqrtbal_sn',wpt) • thr = wthrmngr('wp2ddenoGBL','penalhi',wpt) <p>The wpbmpen function is used with the tuning parameter ALPHA = 6.25.</p> <ul style="list-style-type: none"> • thr = wthrmngr('wp2ddenoGBL','penalme',wpt) <p>The wpbmpen function is used with the tuning parameter ALPHA = 2.</p> <ul style="list-style-type: none"> • thr = wthrmngr('wp2ddenoGBL','penallo',wpt) <p>The wpbmpen function is used with the tuning parameter ALPHA = 1.5.</p>

1-D Stationary Wavelet Decomposition Options

Denoising using level-dependent thresholds is the only option available for a 1-D stationary wavelet decomposition, `swtdec`. In this option, `alpha` is a sparsity parameter and `scale` defines the multiplicative threshold rescaling.

For more information regarding the stationary wavelet decomposition, see `swt`. To learn more about `alpha` and `scale`, see `wbmpen` and `wden` respectively.

opt	Valid Syntaxes
'swlddenoLVL'	<ul style="list-style-type: none"> • thr = wthrmngr('swlddenoLVL','sqrtwolog',swtdec,scale) • thr = wthrmngr('swlddenoLVL','rigrsure',swtdec,scale) • thr = wthrmngr('swlddenoLVL','heursure',swtdec,scale) • thr = wthrmngr('swlddenoLVL','minimaxi',swtdec,scale) • thr = wthrmngr('swlddenoLVL','penalhi',swtdec,alpha), where $2.5 < \alpha < 10$ • thr = wthrmngr('swlddenoLVL','penalme',swtdec,alpha), where $1.5 < \alpha < 2.6$ • thr = wthrmngr('swlddenoLVL','penallo',swtdec,alpha), where $1 < \alpha < 2$

Thresholds are based on a subset of the coefficients in the stationary wavelet decomposition. See “Coefficient Selection” on page 1-1653 for additional information.

2-D Stationary Wavelet Decomposition Options

Denoising using level-dependent thresholds is the only option available for a 2-D stationary wavelet decomposition, swtdec. In this option, alpha is a sparsity parameter and scale defines the multiplicative threshold rescaling.

For more information regarding the stationary wavelet decomposition, see swt2. To learn more about alpha and scale, see wbmopen and wden respectively.

opt	Valid Syntaxes
'sw2ddenoLVL'	<ul style="list-style-type: none"> • thr = wthrmngr('sw2ddenoLVL','sqrtbal_sn',swtdec) • thr = wthrmngr('sw2ddenoLVL','penalhi',swtdec,alpha) where $2.5 < \alpha < 10$ • thr = wthrmngr('sw2ddenoLVL','penalme',swtdec,alpha) where $1.5 < \alpha < 2.5$ • thr = wthrmngr('sw2ddenoLVL','penallo',swtdec,alpha) where $1 < \alpha < 2$ • thr = wthrmngr('sw2ddenoLVL','sqtolog',swtdec,scale)

Thresholds are based on a subset of the coefficients in the stationary wavelet decomposition. See “Coefficient Selection” on page 1-1653 for additional information.

method – Thresholding method

'scarcehi' | 'scarceme' | 'scarcelo' | 'sqtolog' | 'sqtologuwn' | 'sqtologswn' | ...

Thresholding method, specified as one of the values listed here.

method	Description
'scarcehi'	Uses Birgé-Massart strategy on page 1-1654 for determining thresholds.
'scarceme'	Uses Birgé-Massart strategy for determining thresholds.
'scarcelo'	Uses Birgé-Massart strategy for determining thresholds.
'sqtolog'	Uses fixed-form universal threshold. See 'sqtolog' option in wden.
'sqtologuwn'	Uses fixed-form universal threshold. See 'sqtolog' option in wden when used with 'sln' option.

method	Description
'sqtwologswn'	Uses fixed-form universal threshold. See 'sqtwolog' option in wden when used with 'mln' option.
'rigsure'	Uses soft threshold estimator rule based on Stein's Unbiased Estimate of Risk. See 'SURE' option in wdenoise.
'heursure'	Uses mixture of 'rigsure' and 'sqtwolog'. See 'heursure' option in wden.
'minimaxi'	Uses a fixed threshold chosen which yields minimax performance. See 'Minimax' option in wdenoise.
'penalhi'	Used to define Birgé-Massart strategy on page 1-1654 for determining thresholds.
'penalme'	Used to define Birgé-Massart strategy for determining thresholds.
'penallo'	Used to define Birgé-Massart strategy for determining thresholds.
'rem_n0'	Returns a threshold close to 0. A typical THR value is <code>median(abs(coefficients))</code> .
'bal_sn'	Returns a threshold such that the percentages of retained energy and number of zeros are the same.
'sqrtbal_sn'	Returns a threshold equal to the square root of the value such that the percentages of retained energy and number of zeros are the same.

Data Types: char

X — Input data

real-valued vector | real-valued matrix

Input data, specified as a real-valued vector or real-valued matrix.

Data Types: double

C — Wavelet expansion coefficients

real-valued vector

Wavelet expansion coefficients of the data to be compressed or denoised, specified as a real-valued vector. If the data is one-dimensional, C is the output of `wavedec`. If the data is two-dimensional, C is the output of `wavedec2`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: double

L — Size of wavelet expansion coefficients

vector of positive integers | matrix of positive integers

Size of wavelet expansion coefficients of the signal or image to be compressed or denoised, specified as a vector or matrix of positive integers.

For signals, L is the output of `wavedec`. For images, L is the output of `wavedec2`.

Example: `[C,L] = wavedec(randn(1,1024),3,'db4')`

Data Types: double

alpha — Sparsity parameter

positive scalar

Sparsity parameter used for compressing or denoising data, specified as a positive scalar greater than 1 and less than 10. See `wdcbm`, `wdcbm2`, and `wbmpen` for additional information.

Data Types: double

scale — Multiplicative threshold rescaling

'one' | 'sln' | 'mln'

Multiplicative threshold rescaling, specified as one of the following:

- 'one' — No rescaling
- 'sln' — Rescaling using a single estimation of level noise based on first-level coefficients
- 'mln' — Rescaling using a level-dependent estimation of level noise

For more information, see `wden`.

swtdec — Stationary wavelet decomposition structure

real-valued matrix

Stationary wavelet decomposition structure of data to be compressed or denoised, specified as a real-valued matrix. If the data is one-dimensional, `swtdec` is the output of `swt`. If the data is two-dimensional, `swtdec` is the output of `swt2`.

```
Example: swtdec = swt2(randn(256),3,'db1')
```

Data Types: double

wpt — Wavelet packet decomposition structure

wavelet packet object structure

Wavelet packet decomposition structure of the data to be compressed or denoised. If the data is one-dimensional, `wpt` is the output of `wpdec`. If the data is two-dimensional, `wpt` is the output of `wpdec2`.

```
Example: wpt = wpdec(randn(1,1024),5,'db1')
```

Output Arguments

thr — Threshold

real-valued scalar | real-valued vector | real-valued matrix

Threshold, returned as a real-valued scalar for global thresholds, or a real-valued vector or matrix for level-dependent thresholds.

Data Types: double

Tips

- To denoise 1-D signals, consider using the **Wavelet Signal Denoiser**. The app visualizes and denoises real-valued 1-D signals using default parameters. You can also compare results. In addition, you can also recreate the denoised signal in your workspace by generating a MATLAB script, which uses the `wdenoise` function.

Algorithms

Coefficient Selection

A critically sampled wavelet or wavelet packet decomposition involves decimating coefficients by a factor of 2 at each stage of the decomposition. Decimation does not occur in the nondecimated stationary wavelet decomposition.

wthrmngr derives denoising and compression thresholds from the wavelet coefficients. For a critically sampled wavelet or wavelet packet decomposition, the option and method determine whether all wavelet coefficients or only the finest scale coefficients are used.

For the stationary wavelet decomposition, wthrmngr always uses a subset of the wavelet coefficients. When computing the denoising thresholds of an N-level stationary wavelet decomposition, the algorithm first subsamples the wavelet coefficients at level k by a factor of 2^k , for $k = 1, \dots, N$. The algorithm uses this subset of coefficients to determine the thresholds. Most of the coefficients in the stationary wavelet decomposition are not considered.

Birgé-Massart Strategy

The Birgé-Massart strategy for determining thresholds depends on several different parameters. You specify the wavelet decomposition and a thresholding method. You can also specify a sparsity parameter, `alpha`, or a specific multiplicative threshold rescaling, `scale`. Based on your inputs, wthrmngr derives the necessary Birgé-Massart parameters. The parameters depend on the dimension of the signal, and the total number, N, of coefficients at the coarsest scale of wavelet decomposition.

If the thresholding method is 'scarcehi', 'scarceme', or 'scarcelo', the wthrmngr executes either wdcbm or wdcbm2. If the thresholding method is 'penalhi', 'penalme', or 'penallo', then wthrmngr executes either wbmppen or wpbmpen.

Thresholding Method	Description
'scarcehi'	<ul style="list-style-type: none"> If the signal is 1-D, then wdcbm is used with input argument $M = N$. If the signal is 2-D, then wdcbm2 is used with $M = 4*N$.
'scarceme'	<ul style="list-style-type: none"> If the signal is 1-D, then wdcbm is used with input argument $M = 3*N/2$. If the signal is 2-D, then wdcbm2 is used with input argument with $M = 16*N/3$.
'scarcelo'	<ul style="list-style-type: none"> If the signal is 1-D, then wdcbm is used with input argument $M = 2N$. If the signal is 2-D, then wdcbm2 is used with input argument $M = 32*N/3$.

Thresholding Method	Description
'penalhi'	<ul style="list-style-type: none"> If the input is a wavelet decomposition, then <code>wbmpen</code> is used with $\text{ALPHA} = 5 \cdot (3 \cdot \alpha + 1) / 8$. If the input is a wavelet packet decomposition, then <code>wpbmpen</code> is used $\text{ALPHA} = 6.25$.
'penalme'	<ul style="list-style-type: none"> If the input is a wavelet decomposition, then <code>wbmpen</code> is used with $\text{ALPHA} = (\alpha + 5) / 8$. If the input is a wavelet packet decomposition, then <code>wpbmpen</code> is used $\text{ALPHA} = 2$.
'penallo'	<ul style="list-style-type: none"> If the input is a wavelet decomposition, then <code>wbmpen</code> is used with $\text{ALPHA} = (\alpha + 3) / 4$. If the input is a wavelet packet decomposition, then <code>wpbmpen</code> is used $\text{ALPHA} = 1.5$.

References

- [1] Birgé, L., and P. Massart. "From Model Selection to Adaptive Estimation." *Festschrift for Lucien Le Cam: Research Papers in Probability and Statistics* (E. Torgersen, D. Pollard, and G. Yang, eds.). New York: Springer-Verlag, 1997, pp. 55-88.

See Also

Apps

Wavelet Signal Denoiser

Functions

`wbmpen` | `wdcbm` | `wdcbm2` | `wdenoise` | `wpbmpen`

Introduced before R2006a

wtmm

Wavelet transform modulus maxima

Syntax

```
hexp = wtmm(x)
[hexp,tauq] = wtmm(x)
[ ___ ] = wtmm(x,'MinRegressionScale',scale)
[hexp,tauq,structfunc] = wtmm( ___ )

[localhexp,wt,wavscales] = wtmm(x,'ScalingExponent','local')
wtmm( ___ , 'ScalingExponent', 'local')

[ ___ ] = wtmm( ___ ,Name,Value)
```

Description

`hexp = wtmm(x)` returns an estimate of the global Holder exponent, `hexp`, for the real-valued, 1-D input signal, `x`. The global and local Holder exponents are estimated for the linearly-spaced moments of the structure functions from -2 to $+2$ in 0.1 increments.

`[hexp,tauq] = wtmm(x)` also returns an estimate of the partition function scaling exponents, `tauq`.

`[___] = wtmm(x,'MinRegressionScale',scale)` uses only scales greater than or equal to `scale` to estimate the global Holder exponent. This syntax can include any of the output arguments used in previous syntaxes.

`[hexp,tauq,structfunc] = wtmm(___)` also returns the multiresolution structure functions, `structfunc`, for the global Holder exponent estimate. This syntax can include any of the input arguments used in previous syntaxes.

`[localhexp,wt,wavscales] = wtmm(x,'ScalingExponent','local')` returns the local Holder exponent estimates, the continuous wavelet transform `wt`, and the scales, `wavscales`, which are used to calculate the CWT used in the `wtmm` algorithm. The wavelet used in the CWT is the second derivative of a Gaussian.

`wtmm(____, 'ScalingExponent', 'local')` with no output arguments plots the wavelet maxima lines in the current figure. Estimates of the local Holder exponents are displayed in a table to the right of the plot.

`[____] = wtmm(____, Name, Value)` returns the Holder exponent and other specified outputs with additional options specified by one or more `Name, Value` pair arguments.

Examples

Global Holder Exponent for Brownian Motion

Estimate the global Holder exponent for Brownian motion. This monofractal signal has a Holder exponent of approximately 0.5.

```
rng(100);
x = cumsum(randn(2^15,1));
hexp = wtmm(x)

hexp = 0.5010
```

Linearity of Scaling Exponents for Monofractal Signal

Confirm that for a monofractal signal, the scaling exponents are a linear function of the moments. For multifractal signals, the exponents are a nonlinear function of the moments.

Load a signal that contains two time series, each with 8000 samples. `Ts1` is a multifractal signal and `Ts2` is a monofractal fractional Brownian signal. Obtain the exponents using `wtmm`.

```
load RWdata;
[hexp1,tauq1] = wtmm(Ts1);
[hexp2,tauq2] = wtmm(Ts2);
```

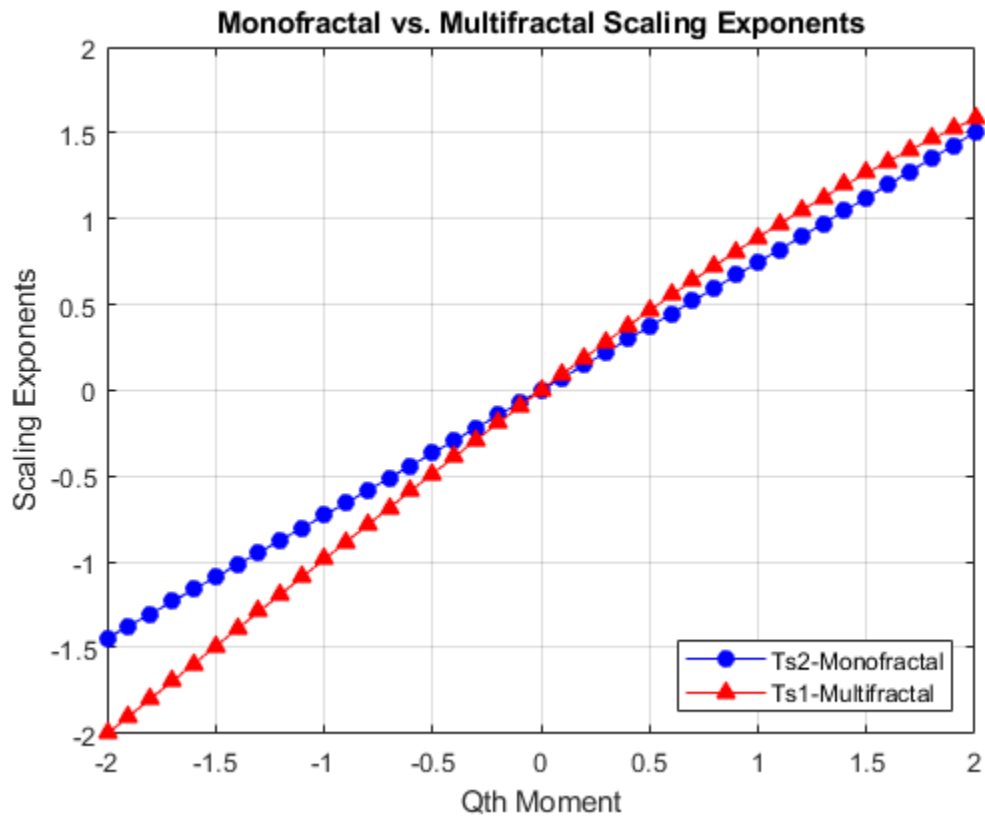
Plot the scaling exponents.

```
expplot = plot(-2:0.1:2,tauq2,'b-o',-2:0.1:2,tauq1,'r-^');
grid on;
expplot(1).MarkerFaceColor = 'b';
```

```

explot(2).MarkerFaceColor = 'r';
legend('Ts2-Monofractal', 'Ts1-Multifractal', 'Location', 'SouthEast');
title('Monofractal vs. Multifractal Scaling Exponents');
xlabel('Qth Moment');
ylabel('Scaling Exponents');

```



Ts2, which is the monofractal signal, is a linear function. Ts1, the multifractal signal, is not linear.

