

## nag\_mv\_cluster\_indicator (g03ejc)

### 1. Purpose

**nag\_mv\_cluster\_indicator (g03ejc)** computes a cluster indicator variable from the results of nag\_mv\_hierar\_cluster\_analysis (g03ecc).

### 2. Specification

```
#include <nag.h>
#include <nagg03.h>

void nag_mv_cluster_indicator(Integer n, double cd[], Integer iord[],
    double dord[], Integer *k, double *dlevel, Integer ic[],
    NagError *fail)
```

### 3. Description

Given a distance or dissimilarity matrix for  $n$  objects, cluster analysis aims to group the  $n$  objects into a number of more or less homogeneous groups or clusters. With agglomerative clustering methods (see nag\_mv\_hierar\_cluster\_analysis (g03ecc)), a hierarchical tree is produced by starting with  $n$  clusters each with a single object and then at each of  $n - 1$  stages, merging two clusters to form a larger cluster until all objects are in a single cluster. nag\_mv\_cluster\_indicator takes the information from the tree and produces the clusters that exist at a given distance. This is equivalent to taking the dendrogram (see nag\_mv\_dendrogram (g03ehc)) and drawing a line across at a given distance to produce clusters.

As an alternative to giving the distance at which clusters are required, the user can specify the number of clusters required and nag\_mv\_cluster\_indicator will compute the corresponding distance. However, it may not be possible to compute the number of clusters required due to ties in the distance matrix.

If there are  $k$  clusters then the indicator variable will assign a value between 1 and  $k$  to each object to indicate to which cluster it belongs. Object 1 always belongs to cluster 1.

### 4. Parameters

#### **n**

Input: the number of objects,  $n$ .

Constraint:  $n \geq 2$ .

#### **cd[n–1]**

Input: the clustering distances in increasing order as returned by nag\_mv\_hierar\_cluster\_analysis (g03ecc).

Constraint:  $cd[i] \geq cd[i - 1]$  for  $i = 1, 2, \dots, n - 2$ .

#### **iord[n]**

Input: the objects in the dendrogram order as returned by nag\_mv\_hierar\_cluster\_analysis (g03ecc).

#### **dord[n]**

Input: the clustering distances corresponding to the order in **iord**.

#### **k**

Input: indicates if a specified number of clusters is required.

If  $k > 0$ , then nag\_mv\_cluster\_indicator (g03ejc) will attempt to find  $k$  clusters.

If  $k \leq 0$ , then nag\_mv\_cluster\_indicator (g03ejc) will find the clusters based on the distance given in **dlevel**.

Constraint:  $k \leq n$ .

Output: the number of clusters produced,  $k$ .

**dlevel**

Input: if  $k \leq 0$ , then **dlevel** must contain the distance at which clusters are produced. Otherwise **dlevel** need not be set.

Constraint: if  $k \leq 0$  then **dlevel**  $> 0.0$ .

Output: if  $k > 0$  on entry, then **dlevel** contains the distance at which the required number of clusters are found. Otherwise **dlevel** remains unchanged.

**ic[n]**

Output: **ic**[ $i - 1$ ] indicates to which of  $k$  clusters the  $i$ th object belongs, for  $i = 1, 2, \dots, n$ .

**fail**

The NAG error parameter, see the Essential Introduction to the NAG C Library.

## 5. Error Indications and Warnings

**NE\_INT\_ARG\_LT**

On entry, **n** must not be less than 2:  $\mathbf{n} = \langle \text{value} \rangle$ .

**NE\_2\_INT\_ARG\_GT**

On entry,  $k = \langle \text{value} \rangle$  while  $\mathbf{n} = \langle \text{value} \rangle$ .

These parameters must satisfy  $k \leq n$ .

**NE\_REAL\_INT**

On entry, **dlevel** =  $\langle \text{value} \rangle$ ,  $\mathbf{k} = \langle \text{value} \rangle$ .

Constraint:  $k \leq 0$  and **dlevel**  $> 0.0$ .

**NE\_NOT\_INCREASING**

The sequence **cd** is not increasing:

$\mathbf{cd}[\langle \text{value} \rangle] = \langle \text{value} \rangle$ ,  $\mathbf{cd}[\langle \text{value} \rangle] = \langle \text{value} \rangle$ .

