Release Notes for Statistics Toolbox™

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Release Notes for Statistics ToolboxTM

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R2012b

Version: 8.1 New Features: Yes Bug Fixes: Yes

Boosting algorithms for imbalanced data, sparse ensembles, and multiclass boosting, with self termination

There are three new boosting algorithms for classification:

- **RUSBoost** (boosting by random undersampling) for imbalanced data (data in which one class has many more observations than the other).
- **LPBoost** (linear programming) and **TotalBoost** (totally corrective boosting) which self-terminate, can lead to a sparse ensemble, and can be used for multiclass boosting.

Burr distribution for expressing a wide range of distribution shapes while preserving a single functional form for the density

There is a new probability distribution object for the Burr Type XII distribution, a three-parameter family of continuous distributions on the real line. Use fitdist to fit this distribution to data. Use ProbDistUnivParam to specify the distribution parameters directly. Either function produces a distribution you can use to generate random samples or compute functions such as pdf and cdf.

Data import to a dataset array with the MATLAB Import Tool

You can now import data from a file directly into a $\tt dataset$ array using the $MATLAB^{\circledast}$ Import Tool.

Principal component analysis enhancements for handling NaN as missing data, weighted PCA, and choosing between EIG or SVD as the underlying algorithm

Compatibility Considerations: Yes

The new pca function includes additional functionality for principal component analysis. Features of pca include:

- Handling of NaN as missing data values.
- Weighted principal component analysis with user-specified weights.
- Choice of SVD or EIG algorithm for computing principal components.
- Option to specify number of components to return.
- Option to not center before computing principal components.

Compatibility Considerations

The new pca function replaces the princomp function.

Speedup of k-means clustering using Parallel Computing Toolbox

Statistics Toolbox[™] now supports parallel execution for kmeans.

One-sided nonparametric hypothesis tests

An option to test one-sided right- or left-tailed alternatives is available for these nonparametric hypothesis tests:

- signrank
- ranksum
- signtest

Reorder nodes in dendrogram plots

- The dendrogram function has new options for reordering the nodes of hierarchical binary cluster trees:
 - The reorder option allows you to specify a permutation vector for the order of nodes in a dendrogram plot.
 - The checkcrossings option checks whether a requested permutation vector leads to crossing branches in a dendrogram plot.
- The function optimalleaforder generates an optimal permutation of nodes.

Nonlinear model enhancements Compatibility Considerations: Yes

You can add a vector of observation weights, or a handle to a function that returns a vector of observation weights, to these functions:

- NonLinearModel.fit.
- predict and random (NonLinearModel methods).
- nlinfit and nlpredci.

For an example of weighted fitting, see Weighted Nonlinear Regression.

Compatibility Considerations

Use either $\tt Weights$ or <code>RobustWgtFun</code> when performing weighted nonlinear regression.

Changes to LinearModel diagnostics Compatibility Considerations: Yes

The diagnostics in the Diagnostics dataset array for LinearModel objects are in a new order, and no longer appear in the Variables editor. The new order is:

- Leverage
- CooksDistance
- Dffits
- S2_i
- CovRatio
- Dfbetas
- HatMatrix

Compatibility Considerations

To access the correct diagnostics, you should update any code that indexes the diagnostics dataset array columns by number.

Functionality being changed Compatibility Considerations: Yes

Functionality	What Happens When You Use This Functionality?	Use This Instead	Compatibility Considerations
princomp	Still runs	рса	Replace instances of princomp with pca

R2012a

Version: 8.0 New Features: Yes Bug Fixes: Yes

Linear, Generalized Linear, and Nonlinear Models for Regression

LinearModel is a new class for performing linear regression. LinearModel.fit creates a model that:

- Lets you fit models with both categorical and continuous predictor variables
- Contains information about the quality of the fit, such as residuals and ANOVA tables
- Lets you easily plot the fit
- Allows for automatic or manual exclusion of unimportant variables
- Enables robust fitting for reduced influence of outliers
- Lets you specify quadratic and other models using a symbolic formula
- Enables stepwise model selection

There are similar improvements for generalized linear and nonlinear modeling using the GeneralizedLinearModel and NonLinearModel classes. For details, see the class reference pages in the reference material, or Linear Regression, Stepwise Regression, Robust Regression — Reduce Outlier Effects, Generalized Linear Regression, or Nonlinear Regression in the User's Guide.

Variable Editor for Dataset Arrays

You can now edit, sort, plot, and select portions of dataset arrays from the MATLAB Variable Editor. For details, see Using Dataset Arrays in the User's Guide.

Lasso for Generalized Linear Regression

The lassoglm function regularizes generalized linear models. Use lassoglm to examine model alternatives and to constrain or remove redundant or unimportant variables in generalized linear regression. For details, see the function reference page, or Lasso Regularization of Generalized Linear Models in the User's Guide.

K-Nearest Neighbor Classification

ClassificationKNN.fit creates a classification model that performs *k*-nearest neighbor classification. You can check the quality of the model with cross validation or resubstitution. For details, see the ClassificationKNN page in the reference material, or Classification Using Nearest Neighbors in the User's Guide.

Random Subspace Ensembles

fitensemble can construct random subspace ensembles to improve the classification accuracy of both k-nearest neighbor classifiers and discriminant analysis classifiers. For details, see Ensemble Methods or Random Subspace Classification in the User's Guide.

