* Encoding: UTF-8.

TITLE RUN weighted Euclidean model on Rosenberg & Kim kinship sorting data.

>Warning # 2003. Command name: TITLE

>The title given exceeds 60 characters in length. The first 60 characters wil

>be used.

COMMENT the data have been converted to dissimilarities.

COMMENT the matrices below represent (females, males) multiple sort 1, multiple sort 2, single sort.

DATA LIST

/1 AUNT 1-3 BROTHR 4-6 COUSIN 7-9 DAUGHT 10-12 FATHER 13-15 GRDAUG 16-18 GRFA TH 19-21 GRMOTH 22-24 GRSON 25-27 MOTHER 28-30

NEPHEW 31-33 NIECE 34-36 SISTER 37-39 SON 40-42 UNCLE 43-45.

Data List will read 1 records from the command file

Variable	Rec	Start	End	Format
AUNT	1	1	3	F3.0
BROTHR	1	4	6	F3.0
COUSIN	1	7	9	F3.0
DAUGHT	1	10	12	F3.0
FATHER	1	13	15	F3.0
GRDAUG	1	16	18	F3.0
GRFATH	1	19	21	F3.0
GRMOTH	1	22	24	F3.0
GRSON	1	25	27	F3.0
MOTHER	1	28	30	F3.0
NEPHEW	1	31	33	F3.0
NIECE	1	34	36	F3.0
SISTER	1	37	39	F3.0
SON	1	40	42	F3.0
UNCLE	1	43	45	F3.0

BEGIN DATA

79

56 70

36 66 71

76 22 78 63

34 73 70 25 77

76 35 75 76 32 61

36 78 75 34 76 17 48

77 27 71 71 31 49 17 63

33 68 76 15 50 31 76 30 77

57 33 55 74 38 74 37 78 30 80

```
13 75 54 32 79 31 77 38 73 39 45
38 48 70 20 67 28 77 34 74 21 77 35
77 20 72 48 16 70 35 77 26 62 32 72 66
47 38 59 79 32 78 37 77 35 75 13 57 79 38
76
55 63
60 56 73
70 42 78 52
57 74 72 44 78
72 58 76 76 45 57
54 79 78 52 70 29 29
74 54 70 68 55 28 30 56
52 64 80 24 29 54 70 45 77
51 52 48 72 61 71 59 78 53 80
26 71 49 53 80 52 78 59 70 61 26
58 23 65 34 65 53 79 58 73 43 71 52
77 35 71 28 25 68 53 75 46 51 54 71 55
28 56 54 77 50 77 55 74 59 69 26 52 75 59
78
53 62
54 66 71
73 33 72 63
52 74 75 49 74
75 52 70 73 48 52
48 77 76 52 73 27 31
78 48 69 72 50 30 25 52
45 60 77 36 31 50 74 47 76
58 48 45 72 54 74 51 76 47 80
30 75 52 49 77 49 75 52 74 53 30
51 32 69 42 61 49 77 52 75 34 74 47
78 41 67 28 36 72 51 74 47 63 47 74 67
31 51 47 77 46 77 48 75 51 74 29 57 78 53
74
60 56
54 57 67
66 36 68 62
```

```
51 69 65 34 78
68 53 69 74 39 62
43 78 74 47 69 34 31
74 45 61 67 50 34 35 62
40 62 74 33 32 50 68 39 76
64 41 45 67 52 65 49 75 39 78
38 67 51 41 80 39 75 49 66 53 30
52 31 62 27 63 40 77 51 69 36 68 40
78 32 63 35 34 68 49 77 37 63 40 67 57
28 50 54 79 40 76 44 69 51 66 39 63 77 53
83
38 77
79 61 83
79 55 84 43
79 82 82 69 83
84 76 84 83 73 48
78 83 84 74 82 38 11
85 73 81 80 74 13 38 48
72 63 84 34 13 74 82 73 83
49 74 53 81 77 80 78 84 73 83
42 83 53 74 85 72 85 79 79 79 12
77 10 78 52 63 73 83 76 82 55 81 76
85 52 82 14 34 80 74 83 69 43 74 81 61
10 77 39 85 72 85 78 84 79 79 42 49 83 79
84
55 73
71 67 80
76 42 82 61
71 77 79 55 82
84 69 81 81 66 60
70 82 82 65 80 46 21
85 60 79 73 66 21 43 61
63 59 83 44 19 66 81 63 82
59 69 58 79 72 79 71 85 64 85
45 82 59 64 85 63 84 72 80 72 20
72 21 75 47 61 59 83 68 76 46 82 67
84 49 80 22 44 75 65 81 56 60 65 80 64
```

17 71 55 84 63 85 70 84 72 76 46 60 84 70 END DATA.

TITLE run PROXSCAL on Rosenberg and Kim kinship data.

