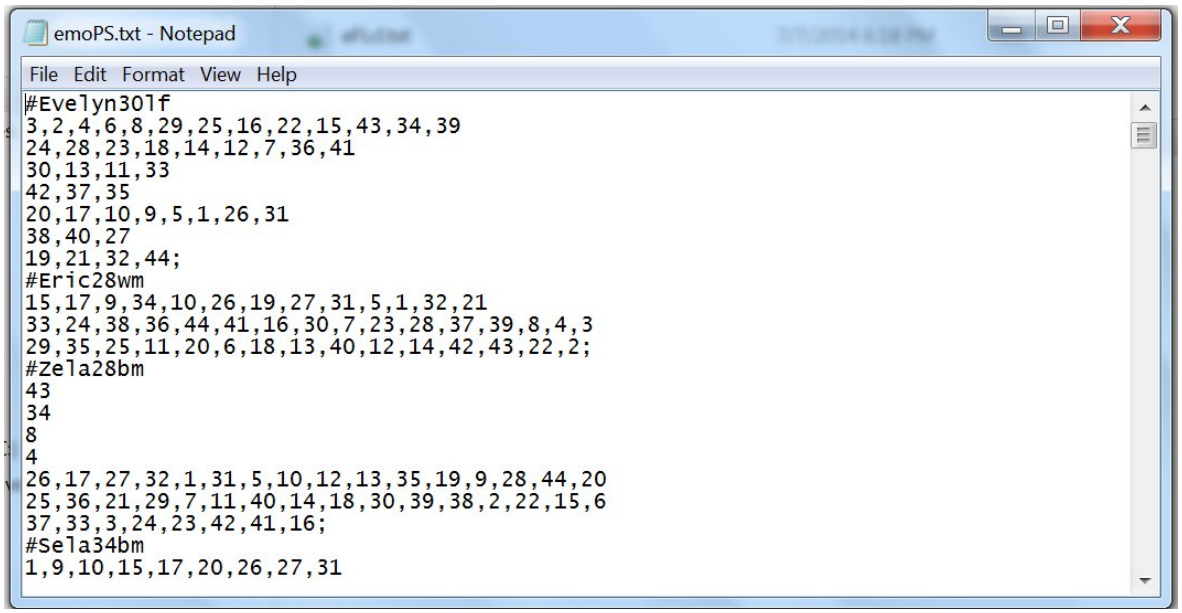
Pile Sorts

Data file: EMOPS.TXT
Label file: EMOLAB.TXT
Respondent label: ERSPLAB.TXT

Input file format:



Import pile sort data

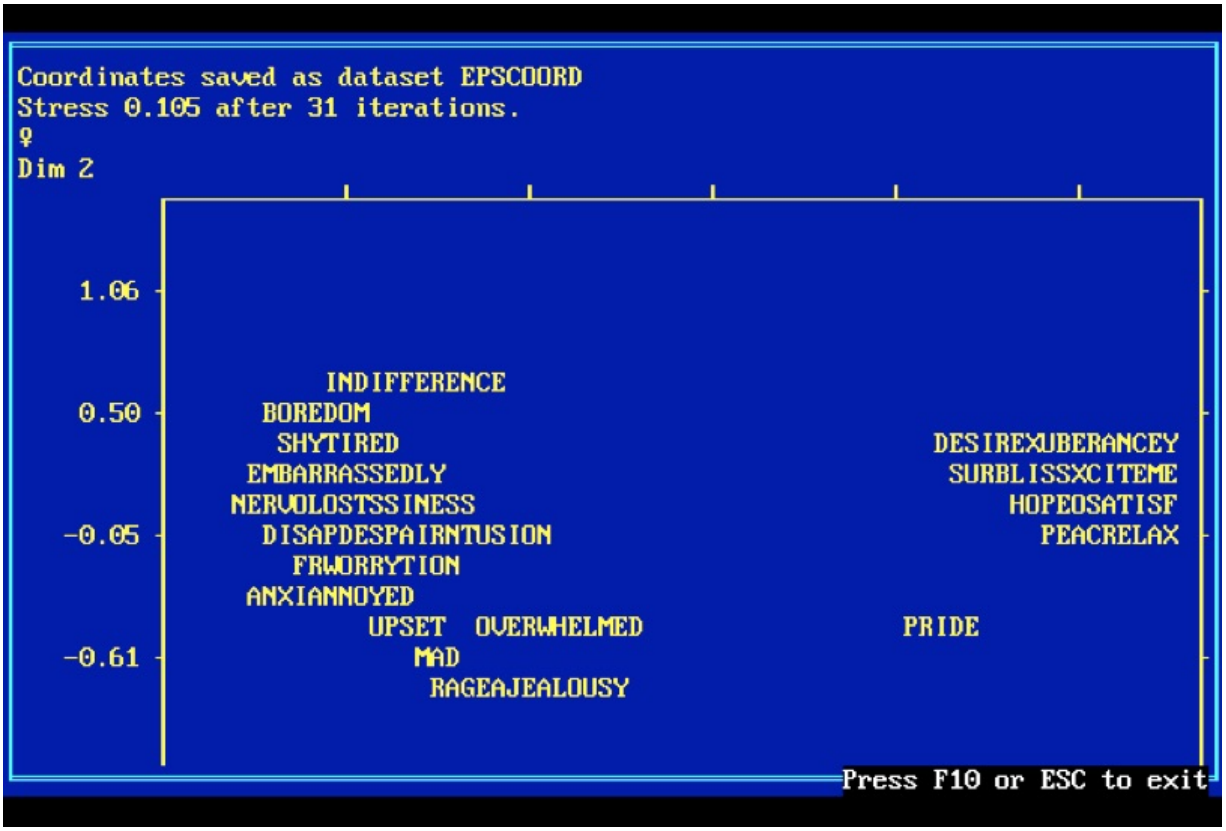
DATA → IMPORT → PILESORT

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)