**NW\_REAL\_REALARR**

On entry, **dlevel** =  $\langle \text{value} \rangle$ ,  $\mathbf{cd}[\langle \text{value} \rangle] = \langle \text{value} \rangle$ .

Trivial solution returned.

**NW\_INT**

On exit,  $k = 1$ .

Trivial solution returned.

**NW\_2\_INT**

On exit,  $k = \langle \text{value} \rangle$ ,  $\mathbf{n} = \langle \text{value} \rangle$ .

Trivial solution returned.

**NE\_INCOMP\_ARRAYS**

Arrays **cd** and **dord** are not compatible.

**NE\_CLUSTER**

The precise number of clusters requested is not possible because of tied clustering distances. The actual number of clusters produced is  $\langle \text{value} \rangle$ .

**NE\_INTERNAL\_ERROR**

An internal error has occurred in this function. Check the function call and any array sizes. If the call is correct then please consult NAG for assistance.

## 6. Further Comments

A fixed number of clusters can be found using the non-hierarchical method used in nag\_mv\_kmeans\_cluster\_analysis (g03efc).

### 6.1. Accuracy

The accuracy will depend upon the accuracy of the distances in **cd** and **dord** (see nag\_mv\_hierar\_cluster\_analysis (g03ecc)).

### 6.2. References

Everitt B S (1974) *Cluster Analysis* Heinemann.

Krzanowski W J (1990) *Principles of Multivariate Analysis* Oxford University Press.

## 7. See Also

`nag_mv_kmeans_cluster_analysis (g03efc)`  
`nag_mv_hierar_cluster_analysis (g03ecc)`

## 8. Example

Data consisting of three variables on five objects are input. Euclidean squared distances are computed using `nag_mv_distance_mat (g03eac)` and median clustering performed using `nag_mv_hierar_cluster_analysis (g03ecc)`. A dendrogram is produced by `nag_mv_dendrogram (g03ehc)` and printed. `nag_mv_cluster_indicator` finds two clusters and the results are printed.

### 8.1. Program Text

```
/* nag_mv_cluster (g03ejc) Example Program.
*
* Copyright 1998 Numerical Algorithms Group.
*
* Mark 5, 1998.
*
* Mark 6 revised, 2000.
*/
#include <nag.h>
#include <stdio.h>
#include <nag_stdlib.h>
#include <nagg03.h>

#define NMAX 10
#define MMAX 10

main()
{
    double cd[NMAX-1], d[NMAX*(NMAX-1)/2], dord[NMAX],
           s[MMAX], x[NMAX][MMAX];
    double dmin_;
    double dstep, ydist;
    double dlevel;

    Integer ic[NMAX], ilc[NMAX-1], iord[NMAX], isx[MMAX],
           iuc[NMAX-1];
    Integer nsym;
    Integer i, j, k;
    Integer m, n;
    Integer int_method;
    Integer tdx=MMAX;

    char **c = 0;
    char name[NMAX][3];
    char char_dist[2];
    char char_scale[2];
    char char_update[2];

    Nag_ClusterMethod method;
    Nag_MatUpdate update;
    Nag_DistanceType dist;
    Nag_VarScaleType scale;

    Vprintf("g03ejc Example Program Results\n\n");

    /* Skip heading in data file */
    Vscanf("%*[^\n]");

    Vscanf("%ld",&n);
    Vscanf("%ld",&m);
    if (n <= NMAX && m <= MMAX)
    {
        Vscanf("%ld",&int_method);
        if (int_method == 1)
```

```

    method = Nag_SingleLink;
else if (int_method == 2)
    method = Nag_CompleteLink;
else if (int_method == 3)
    method = Nag_GroupAverage;
else if (int_method == 4)
    method = Nag_Centroid;
else if (int_method == 5)
    method = Nag_Median;
else
    method = Nag_MinVariance;

Vscanf("%s",char_update);
if (*char_update == 'U')
    update = Nag_MatUp;
else
    update = Nag_NoMatUp;

Vscanf("%s",char_dist);
if (*char_dist == 'A')
    dist = Nag_DistAbs;
else if (*char_dist == 'E')
    dist = Nag_DistEuclid;
else
    dist = Nag_DistSquared;

Vscanf("%s",char_scale);
if (*char_scale == 'S')
    scale = Nag_VarScaleStd;
else if (*char_scale == 'R')
    scale = Nag_VarScaleRange;
else if (*char_scale == 'G')
    scale = Nag_VarScaleUser;
else
    scale = Nag_NoVarScale;

for (j = 0; j < n; ++j)
{
    for (i = 0; i < m; ++i)
        Vscanf("%lf",&x[j][i]);
    Vscanf("%s",name[j]);
}
for (i = 0; i < m; ++i)
    Vscanf("%ld",&isx[i]);
for (i = 0; i < m; ++i)
    Vscanf("%lf",&s[i]);

Vscanf("%ld",&k);
Vscanf("%lf",&dlevel);

/* Compute the distance matrix */
g03eac(update, dist, scale, n, m, (double *)x, tdx, isx, s, d, NAGERR_DEFAULT);

/* Perform clustering */
g03ecc(method, n, d, ilc, iuc, cd, iord, dord, NAGERR_DEFAULT);

Vprintf("\nDistance    Clusters Joined\n\n");

for (i = 0; i < n-1; ++i)
{
    Vprintf("%10.3f      ",cd[i]);
    Vprintf("%3s",name[ilc[i]-1]);
    Vprintf("%3s",name[iuc[i]-1]);
    Vprintf("\n");
}
/* Produce dendrogram */
nsym = 20;
dmin_ = 0.0;
dstep = cd[n - 2] / (double) nsym;
g03ehc(Nag_DendSouth, n, dord, dmin_, dstep, nsym, &c, NAGERR_DEFAULT);