Structure Function of Wavelet Transform Modulus Maxima

Use the structure function output of `wmmm` to analyze a Brownian motion signal.

Create fractional Brownian motion with a Holder exponent of 0.6.

```
Brn = wfbm(0.6,2^15);  
[hexp,tauq,structfunc] = wtmm(Brn);
```

Compare the calculated Holder exponent with the theoretical value of 0.6.

```
hexp
```

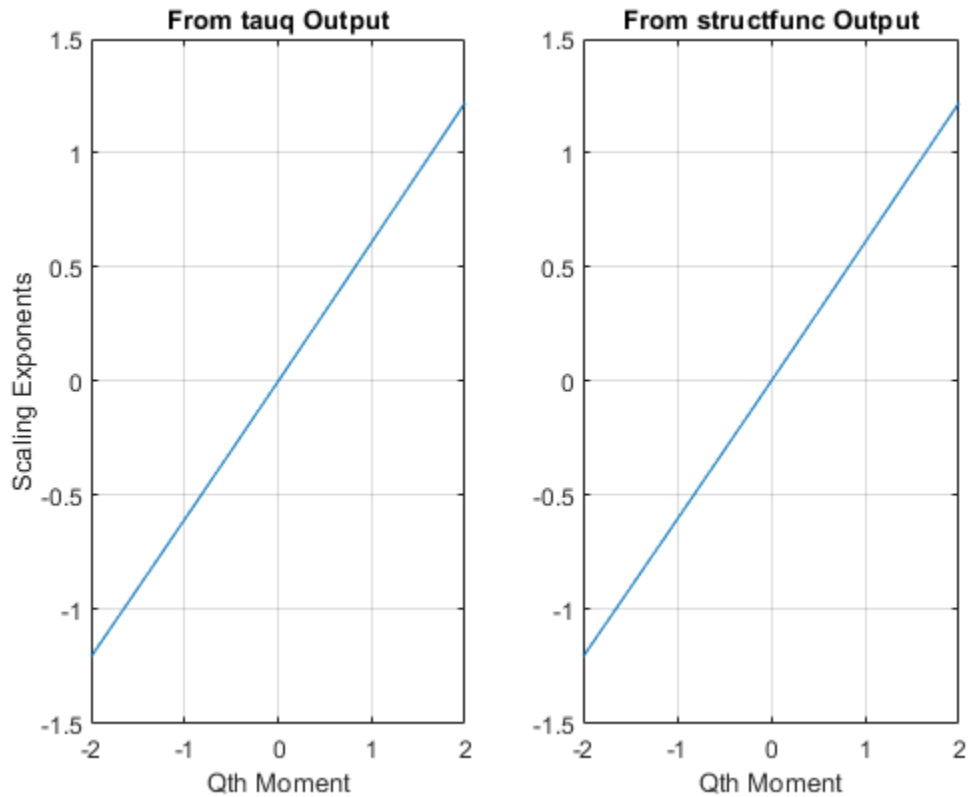
```
hexp = 0.6072
```

Use the data in the `structfunc` output and the `lscov` function to perform the regression on the data.

```
x = ones(length(structfunc.logscals),2);  
x(:,2) = structfunc.logscals;  
betahat = lscov(x,structfunc.Tq,structfunc.weights);  
betahat = betahat(2,:);
```

Plot and compare the scaling exponents from the `tauq` output and from the regressed structure function output.

```
subplot(1,2,1)  
plot(-2:.1:2,tauq)  
grid on  
title('From tauq Output')  
xlabel('Qth Moment')  
ylabel('Scaling Exponents')  
  
subplot(1,2,2)  
plot(-2:.1:2,betahat(1:41))  
grid on  
title('From structfunc Output')  
xlabel('Qth Moment')
```



The plots are the same and show a linear relationship between the moments and the exponents. Therefore, the signal is monofractal. The Holder exponent returned in `hexp` is the slope of this line.

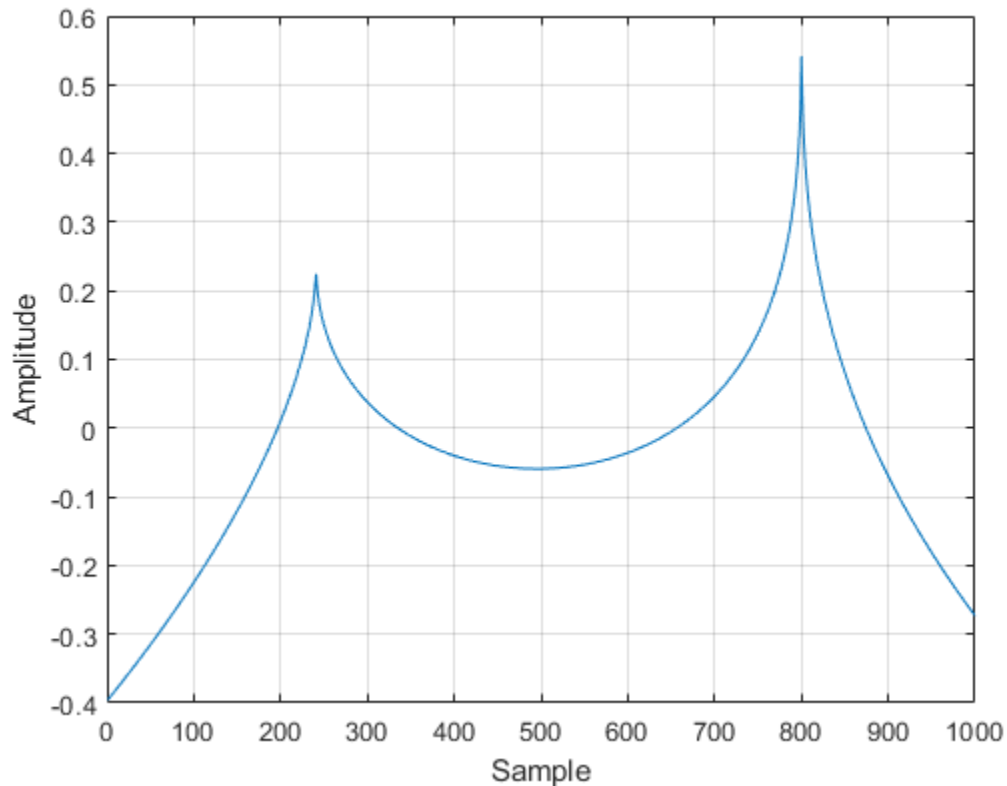
Local Holder Exponents for Cusp Signal and Delta Functions

Using a cusp signal and a signal containing delta functions, generate their local Holder exponents.

Cusp Signal

Load and plot a cusp signal. Note the difference between the two cusps.

```
load cusp;  
plot(cusp)  
grid on  
xlabel('Sample')  
ylabel('Amplitude')
```

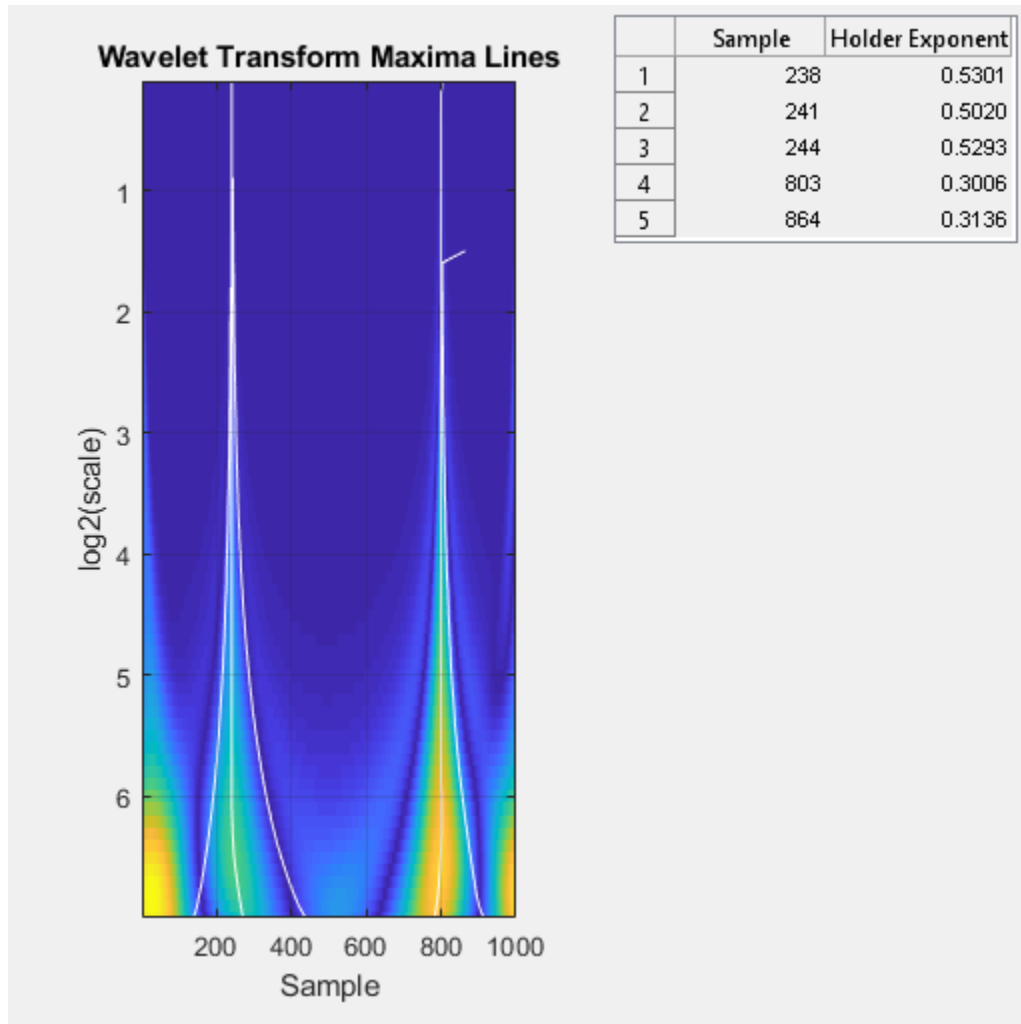


The equation for this cusp signal specifies a Holder exponent of 0.5 at sample 241 and a Holder exponent of 0.3 at sample 803.

$$-0.2*\text{abs}(x-241)^{0.5} - 0.5*\text{abs}(x-803)^{0.3} + 0.00346*x + 1.34$$

Obtain the local Holder exponents and plot the modulus maxima.

```
wtmm(cusp, 'ScalingExponent', 'local');
```

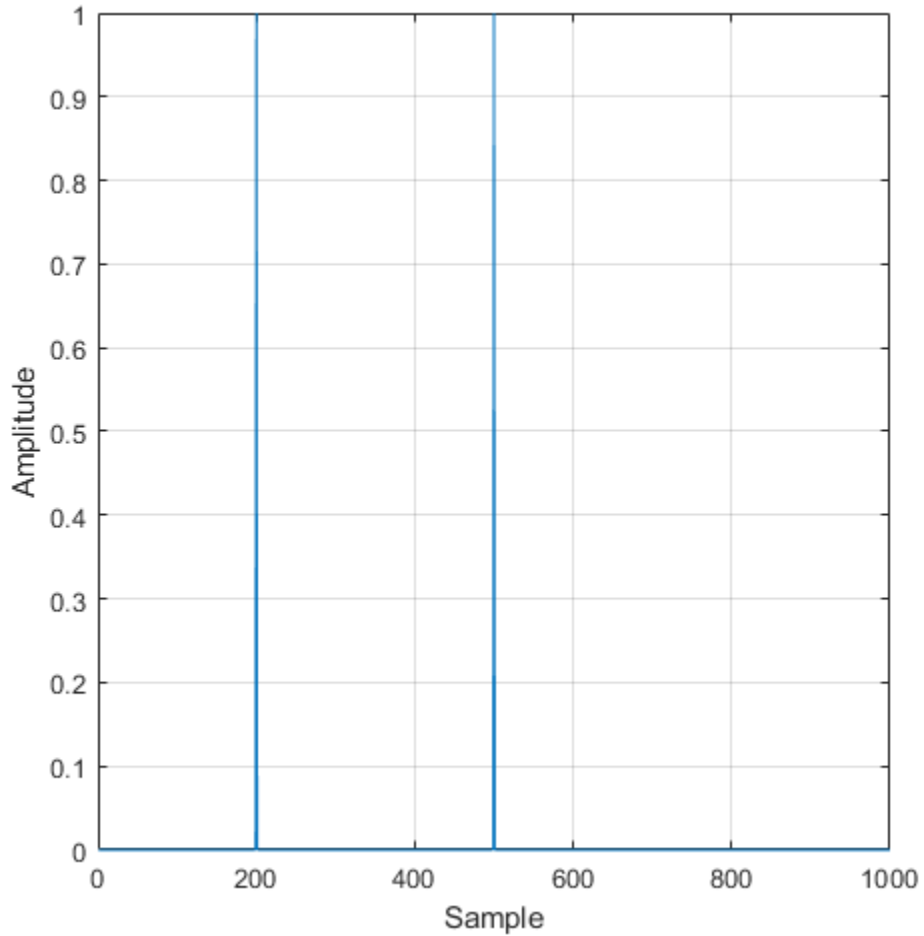


The Holder exponents at samples 241 and 803 are very close to the values specified in the cusp signal equation. The higher Holder value at sample 241 indicates that the signal at that point is closer to being differentiable than the signal at sample 803, which has a smaller Holder value.

