

MULTIDIMENSIONAL SCALING:

Using SPSS/PROXSCAL

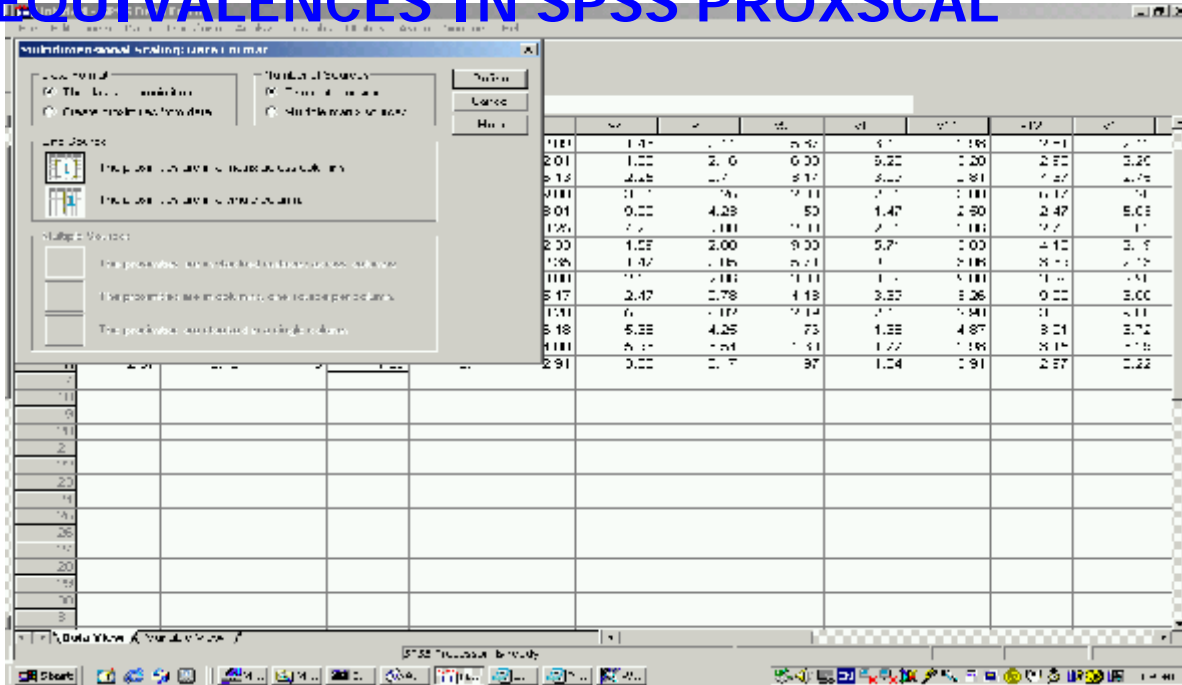
- SPSS 10 offers **PROXSCAL** (PROXimity SCALing) as an alternative to ALSCAL for multidimensional scaling: **USE IT !!** ALSCAL has been shown to be sub-optimal (Ramsay).
- PROXSCAL performs most Distance Model scaling (for scalar products/vector models, see SPSS Categories). (*Pre-SPSS PROXSCAL .pdf Documentation by Busing is available*).
- Data for basic MDS in SPSS10 can be either
 1. input *directly* as a full SSM (*square symmetric matrix*) of proximities=dis/similarities into SPSS editor.
 2. *calculate measure within SPSS* from a raw datafile (separate procedure)
 3. OR (in PROXSCAL only) *matrix can be imported as a SSM. **SPSS will not accept LT matrices directly***, (n.b. universal use of LT matrices in other programs; confusing SPSS documentation suggesting otherwise).
 4. procedures for changing a LT into a SSM for SPSS are contorted but possible

(subject of a separate handout).

SETTING UP

data-transformation-model

EQUIVALENCES IN SPSS PROXSCAL



- n.b. Use Right Click for supplementary information in SPSS10
- PROXSCAL "proximities" = dis/similarities
- PROXSCAL "Transformed proximities" = disparities.

