

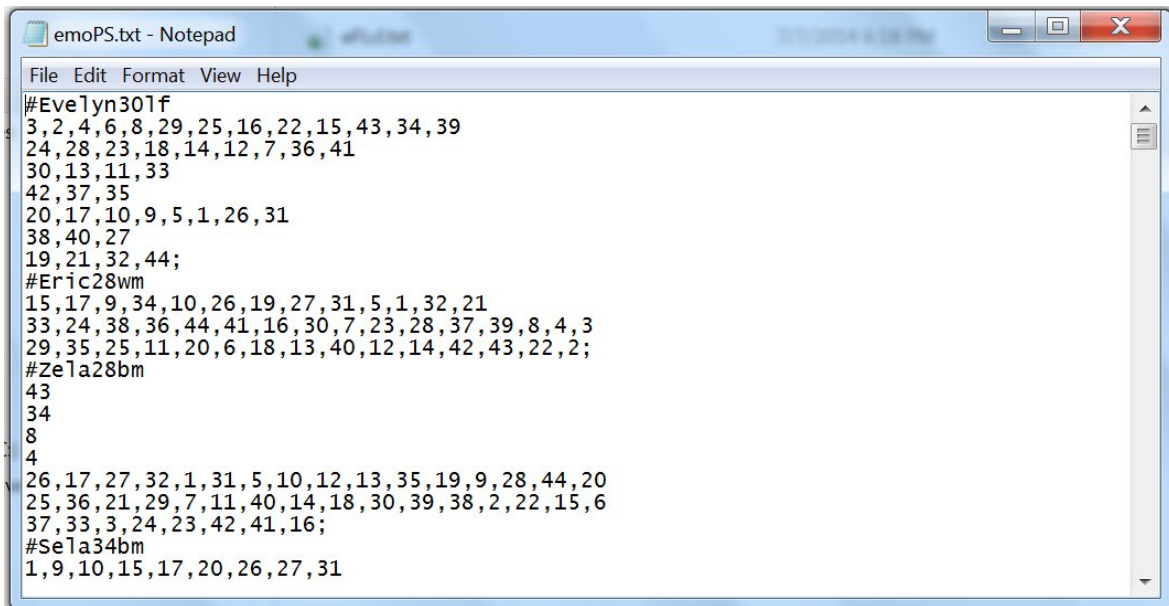
## Pile Sorts

Data file: EMOPS.TXT

Label file: EMOLAB.TXT

Respondent label: ERSPLAB.TXT

Input file format:



```
emoPS.txt - Notepad
File Edit Format View Help
#Evelyn301f
3,2,4,6,8,29,25,16,22,15,43,34,39
24,28,23,18,14,12,7,36,41
30,13,11,33
42,37,35
20,17,10,9,5,1,26,31
38,40,27
19,21,32,44;
#Eric28wm
15,17,9,34,10,26,19,27,31,5,1,32,21
33,24,38,36,44,41,16,30,7,23,28,37,39,8,4,3
29,35,25,11,20,6,18,13,40,12,14,42,43,22,2;
#Zeia28bm
43
34
8
4
26,17,27,32,1,31,5,10,12,13,35,19,9,28,44,20
25,36,21,29,7,11,40,14,18,30,39,38,2,22,15,6
37,33,3,24,23,42,41,16;
#Seia34bm
1,9,10,15,17,20,26,27,31
```

Import pile sort data

DATA → IMPORT → PILESORT

Input data file: EMOPS.TXT

Number of items: 44

Number of respondents: 40

Item labels: EMOLAB.TXT

Respondent labels: ERSPLAB.TXT

Some items more often: NO

Missing items: NO

Individual proximities output: EMOIPX (individual proximity matrices, for each respondent)

Aggregate proximities output: EMOAPX (aggregate proximity matrix)

Agreement dataset: PSCORR

MDS of aggregate proximity matrix

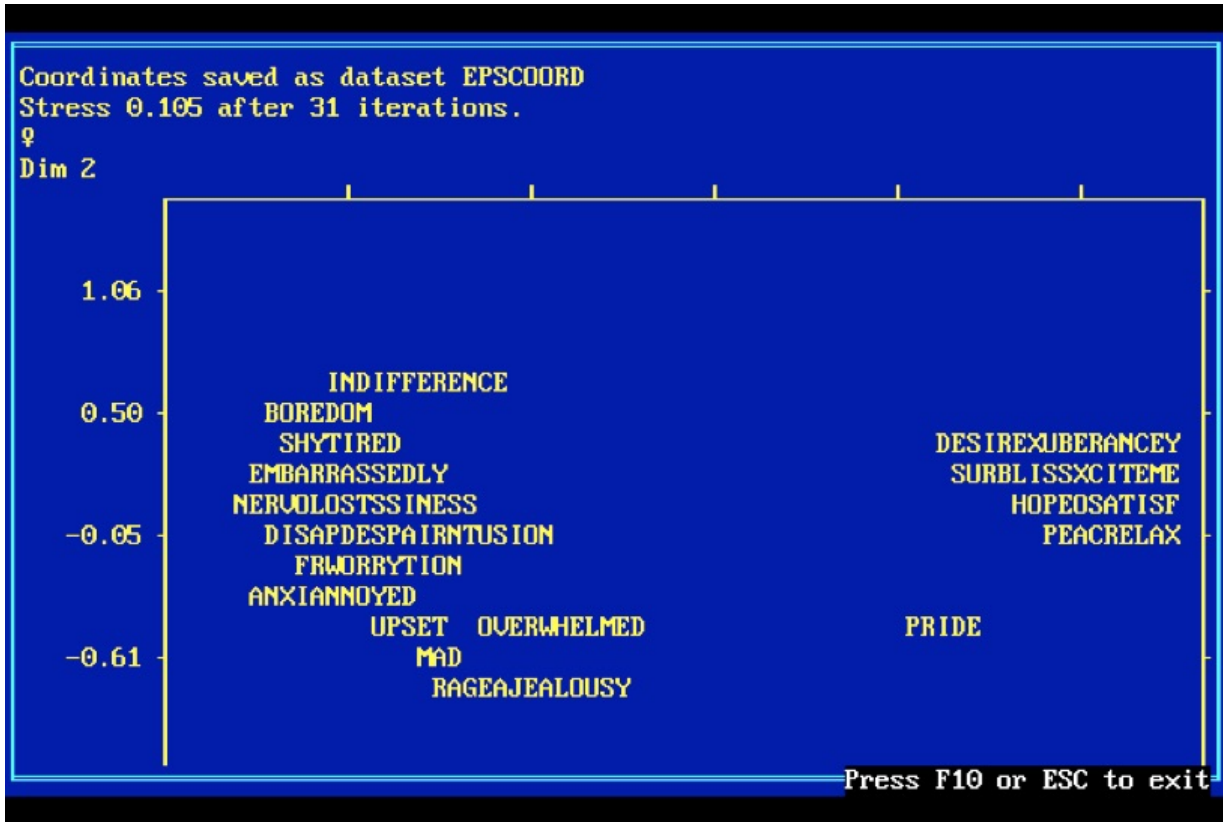
TOOLS → SCALING → NONMETRIC MDS

Dataset: EMOAPX

Dimensions: 2

Similarities

Output: EPSCOORD



To export the coordinates for use in another application:

DATA → EXPORT → EPSCOORD (ends up as EPSCOORD.DAT)