Delta Functions

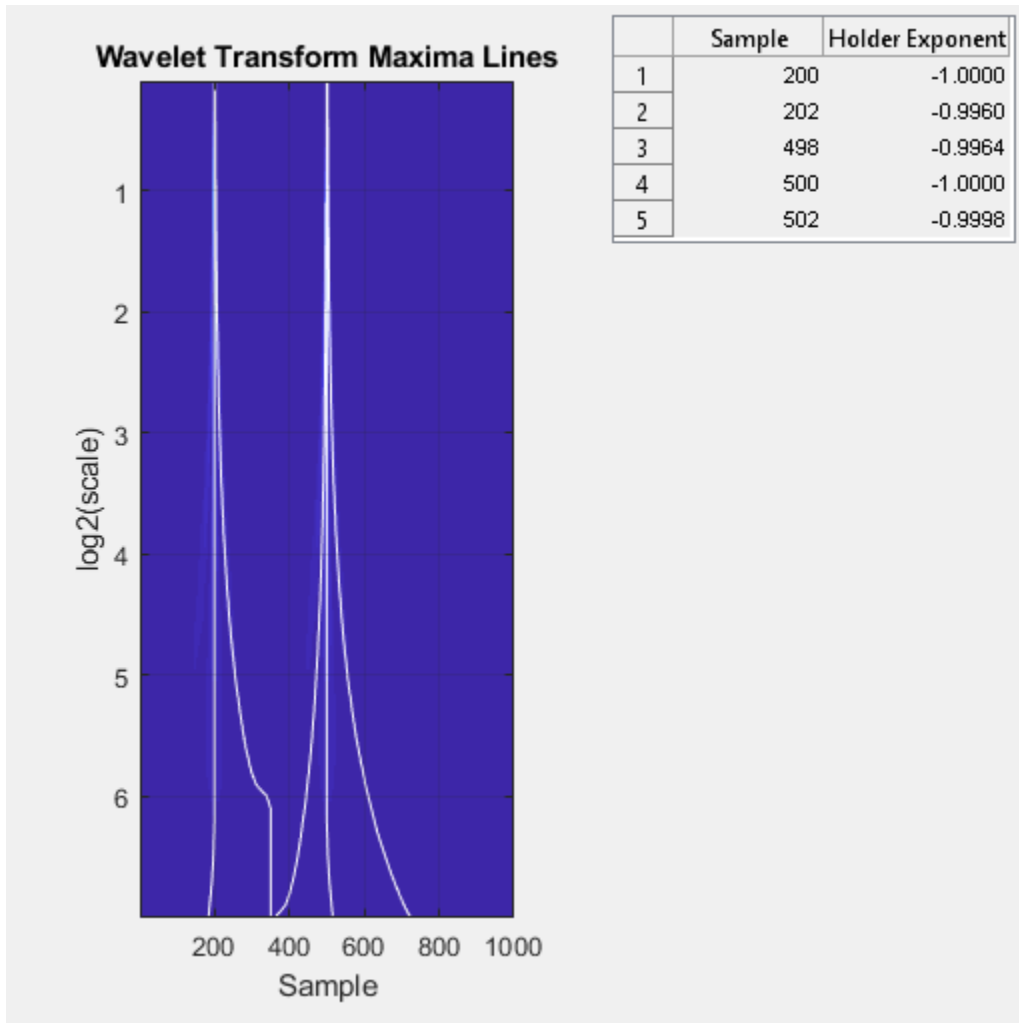
Create and plot two delta functions.

```
x = zeros(1e3,1);  
x([200 500]) = 1;  
plot(x)  
grid on  
xlabel('Sample')  
ylabel('Amplitude')
```



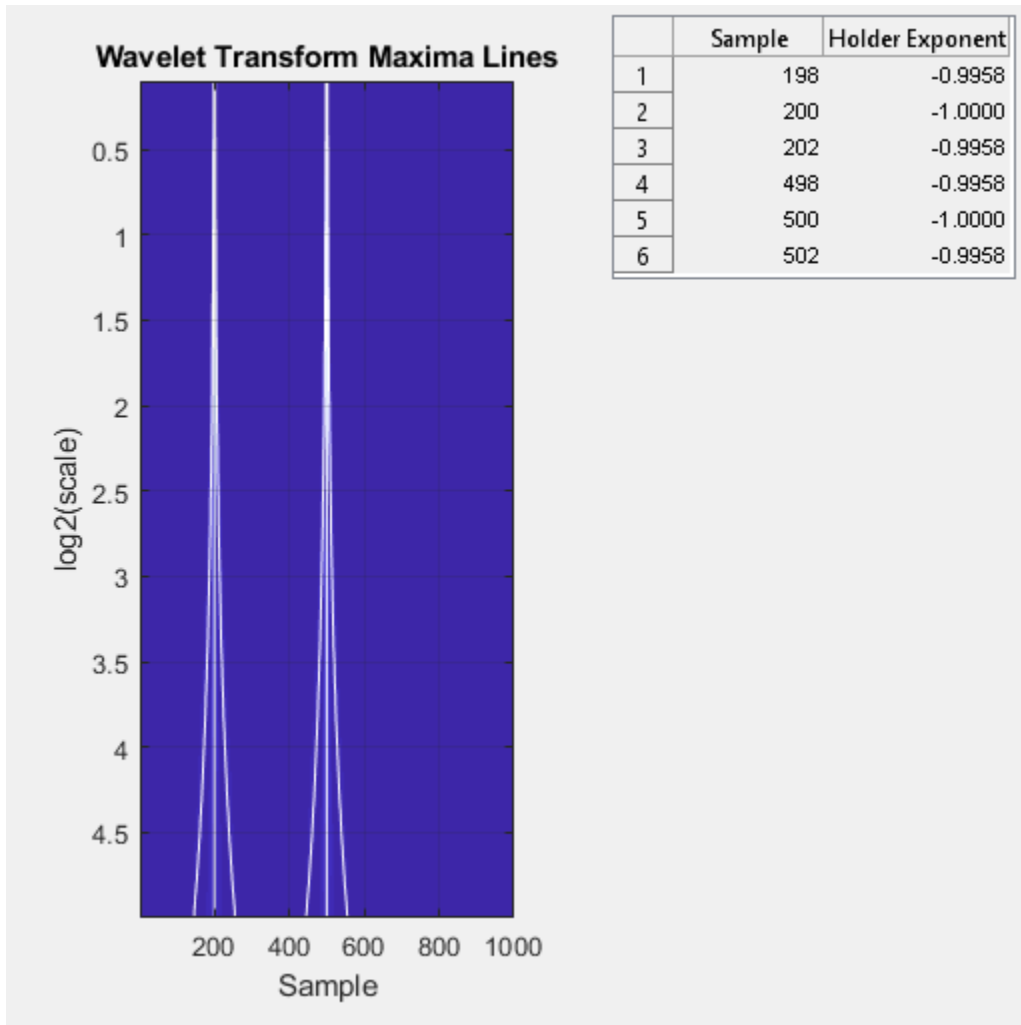
Obtain the local Holder exponents using the default number of octaves, which in this case is 7. Plot the modulus maxima. A delta function has a Holder exponent of -1.

```
wtmm(x, 'ScalingExponent', 'local');
```



Obtain the local Holder exponents using 5 octaves and compare the modulus maxima plot to the plot using the default number of octaves.

```
wtmm(x, 'ScalingExponent', 'local', 'NumOctaves', 5);
```



Reducing the number of scales provides more separation in frequency and less overlap between the modulus maxima lines of the delta functions.

Input Arguments

x — Input signal

real-valued vector

Input signal, specified as a real-valued vector with a minimum of 128 samples. The wavelet transform modulus maxima technique works best for data with 8000 or more samples.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, . . . , NameN, ValueN`.

Example: `'VoicesPerOctave', 18` estimates the global Holder estimate using 18 voices per octave.

MinRegressionScale — Minimum scale for regression

4 (default) | scalar greater than or equal to 4

Minimum scale for regression, specified as the comma-separated pair consisting of `'MinRegressionScale'` and a scalar greater than or equal to 4. This scale is the smallest scale used by the regression. There must be at least two scales with more than 6 CWT maxima. `'MinRegressionScale'` applies only to global Holder exponents.

VoicesPerOctave — Number of voices per octave

10 (default) | even integer from 8 to 32

Number of voices per octave, specified as the comma-separated pair consisting of `'VoicesPerOctave'` and an even integer from 8 to 32. The number of voices per octave and the number of octaves determine the number of scales used in the CWT.

NumOctaves — Number of octaves

minimum of 7 and $\text{floor}(\log_2(\text{numel}(x)/(3*\text{sqrt}(1.1666))))$ (default) | integer greater than or equal to 4

Number of octaves, specified as the comma-separated pair consisting of `'NumOctaves'` and an integer. The number of octaves and the number of voices per octave determine the number of scales used in the CWT. The maximum number of octaves is less than or equal to $\text{floor}(\log_2(\text{numel}(x)/(3*\text{sqrt}(1.1666))))$. The $\text{sqrt}(1.1666)$ factor is the standard deviation of the second derivative of a Gaussian wavelet. If you specify the number of octaves as greater than the maximum number of octaves, `wtmm` uses the maximum supported number of octaves.

ScalingExponent — Type of scaling exponents

`'global'` (default) | `'local'`

Type of scaling exponents, specified as a comma-separated pair consisting of `'ScalingExponent'` and either `'global'` or `'local'`. A global Holder exponent is used for monofractal signals, such as white noise, which are singular everywhere. Global holder exponents give a single estimate of degree of these singularities over the whole signal. Local Holder exponents are useful for signals with cusp singularities.

Output Arguments

hexp — Global Holder exponent

real scalar

Global Holder exponent, returned as a real scalar. Holder exponents are useful for identifying singularities, which are locations where a signal is not differentiable. A global Holder exponent uses a single value to estimate the degree of differentiability of all of the singularities of a signal. Signals with a global Holder exponent are monofractal signals.

tauq — Scaling exponents

column vector

Scaling exponents, returned as a column vector. The exponents are estimated for the linearly-spaced moments of the structure functions from -2 to +2 in 0.1 increments.

structfunc — Multiresolution structure functions

struct

Multiresolution structure functions for the global Holder exponent estimates, returned as a `struct`. The structure function for data `x` is defined as

$$S(q, a) = \frac{1}{n_a} \sum_{k=1}^{n_a} |T_x(a, k)|^q \approx a^{\zeta(q)},$$

where a is the scale, q is the moment, T_x is the maxima at each scale, n_a is the number of maxima at each scale, and $\zeta(q)$ is the scaling exponent. `structfunc` is a structure array containing the following fields:

- `Tq` — Measurements of the input, x , at various scales. `Tq` is a matrix of multiresolution quantities that depend jointly on time and scale. Scaling phenomena in x imply a power-law relationship between the moments of `Tq` and the scale. `Tq` is an N_s -by-44 matrix, where N_s is the number of scales. The first 41 columns of `Tq` contain the scaling exponent estimates for each of the q th from -2:0.1:2, by scale. The last three columns correspond to the first-order, second-order, and third-order cumulants, respectively, by scale. For a monofractal signal, cumulants greater than the first cumulant are zero.
- `weights` — Weights used in the regression estimates. The weights correspond to the number of wavelet maxima at each scale. `weights` is an N_s -by-1 vector.
- `logscals` — Scales used as predictors in the regression. `logscals` is an N_s -by-1 vector with the base-2 logarithm of the scales.

localhexp — Local Holder exponent estimates

array of real values

Local Holder exponent estimates, returned as an M -by-2 array of real values, where M is the number of maxima. If no maxima lines converge to the finest scale in the wavelet transform, then `localhexp` is an empty array. The wavelet transform modulus maxima method (WTMM) identifies cusp-like singularities in a signal. To analyze multifractal signals, use `dwtleader`.

wt — Continuous wavelet transform

matrix

Continuous wavelet transform, returned as a matrix of real values. `wt` is a `numel(wavscals)`-by- N matrix where N is the length of the input signal x .

wavscals — Wavelet scales

column vector

Wavelet scales, returned as a column vector of real values. `wavscals` are the scales used to calculate the CWT.

Algorithms

The WTMM algorithm finds singularities in a signal by determining maxima. The algorithm first calculates the continuous wavelet transform using the second derivative of a Gaussian wavelet with 10 voices per octave. The wavelet that meets this criteria is the Mexican hat, or Ricker, wavelet. Then, the algorithm determines the modulus maxima for each scale. The WTMM is intended to be used with large data sets so that enough samples are available to determine maxima accurately.

The definition of the modulus maximum at point x_0 and scale s_0 is

$$|Wf(s_0, x)| < |Wf(s_0, x_0)|$$

where x is either in the right or left neighborhood of x_0 . When x is in the opposite neighborhood of x_0 , the definition is

$$|Wf(s_0, x)| \leq |Wf(s_0, x_0)|$$

. The algorithm for finding additional maxima repeats for values in that scale. Then, the algorithm continues up through finer scales, checking whether the maxima align between scales. If a maximum converges to the finest scale, it is a true maximum and indicates a singularity at that point.

When each singularity is determined, the algorithm then estimates its Holder exponent. Holder exponents indicate the degree of differentiability for each singularity, which classifies the singularity strength. A Holder exponent less than or equal to 0 indicates a discontinuity at that location. Holder exponents greater than or equal to 1 indicate that the signal is differentiable at that location. Holder values between 0 and 1 indicate continuous, but not differentiable locations. They indicate how close the signal at that sample is to being differentiable. Holder exponents close to 0 indicate signal locations that are less differentiable than locations with exponents closer to 1. The signal is smoother at locations with higher local Holder exponents.

For signals with a few cusp-like singularities and Holder exponents that have large variation, you set the algorithm to return local Holder exponents, which provide individual values for each singularity. For signals with numerous Holder exponents that have relatively small variations, you set the algorithm to return a global Holder exponent. A global Holder exponent applies to the whole signal. For signals with many singularities, you can reduce the number of maxima found by limiting the algorithm to start at or regress to a specific minimum or maximum scale, respectively. For detailed information about the WTMM, see [1] and [3].

References

- [1] Mallat, S., and W. L. Hwang. "Singularity Detection and Processing with Wavelets." *IEEE Transactions on Information Theory*. Vol. 38, No. 2, March 1992, pp. 617-643.
- [2] Wendt, H. and P. Abry. "Multifractality Tests Using Bootstrapped Wavelet Leaders." *IEEE Transactions on. Signal Processing*. Vol. 55, No. 10, 2007, pp. 4811-4820.
- [3] Arneodo, A., B. Audit, N. Decoster, J.-F. Muzy, and C. Vaillant. "Wavelet-Based Multifractal Formalism: Application to DNA Sequences, Satellite Images of the Cloud Structure and Stock Market Data." *The Science of Disasters: Climate Disruptions, Heart Attacks, and Market Crashes*. Bunde, A., J. Kropp, and H. J. Schellnhuber, Eds. 2002, pp. 26-102.

See Also

dwtleader | wfbm

Introduced in R2016b

wtreemgr

NTREE manager

Syntax

Description

wtreemgr is a tree management utility.

This function returns information on the tree T depending on the value of the `OPT` parameter.

Allowed values for `OPT` are listed in the table below.

'allnodes'	Tree nodes
'isnode'	True for existing node
'istnode'	True for terminal nodes
'nodeasc'	Node ascendants
'nodedesc'	Node descendants
'nodepar'	Node parent
'ntnode'	Number of terminal nodes
'tnodes'	Terminal nodes
'leaves'	Terminal nodes
'noleaves'	Not terminal nodes
'order'	Tree order
'depth'	Tree depth

See Also

allnodes | istnode | leaves | nodeasc | nodedesc | nodepar | noleaves | ntnode |
tnodes | treedpth | treeord

Introduced before R2006a

wvarchg

Find variance change points

Syntax

```
[PTS_OPT, KOPT, T_EST] = wvarchg(Y, K, D)
```

Description

`[PTS_OPT, KOPT, T_EST] = wvarchg(Y, K, D)` computes estimated variance change points for the signal `Y` for `j` change points, with $j = 0, 1, 2, \dots, K$.

Integer `D` is the minimum delay between two change points.

Integer `KOPT` is the proposed number of change points ($0 \leq \text{KOPT} \leq K$). The vector `PTS_OPT` contains the corresponding change points.

For $1 \leq k \leq K$, `T_EST(k+1, 1:k)` contains the k instants of the variance change points and then, if $\text{KOPT} > 0$, `PTS_OPT = T_EST(KOPT+1, 1:KOPT)` else `PTS_OPT = []`.

`K` and `D` must be integers such that $1 < K \ll \text{length}(Y)$ and $1 \leq D \ll \text{length}(Y)$.

The signal `Y` should be zero mean.

`wvarchg(Y, K)` is equivalent to `wvarchg(Y, K, 10)`.