Regularized Discriminant Analysis with Variable Selection

ClassificationDiscriminant models now have two parameters, Gamma and Delta, for regularization and lowering the number of variables. Set Gamma to regularize the discriminant. Set Delta to eliminate variables. Use cvshrink to obtain optimal Gamma and Delta parameters by cross validation. For details, see the reference pages, or Regularize a Discriminant Analysis Classifier in the User's Guide.

stepwisefit Coefficient History

The stepwisefit function now returns the fitted coefficient history in the history.B field.

RobustWgtFun Replaces WgtFun Compatibility Considerations: Yes

The WgtFun option is now called RobustWgtFun in the nlinfit, statget, and statset functions. RobustWgtFun also makes the Robust option superfluous.

Compatibility Considerations

The WgtFun and Robust options are currently accepted by all functions. To avoid potential future incompatibilities, update code that uses the WgtFun and Robust options to use the RobustWgtFun option.

ClassificationTree Now Predicts Class with Minimal Misclassification Cost Compatibility Considerations: Yes

The ClassificationTree predict method now chooses the class with minimal expected misclassification cost. Previously, it chose the class with maximal posterior probability. The new behavior is consistent with the cvLoss method. Furthermore, both ClassificationDiscriminant and ClassificationKNN predict using minimal expected misclassification cost. For details, see predict and loss.

Compatibility Considerations

If you use a nondefault cost matrix, some ClassificationTree classification predictions can differ from those in previous versions.

fpdf Improvements

The fpdf function now accepts a wider range of parameter values, including Inf.

R2011b

Version: 7.6 New Features: Yes Bug Fixes: Yes

Lasso Regularization for Linear Regression

The lasso function incorporates both the lasso regularization algorithm and the elastic net regularization algorithm. Use lasso to remove redundant or unimportant variables in linear regression. The lassoPlot function helps you visualize lasso results, with a variety of coefficient trace plots and a cross-validation plot.

For details, see Lasso and Elastic Net.

Discriminant Analysis Classification Object

You can now use the ClassificationDiscriminant and CompactClassificationDiscriminant classes for classification via discriminant analysis. The syntax and methods resemble those in the existing ClassificationTree and CompactClassificationTree classes. The ClassificationDiscriminant class includes the functionality of the classify function. ClassificationDiscriminant provides several benefits compared to the classify function:

- After you fit a classifier, you can predict without refitting.
- ClassificationDiscriminant is built on the same framework as ClassificationTree, so you have a variety of options and methods, including:
 - Cross validation
 - Resubstitution statistics
 - A choice of cost functions
 - Weighted classification
- ClassificationDiscriminant can fit several models, including linear, quadratic, and linear or quadratic with pseudoinverse.

For details, see Discriminant Analysis.

Nearest Neighbor Searching for Points Within a Fixed Distance

The rangesearch function finds all members of a data set that are within a specified distance of members of another data set. As with the knnsearch function, you can set a variety of distance metrics, or program your own. rangesearch has counterparts that are methods of the ExhaustiveSearcher and KDTreeSearcher classes.

datasample Function for Random Sampling

The datasample function samples with or without replacement from a data set. It can also perform weighted sampling, with or without replacement.

Fractional Factorial Design Improvements

The fracfactgen function now allows up to 52 factors, instead of the previous limit of 26 factors. Specify factors as case-sensitive strings, using 'a' through 'z' for the first 26 factors, and 'A' through 'Z' for the remaining factors.

fracfact now checks for an arbitrary level of interaction in confounding, instead of the previous limit of confounding up to products of two factors. Set the MaxInt name-value pair to the level of interaction you want. You can also set names for the factors using the FactorNames name-value pair.

nlmefit Returns the Covariance Matrix of Estimated Coefficients

The nlmefit function now returns the covariance matrix of the estimated coefficients as the covb field of the stats structure.

signrank Change

The signrank test now defines ties to be entries that differ by 2*eps or less. Previously, ties were entries that were identical to machine precision.

Conversion of Error and Warning Message Identifiers Compatibility Considerations: Yes

For R2011b, error and warning message identifiers have changed in Statistics Toolbox.

Compatibility Considerations

If you have scripts or functions that use message identifiers that changed, you must update the code to use the new identifiers. Typically, message identifiers are used to turn off specific warning messages, or in code that uses a try/catch statement and performs an action based on a specific error identifier.

For example, if you use the 'resubstitution' method, the 'stats:plsregress:InvalidMCReps' identifier has changed to 'stats:plsregress:InvalidResubMCReps'. If you use the 'resubstitution' method and your code checks for 'stats:plsregress:InvalidMCReps', you must update it to check for 'stats:plsregress:InvalidResubMCReps' instead.

To determine the identifier for a warning, run the following command just after you see the warning:

[MSG,MSGID] = lastwarn;

This command saves the message identifier to the variable MSGID.

To determine the identifier for an error, run the following command just after you see the error:

```
exception = MException.last;
MSGID = exception.identifier;
```

Tip Warning messages indicate a potential issue with your code. While you can turn off a warning, a suggested alternative is to change your code so it runs warning free.

R2011a

Version: 7.5 New Features: Yes Bug Fixes: Yes

Boosted Decision Trees for Classification and Regression

The new fitensemble function constructs ensembles of decision trees. It provides:

- Several popular boosting algorithms (AdaBoostM1, AdaBoostM2, GentleBoost, LogitBoost, and RobustBoost) for classification
- Least-squares boosting (LSBoost) for regression
- Most TreeBagger functionality for ensembles of bagged decision trees

There is also an improved interface for classification trees (ClassificationTree) and regression trees (RegressionTree), encompassing the functionality of classregtree.