```

```

Vprintf("\n");
Vprintf("Dendrogram ");
Vprintf("\n");
Vprintf("\n");
ydist = cd[n - 2];
for (i = 0; i < nsym; ++i)
{
    if ((i+1) % 3 == 1)
    {
        Vprintf("%10.3f%6s",ydist,"");
        Vprintf("%s",c[i]);
        Vprintf("\n");
    }
    else
    {
        Vprintf("%16s%s","","", c[i]);
        Vprintf("\n");
    }
    ydist -= dstep;
}
Vprintf("\n");
Vprintf("%14s","");
for (i = 0; i < n; ++i)
{
    Vprintf("%3s",name[iord[i]-1]);
}
Vprintf("\n");
g03xzc(&c);
g03ejc(n, cd, iord, dord, &k, &dlevel, ic, NAGERR_DEFAULT);
Vprintf("\n%s%2ld%s\n", "Allocation to ",k," clusters");
Vprintf("Object Cluster\n\n");
for (i = 0; i < n; ++i)
{
    Vprintf("%5s%5s%5s","","",name[i],"");
    Vprintf("%ld      ",ic[i]);
    Vprintf("\n");
}
exit(EXIT_SUCCESS);
}
else
{
    Vprintf("Incorrect input value of n or m.\n");
    exit(EXIT_FAILURE);
}
}

```

## 8.2. Program Data

```

g03ejc Example Program Data
5 3
5
I S U
1 5.0 2.0 A
2 1.0 1.0 B
3 4.0 3.0 C
4 1.0 2.0 D
5 5.0 0.0 E
0 1 1
1.0 1.0 1.0
2 0.0

```

### 8.3. Program Results

g03ejc Example Program Results

Distance Clusters Joined

1.000	B	D
2.000	A	C
6.500	A	E
14.125	A	B

Dendrogram

14.125	-----	
	I      I	
	I      I	
12.006	I      I	
	I      I	
	I      I	
9.887	I      I	
	I      I	
	I      I	
7.769	I      I	
	----*	I
	I    I	I
5.650	I    I	I
	I    I	I
	I    I	I
3.531	I    I	I
	I    I	I
	----*	I      I
1.412	I    I	I    ---*
	I    I	I    I    I
	A    C    E    B    D	

Allocation to 2 clusters

Object Cluster

A	1
B	2
C	1
D	2
E	1

---