`wvarchg(Y)` is equivalent to `wvarchg(Y, 6, 10)`.

Examples

Detect Variance Change Points

Add two variance change points to the blocks signal. Detect the variance change points using `wvarchg`.

Load the blocks signal. Add white noise with two variance change points located at index 180 and 600.

```
x = wnoise(1,10);
rng default;
bb = 1.5*randn(1,length(x));
cp1 = 180; cp2 = 600;
x = x + [bb(1:cp1),bb(cp1+1:cp2)/4,bb(cp2+1:end)];
```

Obtain the level-1 wavelet coefficients. Replace the top 2% of values with the mean value of the wavelet coefficients to remove all signal.

```
wname = 'db3'; lev = 1;
[c,l] = wavedec(x,lev,wname);
det = wrcoef('d',c,l,wname,1);
y = sort(abs(det));
v2p100 = y(fix(length(y)*0.98));
ind = find(abs(det)>v2p100);
det(ind) = mean(det);
```

Estimate the variance change points using the wavelet coefficients.

```
[pts_Opt,kopt,t_est] = wvarchg(det,5);
sprintf('The estimated change points are %d and %d\n',pts_Opt)
```

```
ans =
    'The estimated change points are 181 and 601
    '
```

References

Lavielle, M. (1999), "Detection of multiple changes in a sequence of dependent variables," *Stoch. Proc. and their Applications*, 83, 2, pp. 79-102.

Introduced before R2006a

wvd

Wigner-Ville distribution and smoothed pseudo Wigner-Ville distribution

Syntax

```
d = wvd(x)
```

```
d = wvd(x, fs)
```

```
d = wvd(x, ts)
```

```
d = wvd( ____, 'smoothedPseudo' )
```

```
d = wvd( ____, 'smoothedPseudo', twin, fwin)
```

```
d = wvd( ____, 'smoothedPseudo', Name, Value)
```

```
d = wvd( ____, 'MinThreshold', thresh)
```

```
[d, f, t] = wvd( ____ )
```

```
wvd( ____ )
```

Description

`d = wvd(x)` returns the Wigner-Ville distribution of `x`.

`d = wvd(x, fs)` returns the Wigner-Ville distribution when `x` is sampled at a rate `fs`.

`d = wvd(x, ts)` returns the Wigner-Ville distribution when `x` is sampled with a time interval `ts` between samples.

`d = wvd(____, 'smoothedPseudo')` returns the smoothed pseudo Wigner-Ville distribution of `x`. The function uses the length of the input signal to choose the lengths of the windows used for time and frequency smoothing. This syntax can include any combination of input arguments from previous syntaxes.

`d = wvd(____, 'smoothedPseudo', twin, fwin)` specifies the time window, `twin`, and the frequency window, `fwin`, used for smoothing. To use the default window for either time or frequency smoothing, specify the corresponding argument as empty, `[]`.

`d = wvd(____, 'smoothedPseudo', Name, Value)` specifies additional options for the smoothed pseudo Wigner-Ville distribution using name-value pair arguments. You can specify `twin` and `fwin` in this syntax, or you can omit them.

`d = wvd(____, 'MinThreshold', thresh)` sets to zero those elements of `d` whose amplitude is less than `thresh`. This syntax applies to both the Wigner-Ville distribution and the smoothed pseudo Wigner-Ville distribution.

`[d, f, t] = wvd(____)` also returns a vector of frequencies, `f`, and a vector of times, `t`, at which `d` is computed.

`wvd(____)` with no output arguments plots the Wigner-Ville or smoothed pseudo Wigner-Ville distribution in the current figure.

Examples

Wigner-Ville Distribution of Impulse and Tone

Generate a 1000-sample impulse and a 1000-sample tone with normalized frequency $\pi/2$. Compute the Wigner-Ville distribution of the sum of the two signals.

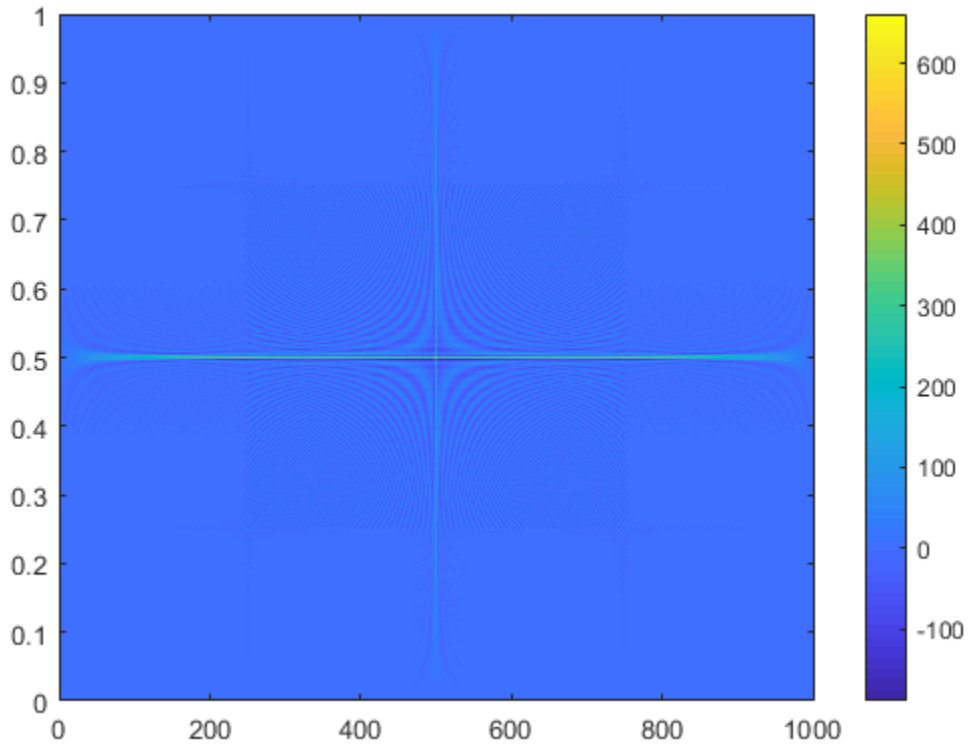
```
x = zeros(1001,1);  
x(500) = 10;
```

```
y = sin(pi*(0:1000)/2)';
```

```
[d, f, t] = wvd(x+y);
```

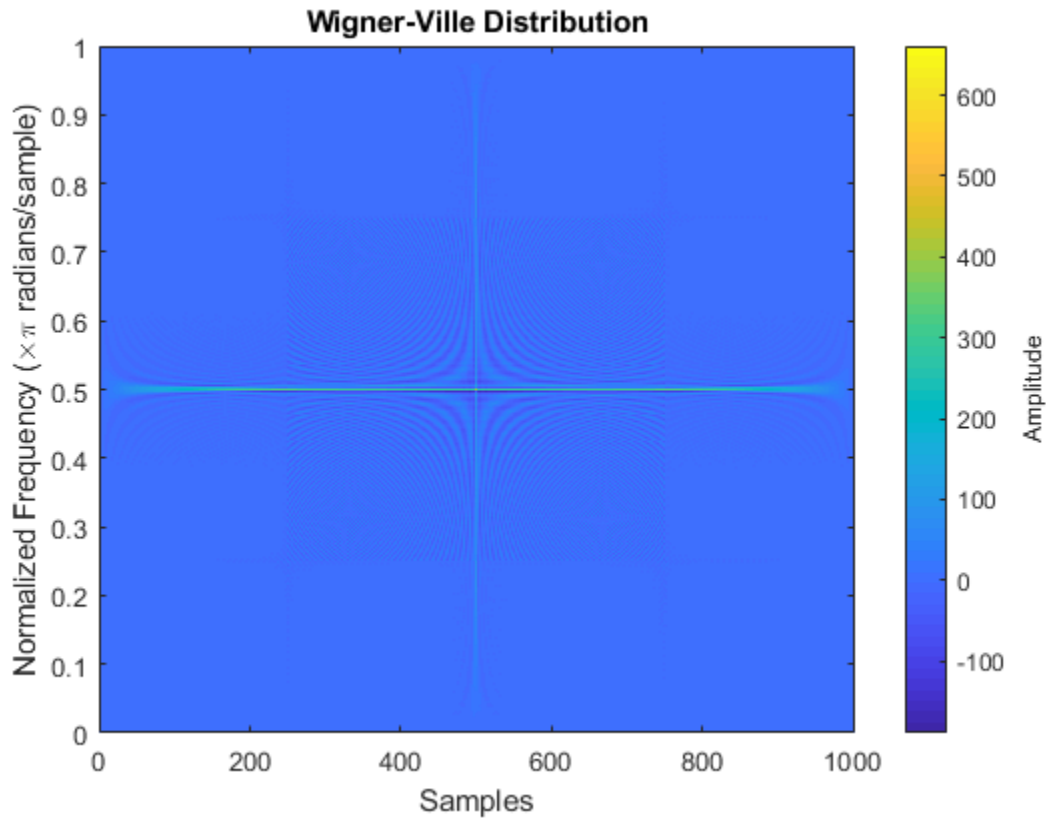
Plot the Wigner-Ville distribution.

```
imagesc(t, f, d)  
axis xy  
colorbar
```



Reproduce the result by calling `wvd` with no output arguments.

`wvd(x+y)`



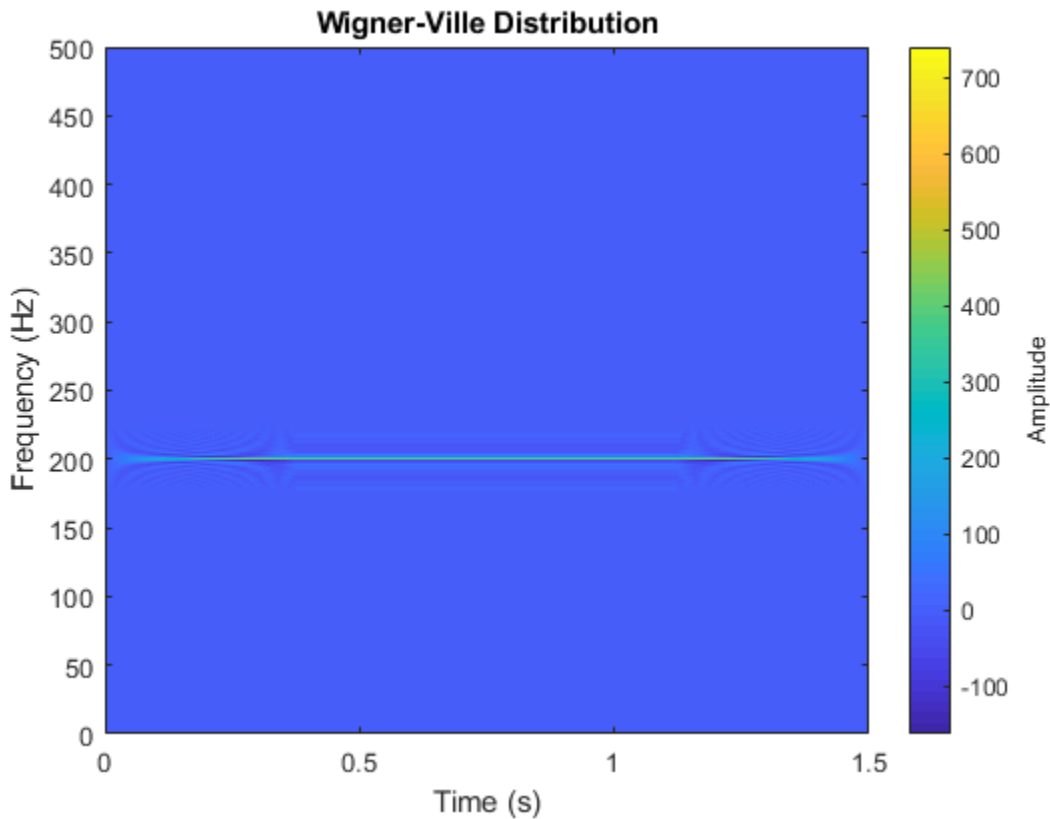
Wigner-Ville Distribution of Sinusoids

Generate a signal consisting of a 200 Hz sinusoid sampled at 1 kHz for 1.5 seconds.

```
fs = 1000;  
t = (0:1/fs:1.5)';  
x = cos(2*pi*t*200);
```

Compute the Wigner-Ville distribution of the signal.

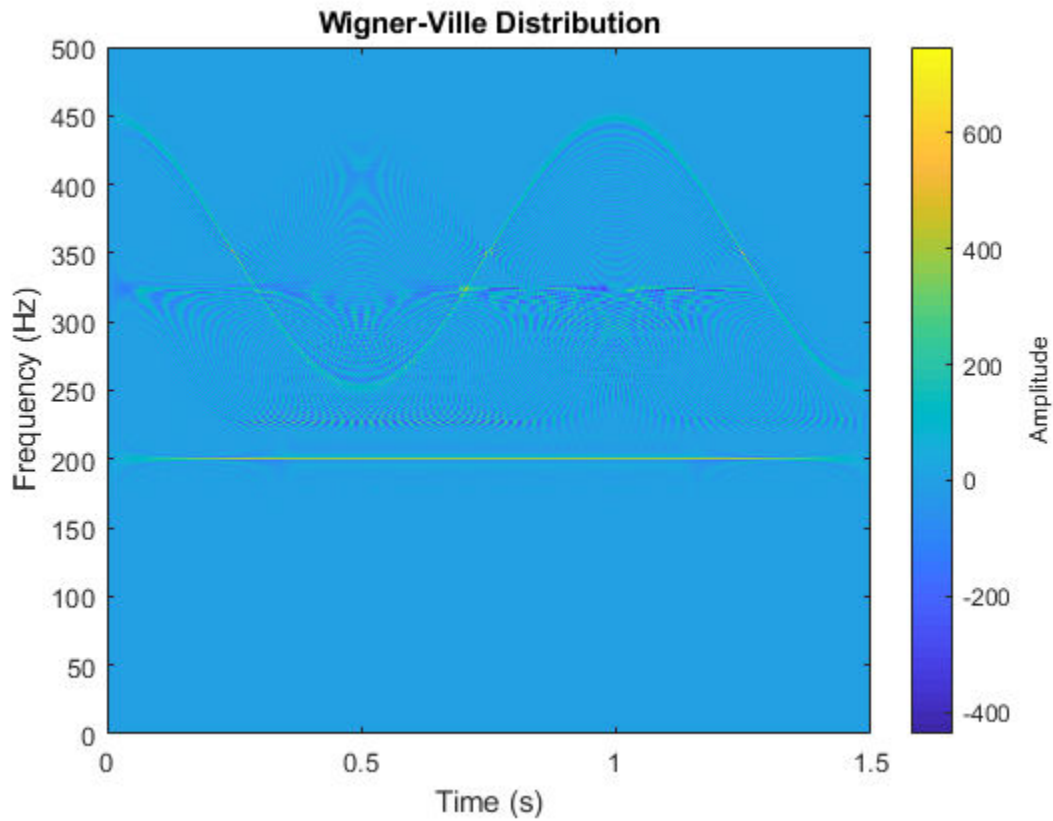
```
wvd(x, fs)
```



Add to the signal a chirp whose frequency varies sinusoidally between 250 Hz and 450 Hz. Convert the signal to a MATLAB® timetable. Compute the Wigner-Ville distribution.