For details, see Ensemble Methods.

Memory and Performance Improvements in Linkage Methods

The linkage and clusterdata functions have a new savememory option that can use less memory than before. With savememory set to 'on', the functions do not build a pairwise distance matrix, so use less memory and, depending on problem size, can use less time. You can use the savememory option when:

- The linkage method is 'ward', 'centroid', or 'median'
- The linkage distance metric is 'euclidean' (default)

For details, see the linkage and clusterdata function reference pages.

Conditional Weighted Residuals and Derivative Step Control in nlmefit and nlmefitsa

The nlmefit and nlmefitsa functions now provide the conditional weighted residuals of the fit. Use this information to assess the quality of the model; see Example: Examining Residuals for Model Verification.

The statset Options structure now includes 'DerivStep', which enables you to set finite differences for gradient estimation.

Detecting Ties in k-Nearest Neighbor Search

knnsearch now optionally returns all kth nearest neighbors of points, instead of just one. The knnsearch methods for ExhaustiveSearcher and KDTreeSearcher also have this option.

Distribution Fitting Tool Uses fitdist Function

MATLAB functions generated with the Distribution Fitting Tool now use the fitdist function to create fitted probability distribution objects. The generated functions return probability distribution objects as output arguments.

Speed and Accuracy Improvements in Noncentral Chi-Square CDF

 ${\tt ncx2cdf}$ is now faster and more accurate for large values of the noncentrality parameter.

Perfect Separation in Binomial Regression

If the two categories in a binomial regression model (such as logit or probit) are perfectly separated, the best-fitting model is degenerate with infinite coefficients. In this case, the glmfit function is likely to exceed its iteration limit. glmfit now tries to detect this perfect separation and display a diagnostic message.

Sign Convention in mdscale

mdscale now enforces that, in each column of the output Y, the value with the largest magnitude has a positive sign. This change makes results consistent across releases and platforms—small changes used to lead to sign reversals.

Demo of Credit Rating Classification Via Bagged Decision Trees

The credit rating demo that used to be exclusively in Financial Toolbox[™] is now available in Statistics Toolbox. The demo uses bagged decision trees for classifying creditworthiness.

To view the demo at the MATLAB command line, enter:

showdemo creditratingdemo

R2010b

Version: 7.4 New Features: Yes Bug Fixes: Yes

Parallel Computing Support for More Functions

Statistics Toolbox now supports parallel execution for the following functions:

- candexch
- cordexch
- daugment
- dcovary
- nnmf
- plsregress
- rowexch
- sequentialfs

For more information, see the Parallel Statistics chapter in the User's Guide.

Algorithm to Rank Features in Classification and Regression

New filter algorithm, relieff, is based on nearest neighbors. The ReliefF algorithm accounts for correlations among predictors by computing the effect of every predictor on the class label (or true response for regression) locally and then integrates these local estimates over the entire predictor space.

nlmefit Support for Error Models, and nlmefitsa changes Compatibility Considerations: Yes

nlmefit now supports the following error models:

- combined
- constant
- exponential
- proportional

You can specify an error model with both nlmefitsa and nlmefit.

The nlmefit bic calculation has changed. Now the degrees of freedom value is based on the number of groups rather than the number of observations. This conforms with the bic definition used by the nlmefitsa function.

Both nlmefit and nlmefitsa now store the estimated error parameters in the errorparm field of the output stats structure. The rmse field of the structure now contains the root mean squared residual for all error models; this value is computed on the log scale for the exponential model.

Compatibility Considerations

In the previous release, the rmse field was used by nlmefitsa for both mean squared residual and the estimated error parameter. Change your code, if necessary, to address the appropriate field in the stats structure.

As described in "nlmefit Support for Error Models, and nlmefits changes" on page 48, nlmefit now calculates different bic values than in previous releases.

Surrogate Splits for Decision Trees

The new surrogate splits feature in classregtree allows for better handling of missing values, more accurate estimation of variable importance, and calculation of the predictive measure of association between variables.

New Bagged Decision Tree Properties

TreeBagger and CompactTreeBagger classes have two new properties:

- NVarSplit provides the number of decision splits for each predictor variable.
- VarAssoc provides a measure of association between pairs of predictor variables.

Enhanced Cluster Analysis Performance

The linkage function has improved performance for the centroid, median, and single linkage methods.

The linkage and pdist hierarchical cluster analysis functions support larger array dimensions with 64-bit platforms, so can handle larger problems.

Export Probability Objects with dfittool Compatibility Considerations: Yes

The distribution fitting GUI (dfittool) now allows you to export fits to the MATLAB workspace as probability distribution fit objects. For more information, see Modeling Data Using the Distribution Fitting Tool.

Compatibility Considerations

If you load a distribution fitting session that was created with previous versions of Statistics Toolbox, you cannot save an existing fit. Fit the distribution again to enable saving.

Compute Partial Correlation of Two Variables Correcting for All Other Variables

partialcorr now accepts a new syntax, RHO = partialcorr(X), which returns the sample linear partial correlation coefficients between pairs of variables in X, controlling for the remaining variables in X. For more information, see the function reference page.

Specify Number of Evenly Spaced Quantiles

quantile now accepts a new syntax, Y = quantile(X,N,...), which returns quantiles at the cumulative probabilities (1:N)/(N+1) where N is a scalar positive integer value.