PROXSCAL

VARIABLES= AUNT BROTHR COUSIN DAUGHT FATHER GRDAUG GRFATH GRMOTH GRSON MOTHE R NEPHEW NIECE SISTER SON UNCLE

/SHAPE=LOWER

/INITIAL=TORGERSON

/CONDITION=MATRIX

/TRANSFORMATION=INTERVAL

/PROXIMITIES=DISSIMILARITIES

/MODEL=WEIGHTED

/CRITERIA=DIMENSIONS(2,2) MAXITER(100) DIFFSTRESS(.0001) MINSTRESS(.0001)

/PRINT=COMMON WEIGHTS INPUT STRESS

/PLOT=COMMON WEIGHTS .

Proxscal

Credit

Proxscal
Version 1.0
by
Leiden SPSS Group
Leiden University
The Netherlands

Case Processing Summary

Cases		90
Sources		6
Objects		15
Proximities	Total Proximities	630 ^a
	Missing Proximities	0
	Active Proximities ^b	630

- a. Sum over sources of all strictly lower-triangular proximities.
- b. Active proximities include all non-missing proximities.

Input Data

Proximities

Source: SRC_1

	AUNT	BROTHR	COUSIN	DAUGHT	FATHER	GRDAUG	GRFATH
AUNT							
BROTHR	79.000						
COUSIN	56.000	70.000					
DAUGHT	36.000	66.000	71.000				
FATHER	76.000	22.000	78.000	63.000			
GRDAUG	34.000	73.000	70.000	25.000	77.000		
GRFATH	76.000	35.000	75.000	76.000	32.000	61.000	
GRMOTH	36.000	78.000	75.000	34.000	76.000	17.000	48.000
GRSON	77.000	27.000	71.000	71.000	31.000	49.000	17.000
MOTHER	33.000	68.000	76.000	15.000	50.000	31.000	76.000
NEPHEW	57.000	33.000	55.000	74.000	38.000	74.000	37.000
NIECE	13.000	75.000	54.000	32.000	79.000	31.000	77.000
SISTER	38.000	48.000	70.000	20.000	67.000	28.000	77.000
SON	77.000	20.000	72.000	48.000	16.000	70.000	35.000
UNCLE	47.000	38.000	59.000	79.000	32.000	78.000	37.000

Proximities

Source: SRC_1

	GRMOTH	GRSON	MOTHER	NEPHEW	NIECE	SISTER	SON
AUNT							
BROTHR							
COUSIN							
DAUGHT							
FATHER							
GRDAUG							
GRFATH							
GRMOTH							
GRSON	63.000						
MOTHER	30.000	77.000					
NEPHEW	78.000	30.000	80.000				
NIECE	38.000	73.000	39.000	45.000			
SISTER	34.000	74.000	21.000	77.000	35.000		
SON	77.000	26.000	62.000	32.000	72.000	66.000	
UNCLE	77.000	35.000	75.000	13.000	57.000	79.000	38.000

Proximities

Source: SRC_1

Course. O	
	UNCLE
AUNT	
BROTHR	
COUSIN	
DAUGHT	
FATHER	
GRDAUG	
GRFATH	
GRMOTH	
GRSON	
MOTHER	
NEPHEW	
NIECE	
SISTER	
SON	
UNCLE	

Goodness of Fit

Stress and Fit Measures

Normalized Raw Stress	.07643
Stress-I	.27646 ^a
Stress-II	.71501 ^a
S-Stress	.16184 ^b
Dispersion Accounted For (D.A.F.)	.92357
Tucker's Coefficient of Congruence	.96103

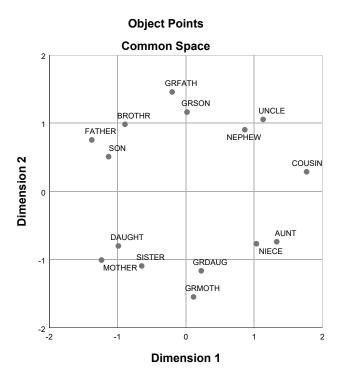
PROXSCAL minimizes Normalized Raw Stress.

- a. Optimal scaling factor = 1.083.
- b. Optimal scaling factor = .957.

Common Space

Final Coordinates

	Dimension		
	1	2	
AUNT	1.331	737	
BROTHR	893	.983	
COUSIN	1.768	.287	
DAUGHT	990	800	
FATHER	-1.379	.754	
GRDAUG	.225	-1.163	
GRFATH	201	1.457	
GRMOTH	.112	-1.546	
GRSON	.016	1.162	
MOTHER	-1.236	-1.006	
NEPHEW	.863	.904	
NIECE	1.033	767	
SISTER	648	-1.092	
SON	-1.133	.509	
UNCLE	1.131	1.055	



Individual Spaces

Dimension Weights

	Dimension		
Source	1	2	
SRC_1	.391	.543	
SRC_2	.489	.441	
SRC_3	.460	.472	
SRC_4	.439	.491	
SRC_5	.524	.377	
SRC_6	.507	.405	