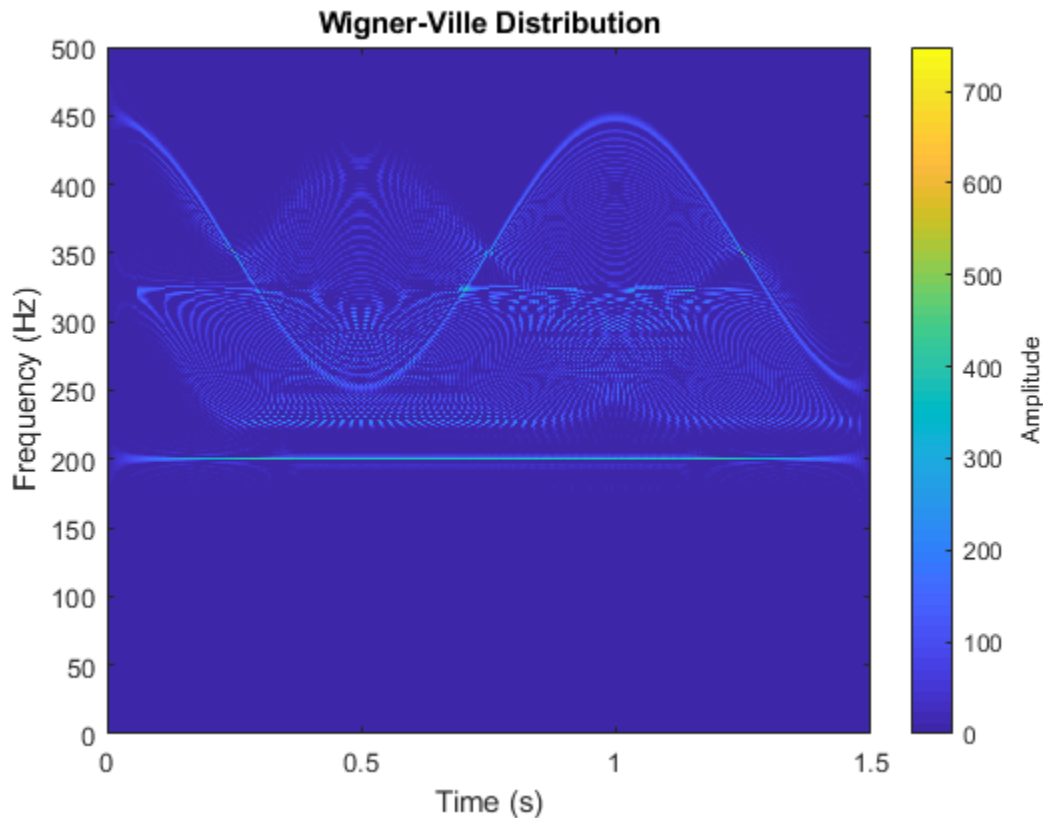
```
x = x + vco(cos(2*pi*t), [250 450], fs);  
xt = timetable(seconds(t), x);
```

```
wvd(xt)
```



Set to zero the distribution elements with amplitude less than 0.

```
wvd(xt, 'MinThreshold', 0)
```



Wigner-Ville Distribution of Chirps

Generate a signal sampled at 1 kHz for 1 second. One component of the signal is a chirp that increases in frequency quadratically from 100 Hz to 400 Hz during the measurement. The other component of the signal is a chirp that decreases in frequency linearly from 350 Hz to 50 Hz in the same lapse.

Store the signal in a timetable.

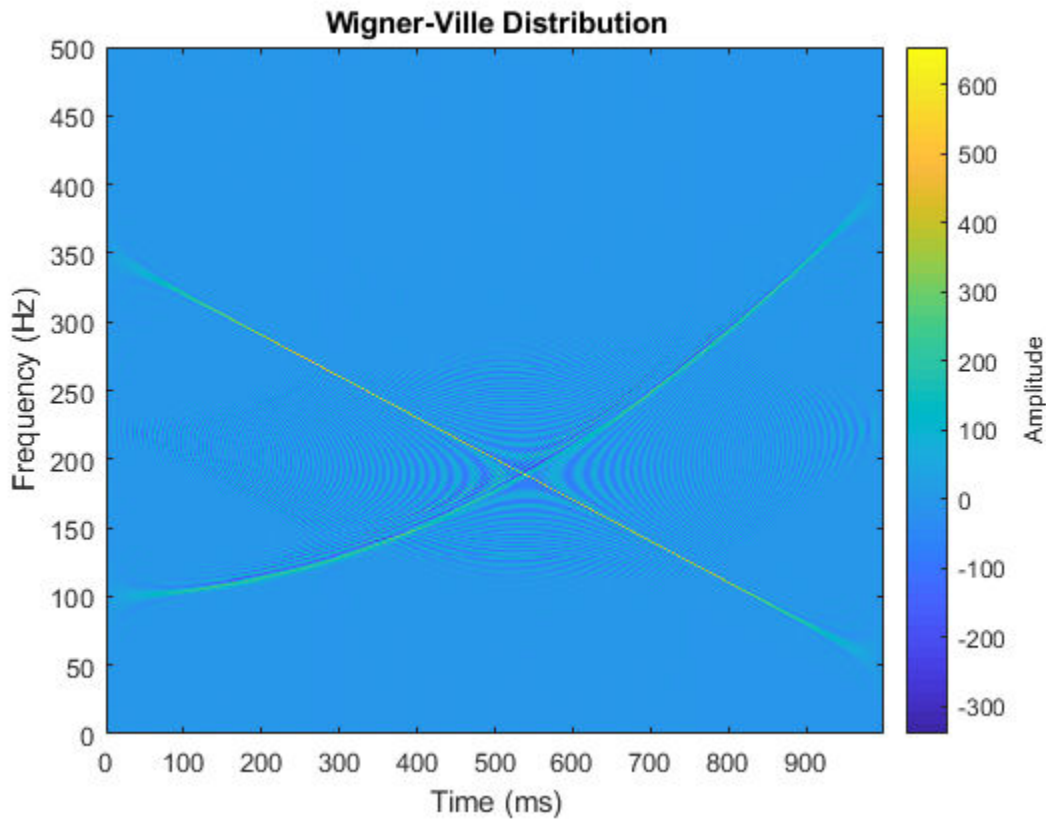
```
fs = 1000;  
t = 0:1/fs:1-1/fs;
```



```
x = chirp(t,100,1,400,'quadratic') + chirp(t,350,1,50);
```

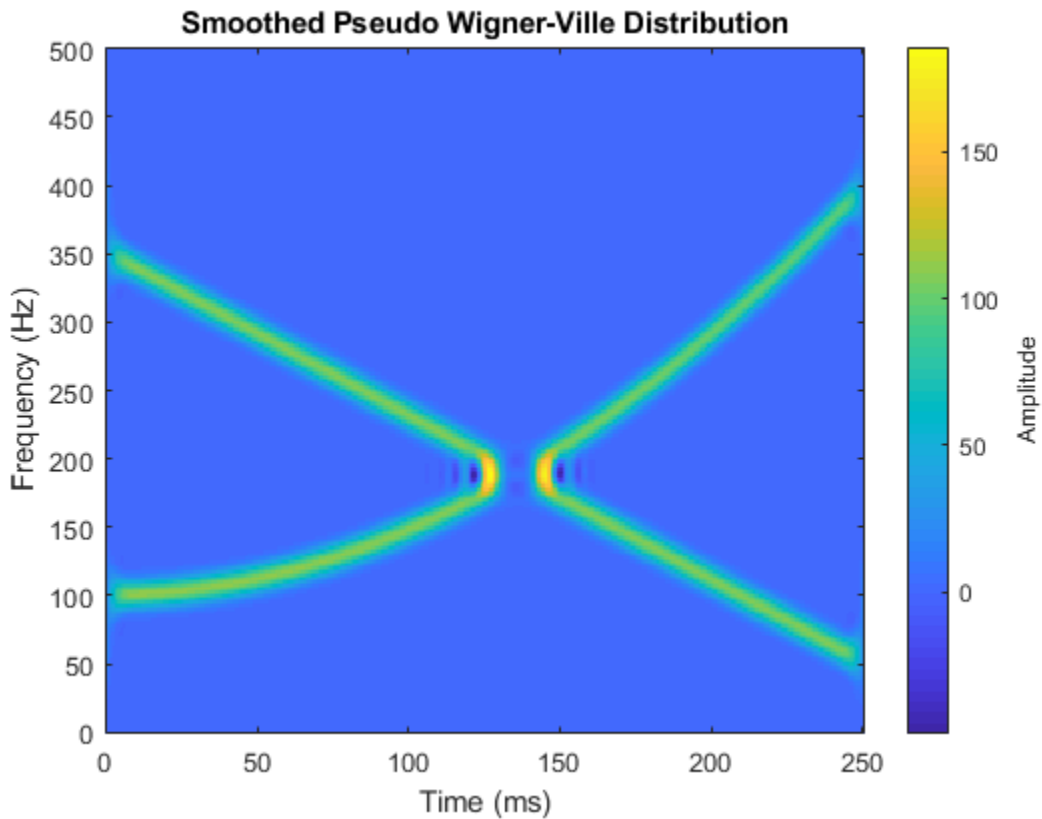
Compute the Wigner-Ville distribution of the signal.

```
wvd(x, fs)
```



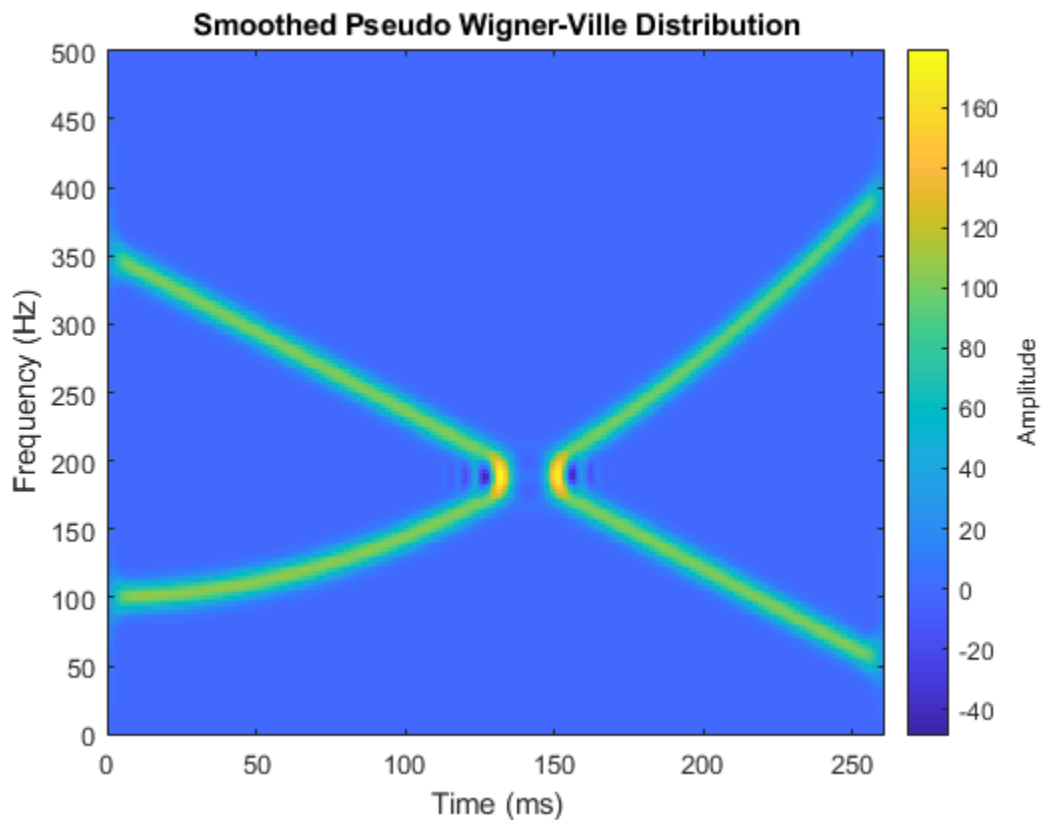
Compute the smoothed pseudo Wigner-Ville distribution of the signal. Specify 201 frequency points and 502 time points.

```
wvd(x, fs, 'smoothedPseudo', 'NumFrequencyPoints', 201, 'NumTimePoints', 502)
```



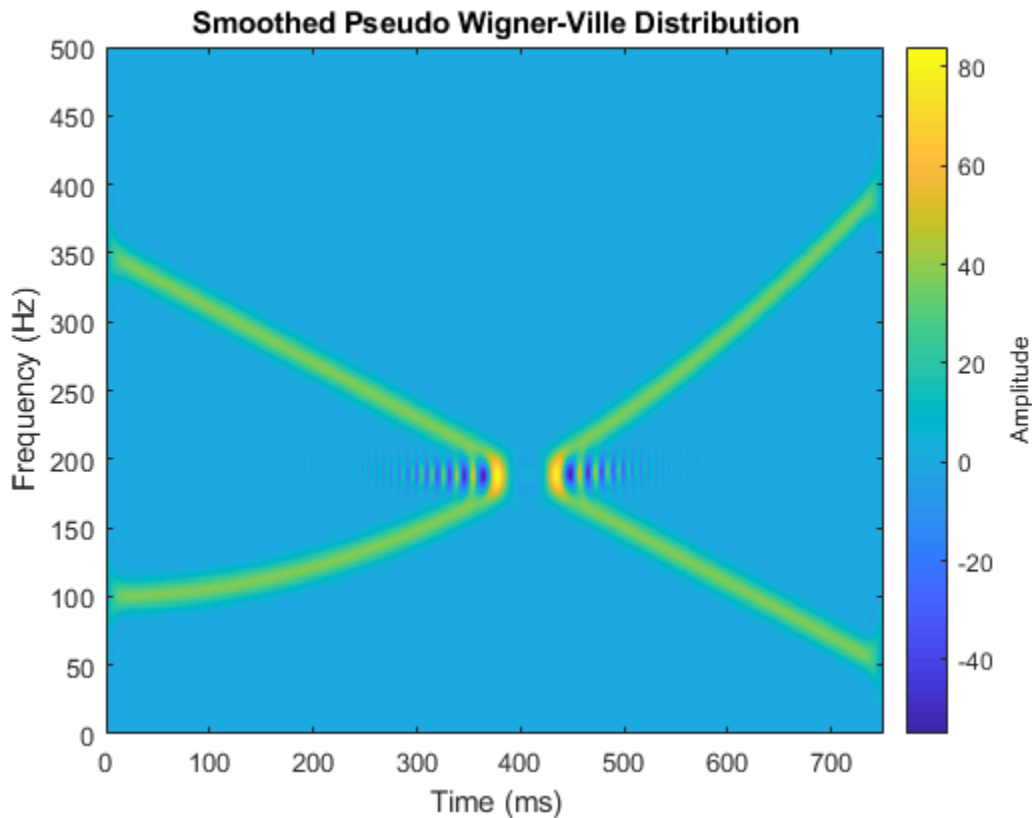
Increase the number of time points so the quadratic chirp becomes visible.

```
wvd(x, fs, 'smoothedPseudo', 'NumFrequencyPoints', 201, 'NumTimePoints', 522)
```



Increase the frequency points and time points to get a sharper image.

```
wvd(x, fs, 'smoothedPseudo', 'NumFrequencyPoints', 1000, 'NumTimePoints', 1502)
```