Control Location and Orientation of Marginal Histograms with scatterhist

scatterhist now accepts three parameter name/value pairs that control where and how the histogram plots appear. The new parameter names are NBins, Location, and Direction. For more information, see the function reference page.

Return Bootstrapped Statistics with bootci

bootci has a new output option which returns the bootstrapped statistic computed for each of the NBoot bootstrap replicate samples. For more information, see the function reference page.

R2010a

Version: 7.3 New Features: Yes Bug Fixes: Yes

Stochastic Algorithm Functionality in NLME Models

New stochastic algorithm for fitting NLME models is more robust with respect to starting values, enables parameter transformations, and relaxes assumption of constant error variance. See nlmefitsa.

k-Nearest Neighbor Searching

New functions for k-Nearest Neighbor (kNN) search efficiently to find the closest points to any query point. For information, see k-Nearest Neighbor Search and Radius Search.

Confidence Intervals Option in perfcurve

A new option in the perfcurve function computes confidence intervals for classifier performance curves.

Observation Weights Options in Resampling Functions

New options to weight resampling probabilities broaden the range of models supported by bootstrp, bootci, and perfcurve functions.

R2009b

Version: 7.2 New Features: Yes Bug Fixes: Yes

New Parallel Computing Support for Certain Functions

Statistics Toolbox now supports parallel execution for the following functions:

- bootci
- bootstrp
- crossval
- jackknife
- TreeBagger

For more information on parallel computing in the Statistics Toolbox, see Parallel Computing Support for Resampling Methods.

New Stack and Unstack Methods for Dataset Arrays

dataset.unstack converts a "tall" dataset array to an equivalent dataset array that is in "wide format", by "unstacking" a single variable in the tall dataset array into multiple variables in wide. dataset.stack reverses this manipulation by converting a "wide" dataset array to an equivalent dataset array that is in "tall format", by "stacking up" multiple variables in the wide dataset array into a single variable in tall.

New Support for SAS Transport (.xpt) Files

Statistics Toolbox now supports importing and exporting files in SAS Transport (.xpt) format. For more information, see the xptread and dataset.export reference pages.

New Output Function in nlmefit for Monitoring or Canceling Calculations

The nlmefit function now supports using an output function to monitor or cancel calculations. For more information, see the nlmefit reference page.

R2009a

Version: 7.1 New Features: Yes Bug Fixes: Yes

Enhanced Dataset Functionality

- An enhanced dataset.join method provides additional types of join operations:
 - join can now perform more complicated inner and outer join operations that allow a many-to-many correspondence between dataset arrays A and B, and allow unmatched observations in either A or B.
 - join can be of Type 'inner', 'leftouter', 'rightouter', 'fullouter', or 'outer' (which is a synonym for 'fullouter'). For an inner join, the dataset array, C, only contains observations corresponding to a combination of key values that occurred in both A and B. For a left (or right) outer join, C also contains observations corresponding to keys in A (or B) that did not match any in B (or A).
 - join can now return index vectors indicating the correspondence between observations in C and those in A and B.
 - join now supports using multiple keys.
 - join now supports an optional parameter for specifying missing key behavior rather than raising an error.
- An enhanced dataset.export method now supports exporting directly to $Microsoft^{\circledast}\ Excel^{\circledast}$ files.

New Naïve Bayes Classification

- The NaiveBayes classification object is suitable for data sets that contain many predictors or features.
- It supports normal, kernel, multinomial, and multivariate multinomial distributions.

New Ensemble Methods for Classification and Regression Trees

- New classification objects, TreeBagger and CompactTreeBagger, provide improved performance through bootstrap aggregation (bagging).
- Includes Breiman's "random forest" method.
- Enhanced classregtree has more options for growing and pruning trees.

New Performance Curve Function

- New perfcurve function provides graphical method to evaluate classification results.
- Includes ROC (receiver operating characteristic) and other curves.

New Probability Distribution Objects

- Provides a consistent interface for working with probability distributions.
- Can be created directly using the ProbDistUnivParam constructor, or fit to data using the fitdist function.
- Option to fit distributions by group.
- Includes kernel object methods and parametric object methods that you can use to analyze the distribution represented by the object.
- Includes kernel object properties and parametric object properties that you can access to determine the fit results and evaluate their accuracy.
- Related enhancements in the chi2gof, histfit, kstest, probplot, and qqplot functions.

R2008b

Version: 7.0 New Features: Yes Bug Fixes: No

Classification

The new confusionmat function tabulates misclassifications by comparing known and predicted classes of observations.

Data Organization Compatibility Considerations: Yes

Dataset arrays constructed by the dataset function can now be written to an external text file using the new export function.

When reading external text files into a dataset array, dataset has a new 'TreatAsEmpty' parameter for specifying strings to be treated as empty.

Compatibility Considerations

In previous versions, dataset used eval to evaluate strings in external text files before writing them into a dataset array. As a result, strings such as $^1/1/2008'$ were treated as numerical expressions with two divides. Now, dataset treats such expressions as strings, and writes a string variable into the dataset array whenever a column in the external file contains a string that does not represent a valid scalar value.

Model Assessment

The cross-validation function, crossval, has new options for directly specifying loss functions for mean-squared error or misclassification rate, without having to provide a separate function M-file.

Multivariate Methods

The procrustes function has new options for computing linear transformations without scale or reflection components.

Probability Distributions Compatibility Considerations: Yes

The multivariate normal functions mvnpdf, mvncdf, and mvnrnd now accept vector specification of diagonal covariance matrices, with corresponding gains in computational efficiency.