Smoothed Pseudo Wigner-Ville Distribution of Complex Signal

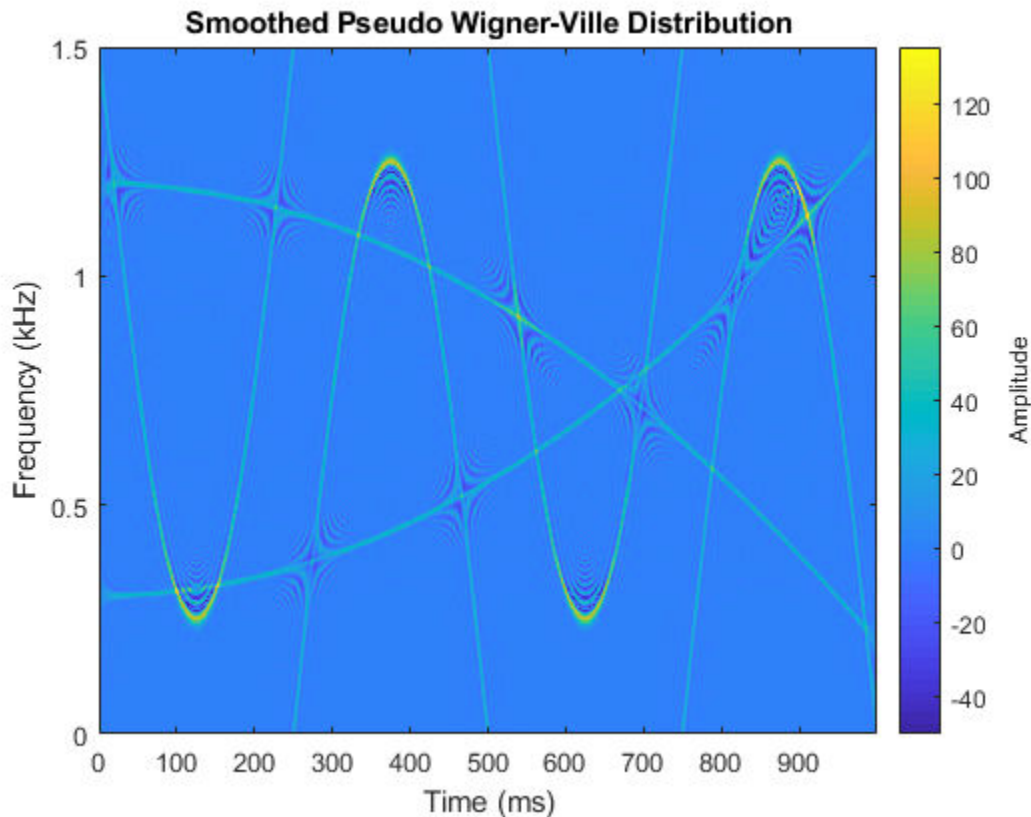
Generate a two-component signal sampled at 3 kHz for 1 second. The first component is a quadratic chirp whose frequency increases from 300 Hz to 1300 Hz during the measurement. The second component is a chirp with sinusoidally varying frequency content. The signal is embedded in white Gaussian noise. Express the time between consecutive samples as a duration scalar.

```
fs = 3000;  
t = 0:1/fs:1-1/fs;  
dt = seconds(t(2)-t(1));
```

```
x1 = chirp(t,300,t(end),1300,'quadratic');  
x2 = exp(2j*pi*100*cos(2*pi*2*t));  
  
x = x1 + x2 + randn(size(t))/10;
```

Compute and plot the smoothed pseudo Wigner Ville of the signal. Window the distribution in time using a 601-sample Hamming window and in frequency using a 305-sample rectangular window. Use 600 frequency points for the display. Set to zero those components of the distribution with amplitude less than -50 .

```
wvd(x,dt,'smoothedPseudo',hamming(601),rectwin(305), ...  
    'NumFrequencyPoints',600,'MinThreshold',-50)
```



Interference Terms

Generate a signal composed of four Gaussian atoms. Each atom consists of a sinusoid modulated by a Gaussian. The sinusoids have frequencies of 100 Hz and 400 Hz. The Gaussians are centered at 150 milliseconds and 350 milliseconds and have a variance of 0.01^2 . All atoms have unit amplitude. The signal is sampled at 1 kHz for half a second.

```
fs = 1000;  
t = (0:1/fs:0.5)';
```

```
f1 = 100;  
f2 = 400;
```

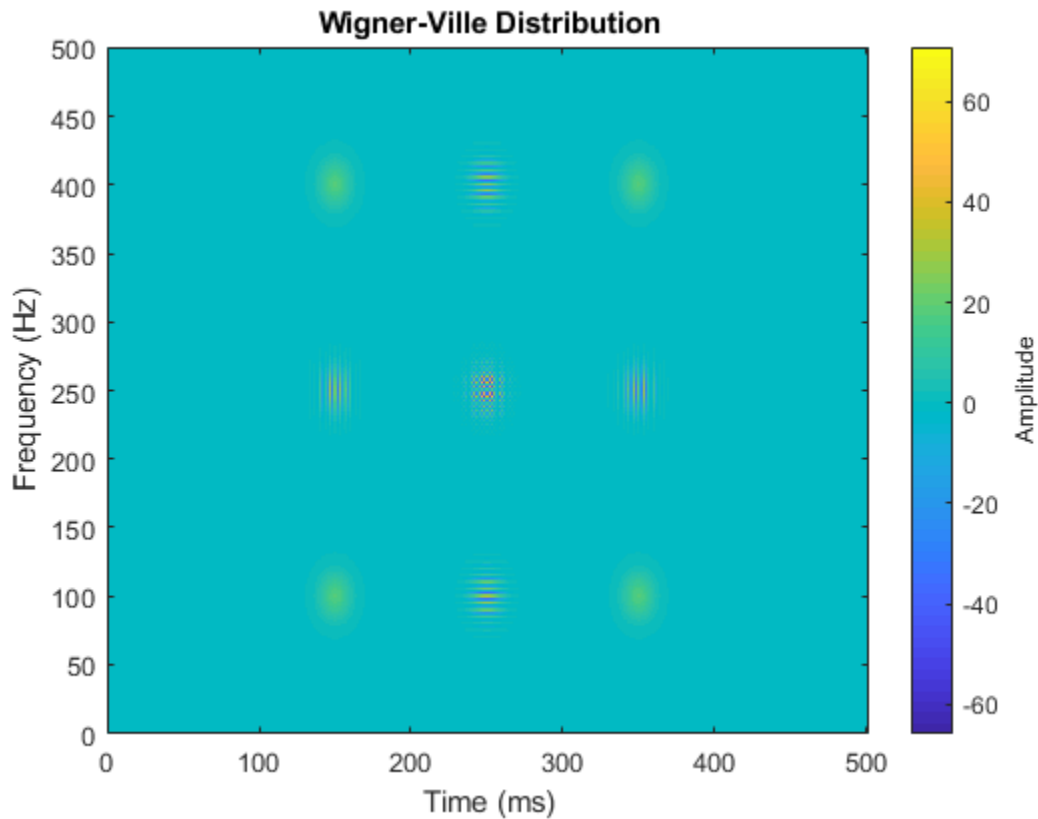
```
mu1 = 0.15;  
mu2 = 0.35;
```

```
gaussFun = @(A,x,mu,f) exp(-(x-mu).^2/(2*0.01^2)).*sin(2*pi*f.*x)*A';
```

```
s = gaussFun([1 1 1 1],t,[mu1 mu1 mu2 mu2],[f1 f2 f1 f2]);
```

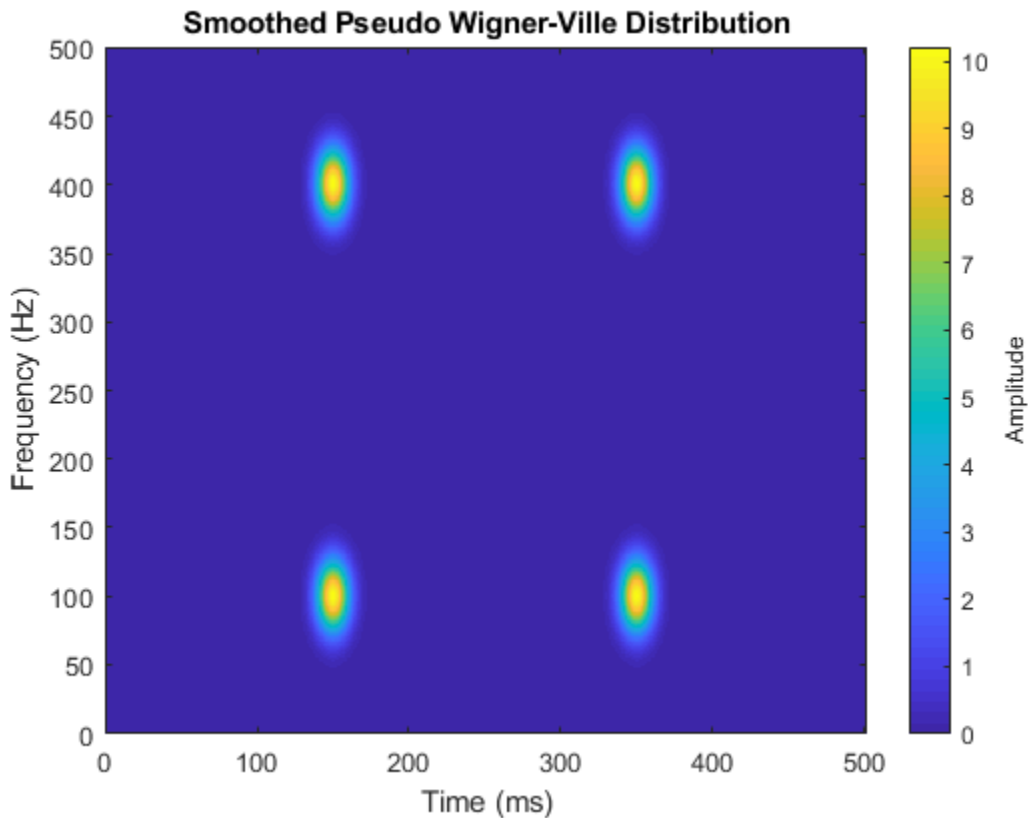
Compute and display the Wigner-Ville distribution of the signal. Interference terms, which can have negative values, appear halfway between each pair of auto-terms.

```
wvd(s, fs)
```



Compute and display the smoothed pseudo Wigner-Ville distribution of the signal. Smoothing in time and frequency attenuates the interference terms.

```
wvd(s, fs, 'SmoothedPseudo')
```



Input Arguments

x — Input signal

vector | timetable

Input signal, specified as a vector or a MATLAB timetable containing a single vector variable.

- If **x** is a timetable, then it must contain increasing finite row times.
- If a timetable has missing or duplicate time points, you can fix it using the tips in “Clean Timetable with Missing, Duplicate, or Nonuniform Times” (MATLAB).

If the input signal has odd length, the function appends a zero to make the length even.

Example: `cos(pi/8*(0:159))'+randn(160,1)/10` specifies a sinusoid embedded in white noise.

Example: `timetable(seconds(0:5)',rand(6,1))` specifies a random variable sampled at 1 Hz for 5 seconds.

Data Types: `single` | `double`

Complex Number Support: Yes

fs — Sample rate

`2*pi` (default) | positive numeric scalar

Sample rate, specified as a positive numeric scalar.

ts — Sample time

duration scalar

Sample time, specified as a duration scalar.

twin, fwin — Time and frequency windows

vectors of odd length

Time and frequency windows used for smoothing, specified as vectors of odd length. By default, `wvd` uses Kaiser windows with shape factor $\beta = 20$.

- The default length of `twin` is the smallest odd integer greater than or equal to `round(length(x)/10)`.
- The default length of `fwin` is the smallest odd integer greater than or equal to `nf/4`, where `nf` is specified using `NumFrequencyPoints`.

Each window must have a length smaller than or equal to `2*ceil(length(x)/2)`.

Example: `kaiser(65,0.5)` specifies a 65-sample Kaiser window with a shape factor of 0.5.

thresh — Minimum nonzero value

`-Inf` (default) | real scalar

Minimum nonzero value, specified as a real scalar. The function sets to zero those elements of `d` whose amplitudes are less than `thresh`.

Name-Value Pair Arguments

Specify optional comma-separated pairs of `Name`, `Value` arguments. `Name` is the argument name and `Value` is the corresponding value. `Name` must appear inside quotes. You can specify several name and value pair arguments in any order as `Name1, Value1, ..., NameN, ValueN`.

Example: `'NumFrequencyPoints', 201, 'NumTimePoints', 300` computes the Wigner-Ville distribution at 201 frequency points and 300 time points.

NumFrequencyPoints — Number of frequency points

`2*ceil(length(x)/2)` (default) | integer

Number of frequency points, specified as the comma-separated pair consisting of `'NumFrequencyPoints'` and an integer. This argument controls the degree of oversampling in frequency. The number of frequency points must be at least $(\text{length}(\text{fwin})+1)/2$ and cannot be greater than the default.

NumTimePoints — Number of time points

`4*ceil(length(x)/2)` (default) | even integer

Number of time points, specified as the comma-separated pair consisting of `'NumTimePoints'` and an even integer. This argument controls the degree of oversampling in time. The number of time points must be at least $2*\text{length}(\text{twin})$ and cannot be greater than the default. If the input signal is large, reduce the number of time points to lower the memory requirements and speed up the computation.

Output Arguments

d — Wigner-Ville distribution

matrix

Wigner-Ville distribution, returned as a matrix. Time increases across the columns of `d`, and frequency increases down the rows. The matrix is of size $N_f \times N_t$, where N_f is the length of `f` and N_t is the length of `t`.

f — Frequencies

vector

Frequencies, returned as a vector.

- If the input has time information, then f contains frequencies expressed in Hz.
- If the input does not have time information, then f contains normalized frequencies expressed in rad/sample.

t — Time instants

vector

Time instants, returned as a vector.

- If the input has time information, then t contains time values expressed in seconds.
- If the input does not have time information, then t contains sample numbers.

Definitions

Wigner-Ville Distribution

The *Wigner-Ville distribution* provides a high-resolution time-frequency representation of a signal. The distribution has applications in signal visualization, detection, and estimation.

For a continuous signal $x(t)$, the Wigner-Ville distribution is defined as

$$\text{WVD}_x(t, f) = \int_{-\infty}^{\infty} x\left(t + \frac{\tau}{2}\right) x^*\left(t - \frac{\tau}{2}\right) e^{-j2\pi f \tau} d\tau.$$

For a discrete signal with N samples, the distribution becomes

$$\text{WVD}_x(n, k) = \sum_{m=-N}^N x(n + m/2) x^*(n - m/2) e^{-j2\pi km/N}.$$

For odd values of m , the definition requires evaluation of the signal at half-integer sample values. It therefore requires interpolation, which makes it necessary to zero-pad the discrete Fourier transform to avoid aliasing.

The Wigner-Ville distribution contains interference terms that often complicate its interpretation. To sharpen the distribution, one can filter the definition with lowpass windows. The smoothed pseudo Wigner-Ville distribution uses independent windows to smooth in time and frequency:

$$\text{SPWVD}_x^{g,H}(t, f) = \int_{-\infty}^{\infty} g(t) H(f) x\left(t + \frac{\tau}{2}\right) x^*\left(t - \frac{\tau}{2}\right) e^{-j2\pi f\tau} d\tau.$$

References

- [1] Cohen, Leon. *Time-Frequency Analysis: Theory and Applications*. Englewood Cliffs, NJ: Prentice-Hall, 1995.
- [2] Mallat, Stéphane. *A Wavelet Tour of Signal Processing*. Second Edition. San Diego, CA: Academic Press, 1999.
- [3] O'Toole, John M., and Boualem Boashash. "Fast and memory-efficient algorithms for computing quadratic time-frequency distributions." *Applied and Computational Harmonic Analysis*. Vol. 35, No. 2, pp. 350-358.

See Also

Functions

xwvd

Introduced in R2018b

xwvd

Cross Wigner-Ville distribution and cross smoothed pseudo Wigner-Ville distribution

Syntax

```
d = xwvd(x,y)
d = xwvd(x,y,fs)
d = xwvd(x,y,ts)

d = xwvd( ____, 'smoothedPseudo' )
d = xwvd( ____, 'smoothedPseudo', twin, fwin)
d = xwvd( ____, 'smoothedPseudo', 'NumFrequencyPoints', nf)

d = xwvd( ____, 'MinThreshold', thresh)

[d,f,t] = xwvd( ____ )

xwvd( ____ )
```

Description

`d = xwvd(x,y)` returns the cross Wigner-Ville distribution of `x` and `y`.

`d = xwvd(x,y,fs)` returns the cross Wigner-Ville distribution when `x` and `y` are sampled at a rate `fs`.

`d = xwvd(x,y,ts)` returns the cross Wigner-Ville distribution when `x` and `y` are sampled with a time interval `ts` between samples.

`d = xwvd(____, 'smoothedPseudo')` returns the cross smoothed pseudo Wigner-Ville distribution of `x` and `y`. The function uses the length of the input signals to choose the lengths of the windows used for time and frequency smoothing. This syntax can include any combination of input arguments from previous syntaxes.

`d = xwvd(____, 'smoothedPseudo', twin, fwin)` specifies the time window, `twin`, and the frequency window, `fwin`, used for smoothing. To use the default window for either time or frequency smoothing, specify the corresponding argument as empty, `[]`.

`d = xwvd(____, 'smoothedPseudo', 'NumFrequencyPoints', nf)` computes the cross smoothed pseudo Wigner-Ville distribution using `nf` frequency points. You can specify `twin` and `fwin` in this syntax, or you can omit them.

`d = xwvd(____, 'MinThreshold', thresh)` sets to zero those elements of `d` whose amplitude is less than `thresh`. This syntax applies to both the cross Wigner-Ville distribution and the cross smoothed pseudo Wigner-Ville distribution.

`[d, f, t] = xwvd(____)` also returns a vector of frequencies, `f`, and a vector of times, `t`, at which `d` is computed.

`xwvd(____)` with no output arguments plots the real part of the cross Wigner-Ville or cross smoothed pseudo Wigner-Ville distribution in the current figure.

Examples

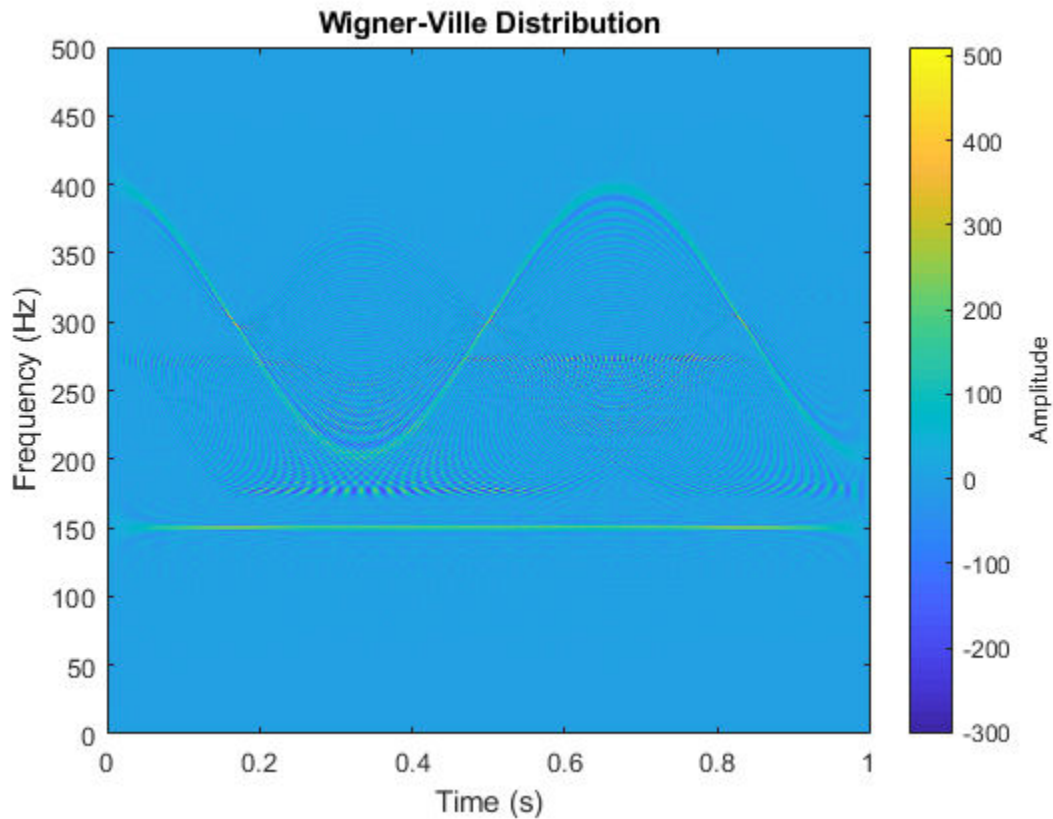
Cross Wigner-Ville Distribution of Signals

Generate two signals sampled at 1 kHz for 1 second and embedded in white noise. One signal is a sinusoid of frequency 150 Hz. The other signal is a chirp whose frequency varies sinusoidally between 200 Hz and 400 Hz. The noise has a variance of 0.1^2 .

```
fs = 1000;  
t = (0:1/fs:1)';  
  
x = cos(2*pi*t*150) + 0.1*randn(size(t));  
y = vco(cos(3*pi*t), [200 400], fs) + 0.1*randn(size(t));
```

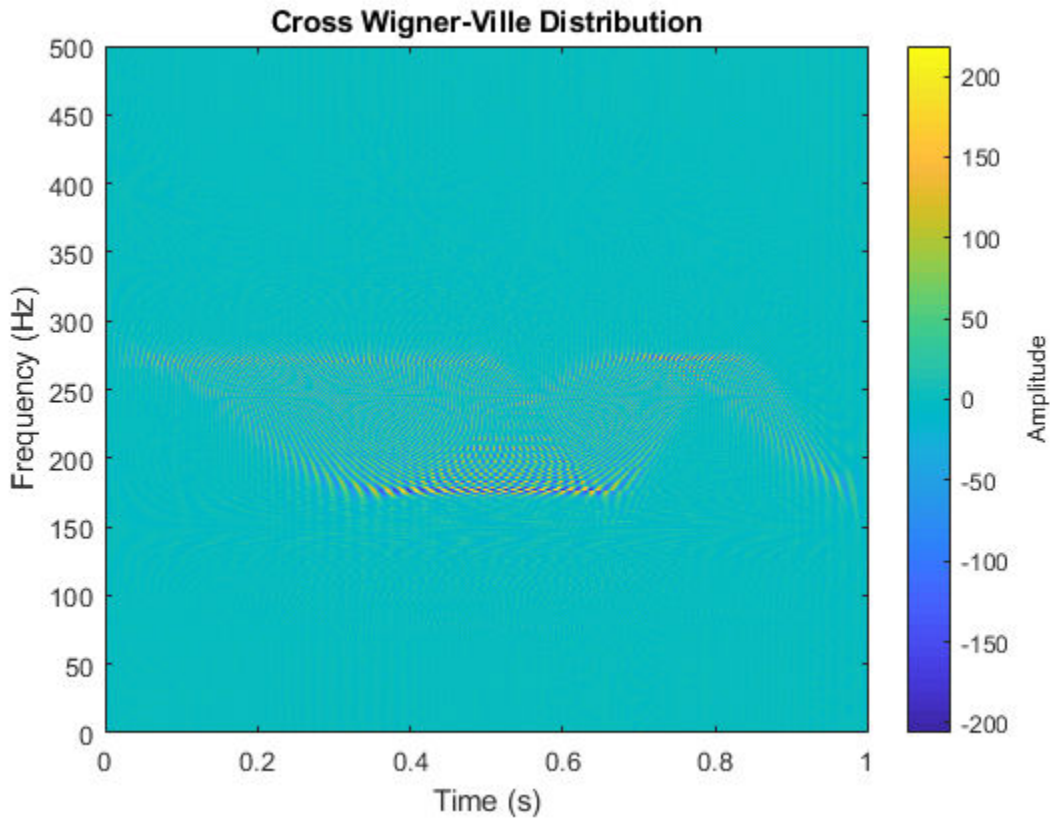
Compute the Wigner-Ville distribution of the sum of the signals.

```
wvd(x+y, fs)
```



Compute and plot the cross Wigner-Ville distribution of the signals. The cross-distribution corresponds to the cross-terms of the Wigner-Ville distribution.

```
xwvd(x, y, fs)
```



Cross Wigner-Ville Distribution of Chirps

Generate a two-channel signal that consists of two chirps. The signal is sampled at 3 kHz for one second. The first chirp has an initial frequency of 400 Hz and reaches 800 Hz at the end of the sampling. The second chirp starts at 500 Hz and reaches 1000 Hz at the end. The second chirp has twice the amplitude of the first chirp.

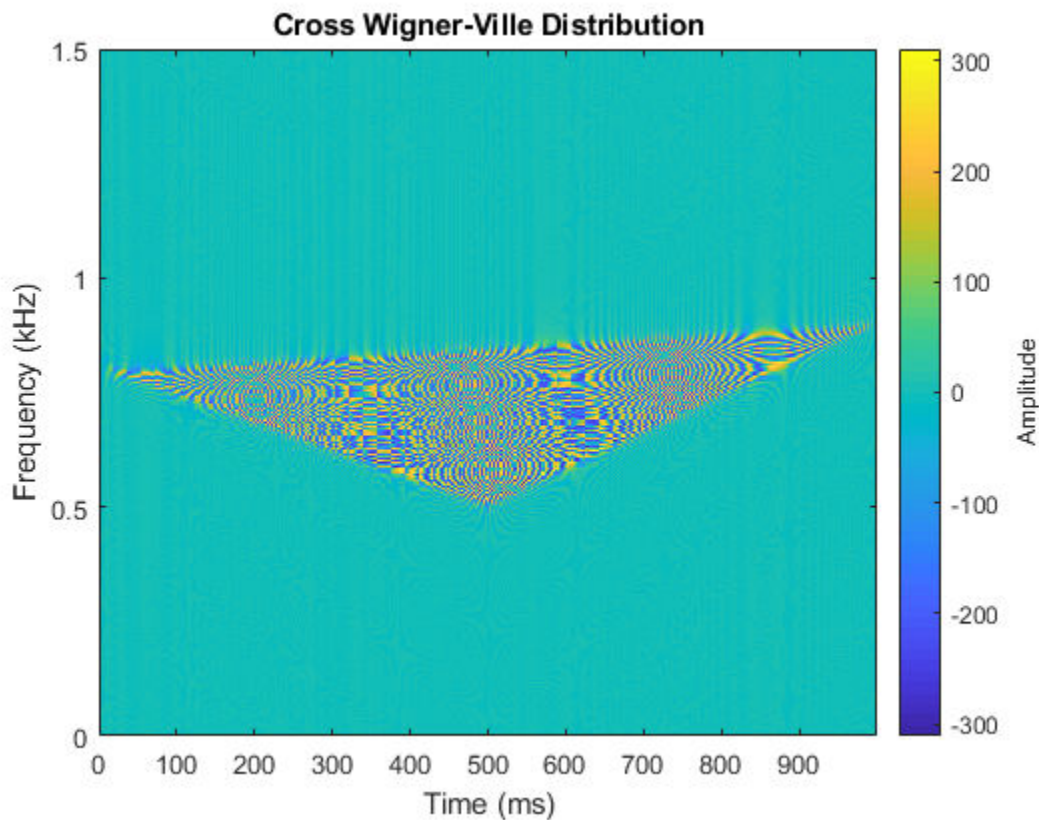
```
fs = 3000;  
t = (0:1/fs:1-1/fs)';
```



```
x1 = chirp(t,1400,t(end),800);  
x2 = 2*chirp(t,200,t(end),1000);
```

Store the signal as a timetable. Compute and plot the cross Wigner-Ville distribution of the two channels.

```
xt = timetable(seconds(t),x1,x2);  
xwvd(xt(:,1),xt(:,2))
```



Use Cross Wigner-Ville Distribution to Estimate Instantaneous Frequency

Compute the instantaneous frequency of a signal by using a known reference signal and the cross Wigner-Ville distribution.

Create a reference signal consisting of a Gaussian atom sampled at 1 kHz for 1 second. A Gaussian atom is a sinusoid modulated by a Gaussian. Specify a sinusoid frequency of 50 Hz. The Gaussian is centered at 64 milliseconds and has a variance of 0.01^2 .

```
fs = 1e3;
t = (0:1/fs:1-1/fs)';

mu = 0.064;
sigma = 0.01;
fsin = 50;

xr = exp(-(t-mu).^2/(2*sigma^2)).*sin(2*pi*fsin*t);
```

Create the "unknown" signal to analyze, consisting of a chirp. The signal starts suddenly at 0.4 second and ends suddenly half a second later. In that lapse, the frequency of the chirp decreases linearly from 400 Hz to 100 Hz.

```
f0 = 400;
f1 = 100;

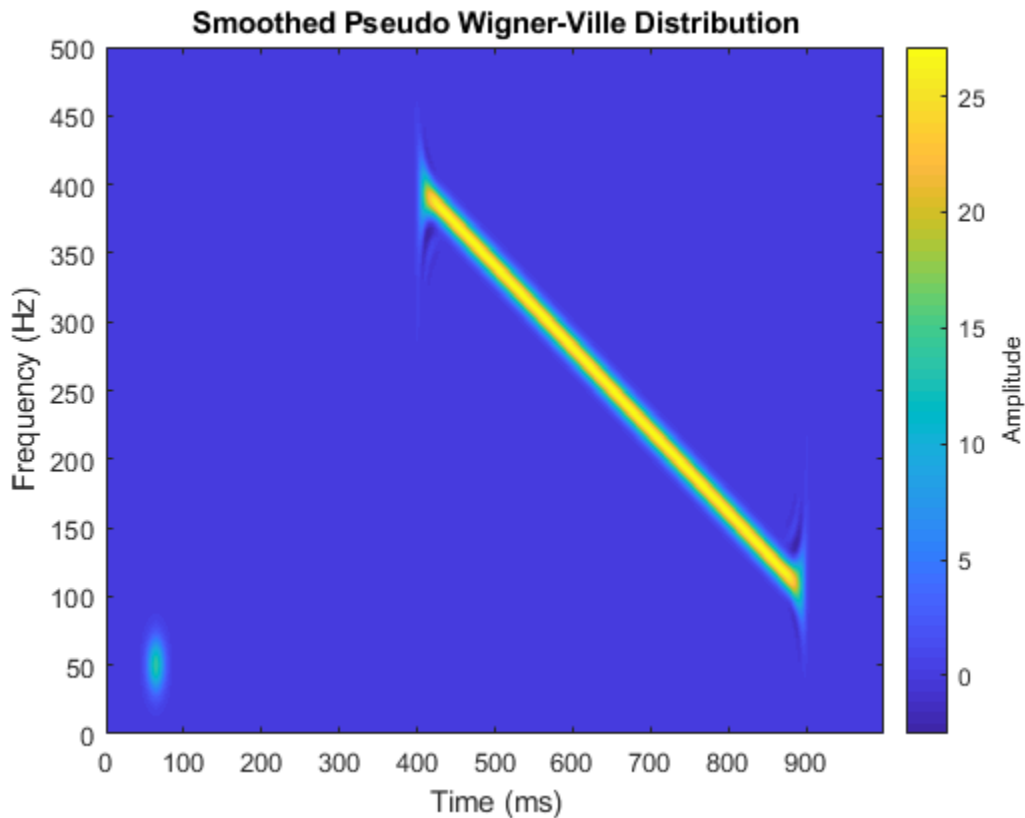
xa = zeros(size(t));
xa(t>0.4 & t<=0.9) = chirp((0:1/fs:0.5-1/fs)', f0, 0.5, f1);
```

Create a two-component signal consisting of the sum of the unknown and reference signals. The smoothed pseudo Wigner-Ville distribution of the result provides an "ideal" time-frequency representation.