The hypergeometric distribution has been added to both the disttool and randtool graphical user interfaces.

Compatibility Considerations

The ksdensity function may give different answers for the case where there are censoring times beyond the last observed value. In this case, ksdensity tries to reduce the bias in its density estimate by folding kernel functions across a folding point so that they do not extend into the area that is completely censored. Two things have changed for this release:

- **1** In previous releases the folding point was the last observed value. In this release it is the first censoring time after the last observed value.
- **2** The folding procedure is applied not just when the 'function' parameter is 'pdf', but for all 'function' values.

Regression Analysis

The new nlmefit function fits nonlinear mixed-effects models to data with both fixed and random sources of variation. Mixed-effects models are commonly used with data over multiple groups, where measurements are correlated within groups but independent between groups.

Statistical Visualization Compatibility Considerations: Yes

The boxplot function has new options for handling multiple grouping variables and extreme outliers.

The lsline, gline, refline, and refcurve functions now work with scatter plots produced by the scatter function. In previous versions, these functions worked only with scatter plots produced by the plot function.

The following visualization functions now have custom data cursors, displaying information such as observation numbers, group numbers, and the values of related variables:

- andrewsplot
- biplot
- ecdf
- glyphplot
- gplotmatrix
- gscatter
- normplot
- parallelcoords
- probplot
- qqplot
- scatterhist
- wblplot

Compatibility Considerations

Changes to boxplot have altered a number of default behaviors:

• Box labels are now drawn as text objects rather than tick labels. Any code that customizes the box labels by changing tick marks should now set the tick locations as well as the tick labels.

- The function no longer returns a handles array with a fixed number handles, and the order and meaning of the handles now depends on which options are selected. To locate a handle of interest, search for its 'Tag' property using findobj. 'Tag' values for box plot components are listed on the boxplot reference page.
- There are now valid handles for outliers, even when boxes have no outliers. In previous releases, the handles array returned by the function had NaN values in place of handles when boxes had no outliers. Now the 'xdata' and 'ydata' for outliers are NaN when there are no outliers.
- For small groups, the 'notch' parameter sometimes produces notches that extend outside of the box. In previous releases, the notch was truncated to the extent of the box, which could produce a misleading display. A new value of 'markers' for this parameter avoids the display issue.

As a consequence, the anoval function, which displays notched box plots for grouped data, may show notches that extend outside the boxes.

Utility Functions

The statistics options structure created by statset now includes a Jacobian field to specify whether or not an objective function can return the Jacobian as a second output.

R2008a

Version: 6.2 New Features: Yes Bug Fixes: Yes

Descriptive Statistics Compatibility Considerations: Yes

Bootstrap confidence intervals computed by **bootci** are now more accurate for lumpy data.

Compatibility Considerations

The formula for bootci confidence intervals of type 'bca' or 'cper' involves the proportion of bootstrap statistics less than the observed statistic. The formula now takes into account cases where there are many bootstrap statistics exactly equal to the observed statistic.

Model Assessment

Two new cross-validation functions, cvpartition and crossval, partition data and assess models in regression, classification, and clustering applications.

Multivariate Methods

A new sequential feature selection function, sequentialfs, selects predictor subsets that optimize user-defined prediction criteria.

The new nnmf function performs nonnegative matrix factorization (NMF) for dimension reduction.

Probability Distributions

The new sobolset and haltonset functions produce quasi-random point sets for applications in Monte Carlo integration, space-filling experimental designs, and global optimization. Options allow you to skip, leap over, and scramble the points. The qrandstream function provides corresponding quasi-random number streams for intermittent sampling.

Regression Analysis

The new plsregress function performs partial least-squares regression for data with correlated predictors.

Statistical Visualization

The normspec function now shades regions of a normal density curve that are either inside or outside specification limits.

Utility Functions

The statistics options structure created by statset now includes fields for TolTypeFun and TolTypeX, to specify tolerances on objective functions and parameter values, respectively.

R2007b

Version: 6.1 New Features: Yes Bug Fixes: Yes

Cluster Analysis Compatibility Considerations: Yes

The new gmdistribution class represents Gaussian mixture distributions, where random points come from different multivariate normal distributions with certain probabilities. The gmdistribution constructor creates mixture models with specified means, covariances, and mixture proportions, or by fitting a mixture model with a specified number of components to data. Methods for the class include:

- fit Distribution fitting function
- pdf Probability density function
- cdf Cumulative distribution function
- random Random number generator
- cluster Data clustering
- posterior Cluster posterior probabilities
- mahal Mahalanobis distance

The cluster function for hierarchical clustering now accepts a vector of cutoff values, and returns a matrix of cluster assignments, with one column per cutoff value.

Compatibility Considerations

The kmeans function now returns a vector of cluster indices of length n, where n is the number of rows in the input data matrix X, even when X contains NaN values. In the past, rows of X with NaN values were ignored, and the vector of cluster indices was correspondingly reduced in size. Now the vector of cluster indices contains NaN values where rows have been ignored, consistent with other toolbox functions.

Design of Experiments

A new option in the *D*-optimal design function candexch specifies fixed design points in the row-exchange algorithm. A similar feature is already available for the daugment function, which uses the coordinate-exchange algorithm.

Hypothesis Tests Compatibility Considerations: Yes

The kstest function now uses a more accurate method to calculate the *p*-value for a single-sample Kolmogorov-Smirnov test.