Compute and display the smoothed pseudo Wigner-Ville distribution.

```
w = wvd(xa+xr, fs, 'smoothedPseudo');

wvd(xa+xr, fs, 'smoothedPseudo')
```

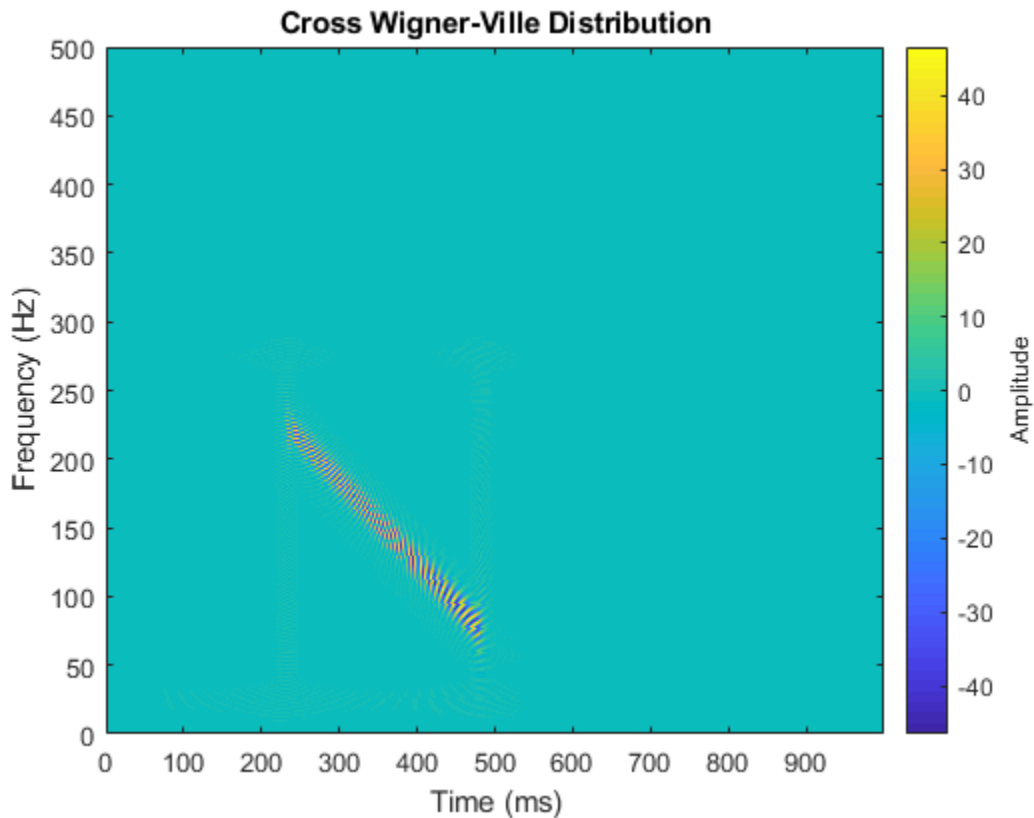


Compute the cross Wigner-Ville distribution of the unknown and reference signals. Take the absolute value of the distribution and set to zero the elements with amplitude less than 10. The cross Wigner-Ville distribution is equal to the cross-terms of the two-component signal.

Plot the real part of the cross Wigner-Ville distribution.

```
[c, fc, tc] = xwvd(xa, xr, fs);  
c = abs(c);  
c(c < 10) = 0;
```

```
xwvd(xa, xr, fs)
```



Enhance the Wigner-Ville cross-terms by adding the ideal time-frequency representation to the cross Wigner-Ville distribution. The cross-terms of the Wigner-Ville distribution occur halfway between the reference signal and the unknown signal.

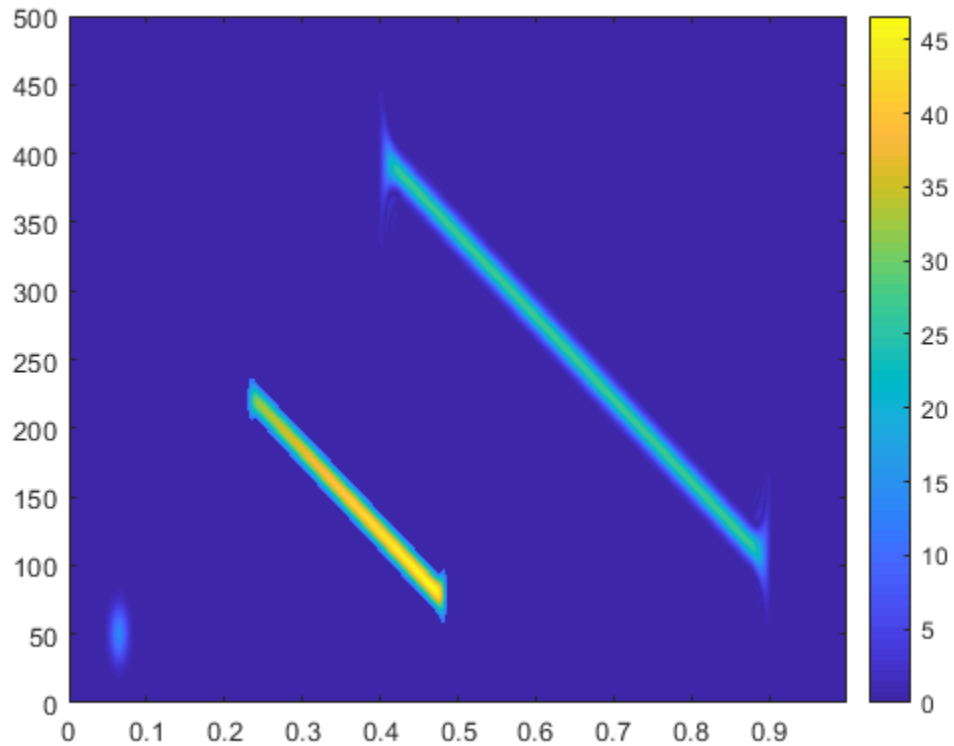
```
d = w + c;
```

```
d = abs(real(d));
```

```
imagesc(tc,fc,d)
```

```
axis xy
```

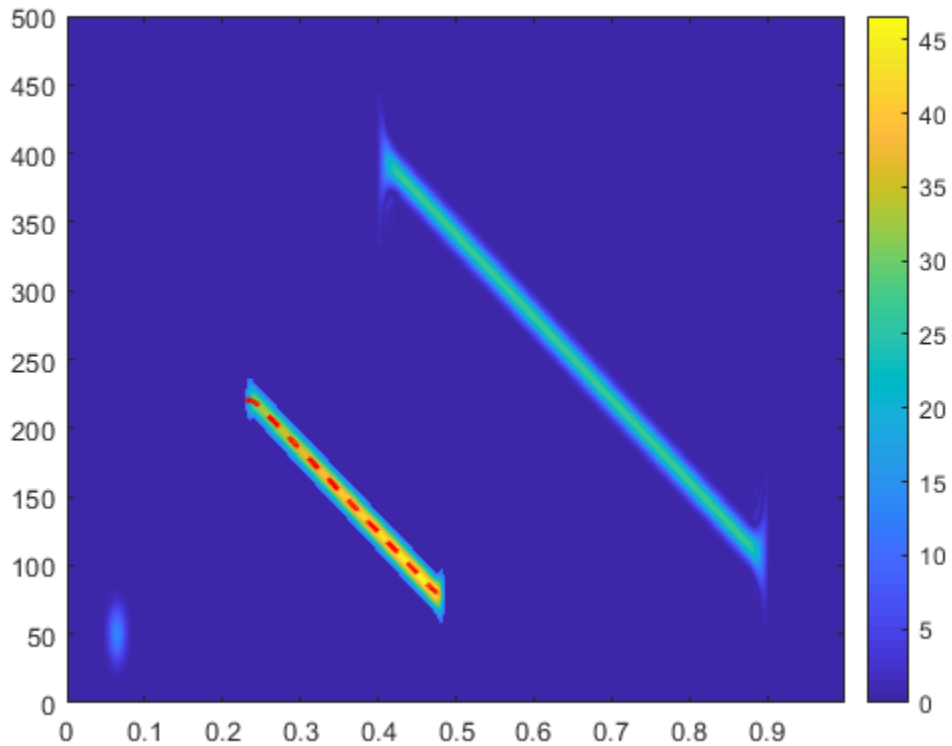
```
colorbar
```



Identify and plot the high-energy ridge corresponding to the cross-terms. To isolate the ridge, find the time values where the cross-distribution has nonzero energy.

```
ff = tfridge(c,fc);
tv = sum(c)>0;
ff = ff(tv);
tc = tc(tv);

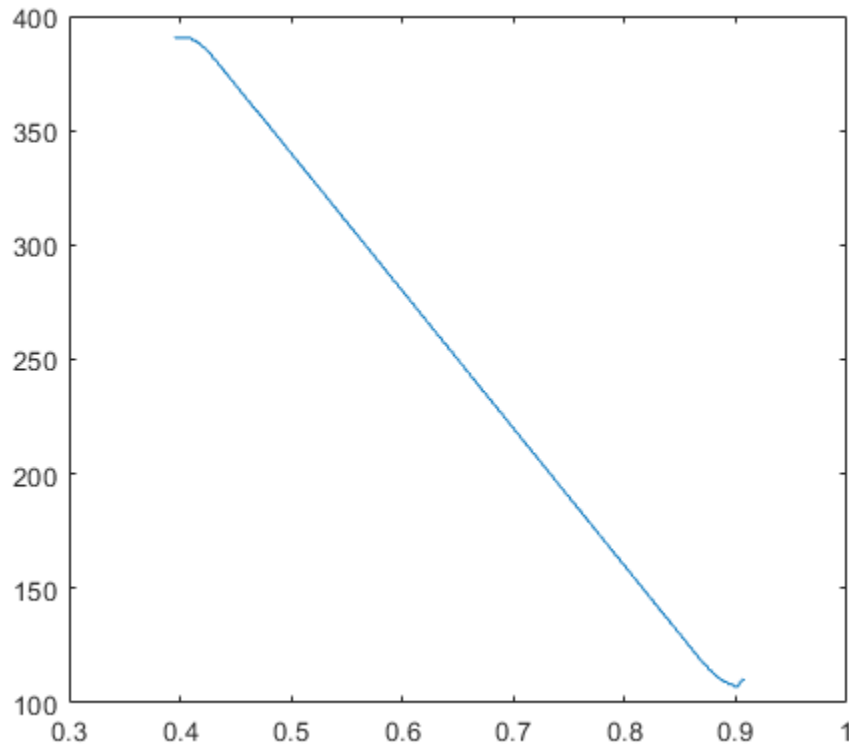
hold on
plot(tc,ff,'r--','linewidth',2)
hold off
```



Reconstruct the instantaneous frequency of the unknown signal by using the ridge and the reference function. Plot the instantaneous frequency as a function of time.

```
tEst = 2*tc - mu;  
fEst = 2*ff - fsin;
```

```
plot(tEst, fEst)
```



Input Arguments

x, y — Input signals

vectors | timetables

Input signals, specified as vectors or MATLAB timetables each containing a single vector variable. x and y must both be vectors or both be timetables and must have the same length.

- If x and y are timetables, then they must contain increasing finite row times.

- If a timetable has missing or duplicate time points, you can fix it using the tips in “Clean Timetable with Missing, Duplicate, or Nonuniform Times” (MATLAB).

If the input signals have odd length, the function appends a zero to make the length even.

Example: `cos(pi/8*(0:159))'+randn(160,1)/10` specifies a sinusoid embedded in white noise.

Example: `timetable(seconds(0:5)', rand(6,1))` specifies a random variable sampled at 1 Hz for 4 seconds.

Data Types: `single` | `double`
Complex Number Support: Yes

fs — Sample rate

`2*pi` (default) | positive numeric scalar

Sample rate, specified as a positive numeric scalar.

ts — Sample time

duration scalar

Sample time, specified as a duration scalar.

twin, fwin — Time and frequency windows

vectors of odd length

Time and frequency windows used for smoothing, specified as vectors of odd length. By default, `xwvd` uses Kaiser windows with shape factor $\beta = 20$.

- The default length of `twin` is the smallest odd integer greater than or equal to `round(length(x)/10)`.
- The default length of `fwin` is the smallest odd integer greater than or equal to `nf/4`.

Each window must have a length smaller than or equal to `2*ceil(length(x)/2)`.

Example: `kaiser(65,0.5)` specifies a 65-sample Kaiser window with a shape factor of 0.5.

nf — Number of frequency points

`2*ceil(length(x)/2)` (default) | integer

Number of frequency points, specified as an integer. This argument controls the degree of oversampling in frequency. The number of frequency points must be at least $(\text{length}(\text{fwin})+1)/2$ and cannot be greater than the default.

thresh — Minimum nonzero value

-Inf (default) | real scalar

Minimum nonzero value, specified as a real scalar. The function sets to zero those elements of \mathbf{d} whose amplitudes are less than `thresh`.

Output Arguments

d — Cross Wigner-Ville distribution

matrix

Cross Wigner-Ville distribution, returned as a matrix. Time increases across the columns of \mathbf{d} , and frequency increases down the rows. The matrix is of size $N_f \times N_t$, where N_f is the length of \mathbf{f} and N_t is the length of \mathbf{t} .

f — Frequencies

vector

Frequencies, returned as a vector.

- If the input has time information, then \mathbf{f} contains frequencies expressed in Hz.
- If the input does not have time information, then \mathbf{f} contains normalized frequencies expressed in rad/sample.

t — Time instants

vector

Time instants, returned as a vector.

- If the input has time information, then \mathbf{t} contains time values expressed in seconds.
- If the input does not have time information, then \mathbf{t} contains sample numbers.

The number of time points is fixed as $4 \cdot \text{ceil}(\text{length}(x)/2)$.

Definitions

Cross Wigner-Ville Distribution

For continuous signals $x(t)$ and $y(t)$, *cross Wigner-Ville distribution* is defined as

$$\text{XWVD}_{x,y}(t, f) = \int_{-\infty}^{\infty} x\left(t + \frac{\tau}{2}\right) y^*\left(t - \frac{\tau}{2}\right) e^{-j2\pi f\tau} d\tau.$$

For a discrete signal with N samples, the distribution becomes

$$\text{XWVD}_{x,y}(n, k) = \sum_{m=-N}^N x(n + m/2) y^*(n - m/2) e^{-j2\pi km/N}.$$

For odd values of m , the definition requires evaluation of the signal at half-integer sample values. It therefore requires interpolation, which makes it necessary to zero-pad the discrete Fourier transform to avoid aliasing.

The cross Wigner-Ville distribution contains interference terms that often complicate its interpretation. To sharpen the distribution, one can filter the definition with lowpass windows. The cross smoothed pseudo Wigner-Ville distribution uses independent windows to smooth in time and frequency:

$$\text{XSPWVD}_{x,y}^{g,H}(t, f) = \int_{-\infty}^{\infty} g(t) H(f) x\left(t + \frac{\tau}{2}\right) y^*\left(t - \frac{\tau}{2}\right) e^{-j2\pi f\tau} d\tau.$$

References

- [1] Cohen, Leon. *Time-Frequency Analysis: Theory and Applications*. Englewood Cliffs, NJ: Prentice-Hall, 1995.
- [2] Mallat, Stéphane. *A Wavelet Tour of Signal Processing*. Second Edition. San Diego, CA: Academic Press, 1999.
- [3] Malnar, Damir, Victor Sucic, and Boualem Boashash. "A cross-terms geometry based method for components instantaneous frequency estimation using the cross Wigner-Ville distribution." In *11th International Conference on Information Sciences, Signal Processing and their Applications (ISSPA)*, pp. 1217-1222. Montréal: IEEE, 2012.

See Also

Functions

wvd | xspectrogram

Introduced in R2018b