Compatibility Considerations

kstest now compares the computed *p*-value to the desired cutoff, rather than comparing the test statistic to a table of values. Results may differ from those in previous releases, especially for small samples in two-sided tests where an asymptotic formula was used in the past.

Probability Distributions Compatibility Considerations: Yes

A new fitting function, copulafit, has been added to the family of functions that describe dependencies among variables using copulas. The function fits parametric copulas to data, providing a link between models of marginal distributions and models of data correlations.

A number of probability functions now have improved accuracy, especially for extreme parameter values. The functions are:

- betainv More accurate for probabilities in P near 1.
- binocdf More efficient and less likely to run out of memory for large values in X.
- binopdf More accurate when the probabilities in P are on the order of eps.
- fcdf More accurate when the parameter ratios V2./V1 are much less than the values in X.
- ncx2cdf More accurate in some extreme cases that previously returned
 0.
- poisscdf More efficient and less likely to run out of memory for large values in X.
- tcdf More accurate when the squares of the values in X are much less than the parameters in V.
- tinv More accurate when the probabilities in P are very close to 0.5 and the outputs are very small in magnitude.

Function-style syntax for paretotails objects has been removed.

Compatibility Considerations

The changes to the probability functions listed above may lead to different, but more accurate, outputs than in previous releases.

In previous releases, syntax of the form obj(x) for a paretotails objects obj invoked the cdf method. This syntax now produces a warning. To evaluate the cumulative distribution function, use the syntax cdf(obj,x).

Regression Analysis Compatibility Considerations: Yes

The new corrcov function converts a covariance matrix to the corresponding correlation matrix.

The mvregress function now supports an option to force the estimated covariance matrix to be diagonal.

Compatibility Considerations

In previous releases the mvregress function, when using the 'cwls' algorithm, estimated the covariance of coefficients COVB using the estimated, rather than the initial, covariance of the responses SIGMA. The initial SIGMA is now used, and COVB differs to a degree dependent on the difference between the initial and final estimates of SIGMA.

Statistical Visualization

The boxplot function has a new 'compact' plot style suitable for displaying large numbers of groups.

R2007a

Version: 6.0 New Features: Yes Bug Fixes: Yes

Data Organization

New categorical and dataset arrays are available for organizing and processing statistical data.

- Categorical arrays facilitate the use of nominal and ordinal categorical data.
- Dataset arrays provide a natural way to encapsulate heterogeneous statistical data and metadata, so that it can be accessed and manipulated using familiar methods analogous to those for numerical matrices.
- Categorical and dataset arrays are supported by a variety of new functions for manipulating the encapsulated data.
- Categorical arrays are now accepted as input arguments in all Statistics Toolbox functions that make use of grouping variables.

Hypothesis Testing

Expanded options are available for linear hypothesis testing.

- The new linhyptest function performs linear hypothesis tests on parameters such as regression coefficients. These tests have the form H*b = c for specified values of H and c, where b is a vector of unknown parameters.
- The covb output from regstats and the SIGMA output from nlinfit are suitable for use as the covariance matrix input argument required by linhyptest. The following functions have been modified to return a covb output for use with linhyptest: coxphfit, glmfit, mnrfit, robustfit.

Multivariate Statistics Compatibility Considerations: Yes

The new cholcov function computes a Cholesky-like decomposition of a covariance matrix, even if the matrix is not positive definite. Factors are useful in many of the same ways as Cholesky factors, such as imposing correlation on random number generators.

The classify function for discriminant analysis has been improved.

- The function now computes the coefficients of the discriminant functions that define boundaries between classification regions.
- The output of the function is now of the same type as the input grouping variable group.

Compatibility Considerations

The classify function now returns outputs of different type than it did in the past. If the input argument group is a logical vector, output is now converted to a logical vector. In the past, output was returned as a cell array of 0s and 1s. If group is numeric, the output is now converted to the same type. For example, if group is of type uint8, the output will be of type uint8.

Probability Distributions

New paretotails objects are available for modeling distributions with an empirical cdf or similar distribution in the center and generalized Pareto distributions in the tails.

- The paretotails function converts a data sample to a paretotails object. The objects are useful for generating random samples from a distribution similar to the data, but with tail behavior that is less discrete than the empirical distribution.
- Objects from the paretotails class are supported by a variety of new methods for working with the piecewise distribution.
- The paretotails class provides function-like behavior, so that p(x) evaluates the cdf of p at values x.

Regression Analysis Compatibility Considerations: Yes

The new mvregresslike function is a utility related to the mvregress function for fitting regression models to multivariate data with missing values. The new function computes the objective (log likelihood) function, and can also compute the estimated covariance matrix for the parameter estimates.

New classregtree objects are available for creating and analyzing classification and regression trees.

- The classregtree function fits a classification or regression tree to training data. The objects are useful for predicting response values from new predictors.
- Objects from the classregtree class are supported by a variety of new methods for accessing information about the tree.
- The classregtree class provides function-like behavior, so that t(X) evaluates the tree t at predictor values in X.
- The following functions now create or operate on objects from the new classregtree class: treefit, treedisp, treeval, treefit, treeprune, treetest.

Compatibility Considerations

Objects from the classregtree class are intended to be compatible with the structure arrays that were produced in previous versions by the classification and regression tree functions listed above. In particular, classregtree supports dot indexing of the form t.property to obtain properties of the object t. The class also provides function-like behavior through parenthesis indexing, so that t(x) uses the tree t to classify or compute fitted values for predictors x, rather than index into t as a structure array as it did in the past. As a result, cell arrays should now be used to aggregate classregtree objects.

Statistical Visualization

The new scatterhist function produces a scatterplot of 2D data and illustrates the marginal distributions of the variables by drawing histograms along the two axes. The function is also useful for viewing properties of random samples produced by functions such as copularnd, mvnrnd, and lhsdesign.

Other Improvements

- The mvtrnd function now produces a single random sample from the multivariate *t* distribution if the cases input argument is absent.
- The zscore function, which centers and scales input data by mean and standard deviation, now returns the means and standard deviations as additional outputs.

R2006b

Version: 5.3 New Features: Yes Bug Fixes: Yes

Demos

The following demo has been updated:

• Selecting a Sample Size — Modified to highlight the new sampsizepwr function

Design of Experiments

The following visualization functions, commonly used in the design of experiments, have been added:

- \bullet interactionplot Two-factor interaction plot for the mean
- maineffectsplot Main effects plot for the mean
- $\bullet \ {\tt multivarichart} {\tt Multivari} \ {\tt chart} \ {\tt for} \ {\tt the} \ {\tt mean}$

Hypothesis Tests Compatibility Considerations: Yes

The following functions for hypothesis testing have been added or improved:

- jbtest Replaces the chi-square approximation of the test statistic, which is asymptotic, with a more accurate algorithm that interpolates *p*-values from a table of quantiles. A new option allows you to run Monte Carlo simulations to compute *p*-values outside of the table.
- lillietest Uses an improved version of Lilliefors' table of quantiles, covering a wider range of sample sizes and significance levels, with more accurate values. New options allow you to test for exponential and extreme value distributions, as well as normal distributions, and to run Monte Carlo simulations to compute *p*-values outside of the tables.
- runstest Adds a test for runs up and down to the existing test for runs above or below a specified value.
- sampsizepwr New function to compute the sample size necessary for a test to have a specified power. Options are available for choosing a variety of test types.

Compatibility Considerations

If the significance level for a test lies outside the range of tabulated values, [0.001, 0.5], then both jbtest and lillietest now return an error. In previous versions, jbtest returned an approximate *p*-value and lillietest returned an error outside a smaller range, [0.01, 0.2]. Error messages suggest using the new Monte Carlo option for computing values outside the range of tabulated values.

If the data sample for a test leads to a *p*-value outside the range of tabulated values, then both jbtest and lillietest now return, with a warning, either the smallest or largest tabulated value. In previous versions, jbtest returned an approximate *p*-value and lillietest returned NaN.

Multinomial Distribution

The multinomial distribution has been added to the list of almost 50 probability distributions supported by the toolbox.

- $\bullet \ {\tt mnpdf} {\tt Multinomial probability density function} \\$
- mnrnd Multinomial random number generator

Regression Analysis

Multinomial Regression

Support has been added for multinomial regression modeling of discrete multi-category response data, including multinomial logistic regression. The following new functions supplement the regression models in glmfit and glmval by providing for a wider range of response values:

- mnrfit Fits a multinomial regression model to data
- mnrval Computes predicted probabilities for the multinomial regression model

Multivariate Regression

The new mvregress function carries out multivariate regression on data with missing response values. An option allows you to specify how missing data is handled.

Survival Analysis

coxphfit — A new option allows you to specify the values at which the baseline hazard is computed.

Statistical Process Control Compatibility Considerations: Yes

The following new functions consolidate and expand upon existing functions for statistical process control:

- capability Computes a wider range of probabilities and capability indices than the capable function found in previous releases
- controlchart Displays a wider range of control charts than the ewmaplot, schart, and xbarplot functions found in previous releases
- controlrules Supplements the new controlchart function by providing for a wider range of control rules (Western Electric and Nelson)
- gagerr Performs a gage repeatability and reproducibility study on measurements grouped by operator and part

Compatibility Considerations

The capability function subsumes the capable function that appeared in previous versions of Statistics Toolbox software, and the controlchart function subsumes the functions ewmaplot, schart, and xbarplot. The older functions remain in the toolbox for backwards compatibility, but they are no longer documented or supported.

R2006a

Version: 5.2 New Features: Yes Bug Fixes: Yes

Analysis of Variance

Support for nested and continuous factors has been added to the anovan function for N-way analysis of variance.

Bootstrapping

The following functions have been added to supplement the existing bootstrp function for bootstrap estimation:

- bootci Computes confidence intervals of a bootstrapped statistic. An option allows you to choose the type of the bootstrap confidence interval.
- jackknife Draws jackknife samples from a data set and computes statistics on each sample

Demos

The following demos have been added to the toolbox:

- Bayesian Analysis for a Logistic Regression Model
- Time Series Regression of Airline Passenger Data

The following demo has been updated to demonstrate new features:

• Random Number Generation

Design of Experiments

The new fracfactgen function finds a set of fractional factorial design generators suitable for fitting a specified model.

The following functions for *D*-optimal designs have been enhanced:

- cordexch, daugment, dcovary, rowexch New options specify the range of values and the number of levels for each factor, exclude factor combinations, treat factors as categorical rather than continuous, control the number of iterations, and repeat the design generation process from random starting points
- candexch New options control the number of iterations and repeat the design generation process from random starting points
- candgen New options specify the range of values and the number of levels for each factor, and treat factors as categorical rather than continuous
- x2fx New option treats factors as categorical rather than continuous

Hypothesis Tests

The new ${\tt dwtest}$ function performs a Durbin-Watson test for autocorrelation in linear regression.

Multivariate Distributions

Two new functions have been added to compute multivariate cdfs. These supplement existing functions for pdfs and random number generators for the same distributions.

- $\bullet\,$ mvncdf Cumulative distribution function for the multivariate normal distribution
- mvtcdf Cumulative distribution function for the multivariate t distribution

Random Number Generation

Copulas

New functions have been added to the toolbox that allow you to use copulas to model correlated multivariate data and generate random numbers from multivariate distributions.

- copulacdf Cumulative distribution function for a copula
- copulaparam Copula parameters as a function of rank correlation
- copulapdf Probability density function for a copula
- copularnd Random numbers from a copula
- copulastat Rank correlation for a copula

Markov Chain Monte Carlo Methods

The following functions generate random numbers from nonstandard distributions using Markov Chain Monte Carlo methods:

- mhsample Generate random numbers using the Metropolis-Hasting algorithm
- slicesample Generate random numbers using a slice sampling algorithm

Pearson and Johnson Systems of Distributions

Support has been added for random number generation from Pearson and Johnson systems of distributions.

- pearsrnd Random numbers from a distribution in the Pearson system
- johnsrnd Random numbers from a distribution in the Johnson system

Robust Regression

To supplement the robustfit function, the following functions now have options for robust fitting:

- nlinfit Nonlinear least-squares regression
- nlparci Confidence intervals for parameters in nonlinear regression
- nlpredci Confidence intervals for predictions in nonlinear regression

Statistical Process Control

The following control chart functions now support time-series objects:

- xbarplot Xbar plot
- schart Standard deviation chart
- ewmaplot Exponentially weighted moving average plot

R14SP3

Version: 5.1 New Features: Yes Bug Fixes: No

Demos

The following demos have been added to the toolbox:

- Curve Fitting and Distribution Fitting
- Fitting a Univariate Distribution Using Cumulative Probabilities
- Fitting an Orthogonal Regression Using Principal Components Analysis
- Modelling Tail Data with the Generalized Pareto Distribution
- Pitfalls in Fitting Nonlinear Models by Transforming to Linearity
- Weighted Nonlinear Regression

The following demo has been updated:

• Modelling Data with the Generalized Extreme Value Distribution

Descriptive Statistics

The new partialcorr function computes the correlation of one set of variables while controlling for a second set of variables.

The grpstats function now computes a wider variety of descriptive statistics for grouped data. Choices include the mean, standard error of the mean, number of elements, group name, standard deviation, variance, confidence interval for the mean, and confidence interval for new observations. The function also supports the computation of user-defined statistics.

Hypothesis Tests

Chi-Square Goodness-of-Fit Test

The new chi2gof function tests if a sample comes from a specified distribution, against the alternative that it does not come from that distribution, using a chi-square test statistic.

Variance Tests

Three functions have been added to test sample variances:

- vartest One-sample chi-square variance test. Tests if a sample comes from a normal distribution with specified variance, against the alternative that it comes from a normal distribution with a different variance.
- vartest2 Two-sample *F*-test for equal variances. Tests if two independent samples come from normal distributions with the same variance, against the alternative that they come from normal distributions with different variances.
- vartestn Bartlett multiple-sample test for equal variances. Tests if multiple samples come from normal distributions with the same variance, against the alternative that they come from normal distributions with different variances.

Ansari-Bradley Test

The new ansaribradley function tests if two independent samples come from the same distribution, against the alternative that they come from distributions that have the same median and shape but different variances.

Tests of Randomness

The new runstest function tests if a sequence of values comes in random order, against the alternative that the ordering is not random.

Probability Distributions

Support has been added for two new distributions:

- "Generalized Extreme Value Distribution" on page 131
- "Generalized Pareto Distribution" on page 131

Generalized Extreme Value Distribution

The Generalized Extreme Value distribution combines the Gumbel, Frechet, and Weibull distributions into a single distribution. It is used to model extreme values in data.

The following distribution functions have been added:

- gevcdf Cumulative distribution function
- gevfit Parameter estimation function
- gevinv Inverse cumulative distribution function
- gevlike Negative log-likelihood function
- gevpdf Probability density function
- gevrnd Random number generator
- gevstat Distribution statistics

Generalized Pareto Distribution

The Generalized Pareto distribution is used to model the tails of a data distribution.

The following distribution functions have been added:

- gpcdf Cumulative distribution function
- gpfit Parameter estimation function
- $\bullet ~ {\tt gpinv} {\tt Inverse} ~ {\tt cumulative} ~ {\tt distribution} ~ {\tt function} \\$
- gplike Negative log-likelihood function
- gppdf Probability density function

- $\bullet \ {\tt gprnd} {\tt Random \ number \ generator}$
- $\bullet \ {\tt gpstat} {\tt Distribution \ statistics}$

Regression Analysis

- The new coxphfit function fits Cox's proportional hazards regression model to data.
- The new invpred function estimates the inverse prediction intervals for simple linear regression.
- The polyconf function has new options to let you specify the confidence interval computed.

Statistical Visualization

Both the $\verb|ecdf|$ and $\verb|ksdensity|$ functions now produce plots when no output arguments are specified.

R14SP2

Version: 5.0.2 New Features: Yes Bug Fixes: Yes

Multivariate Statistics

The cophenet function now returns cophenetic distances as well as the cophenetic correlation coefficient.