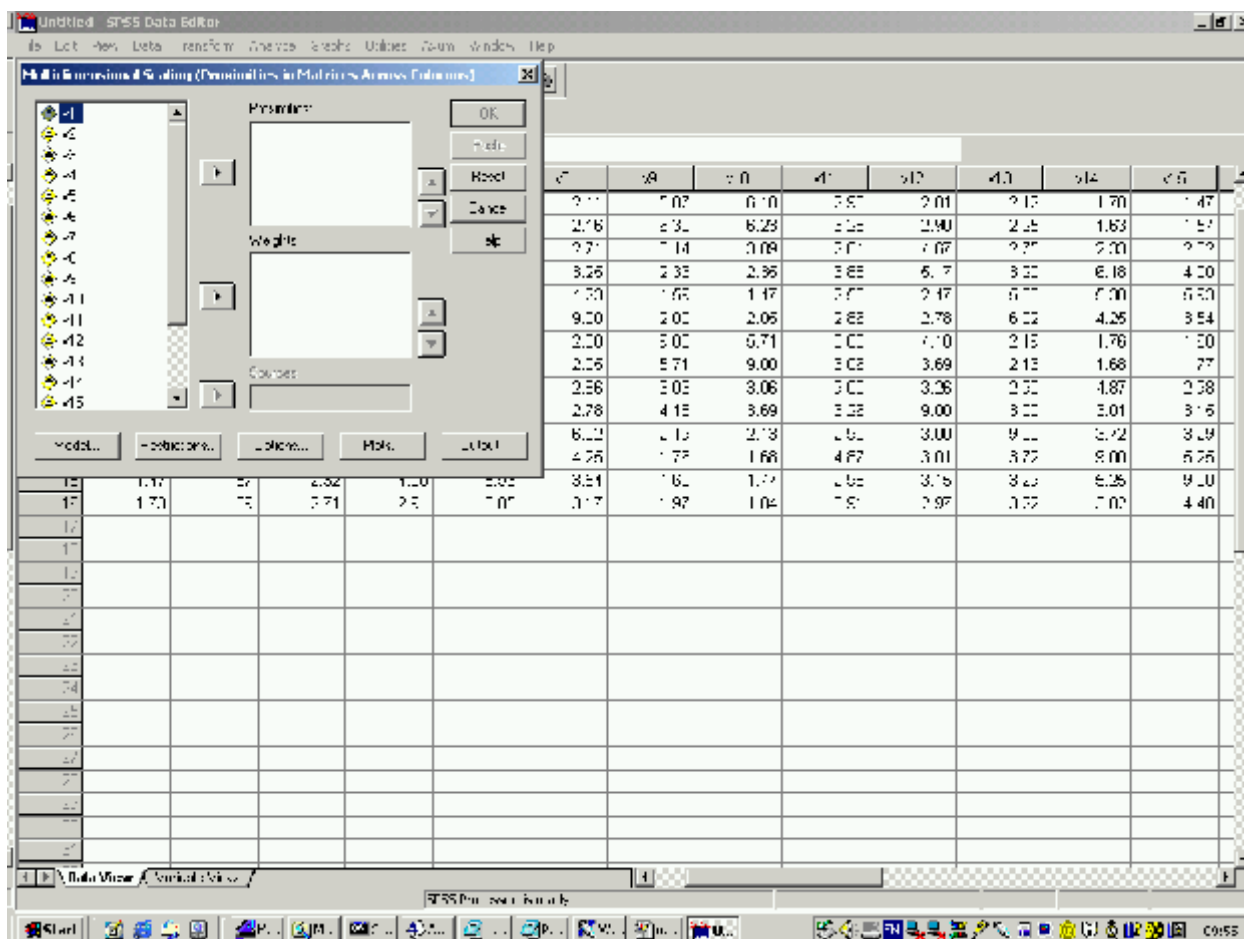
1. DATA

Analyze Y Scale Y Multidimensional scaling (PROXSCAL)

First Window (Data Format)

- Data Format: (T data are proximities (or create from raw data – initiates "Create proximities from data" procedure)
- Number of sources: (T One for 2W1M data; Multiple for INDSCAL etc)

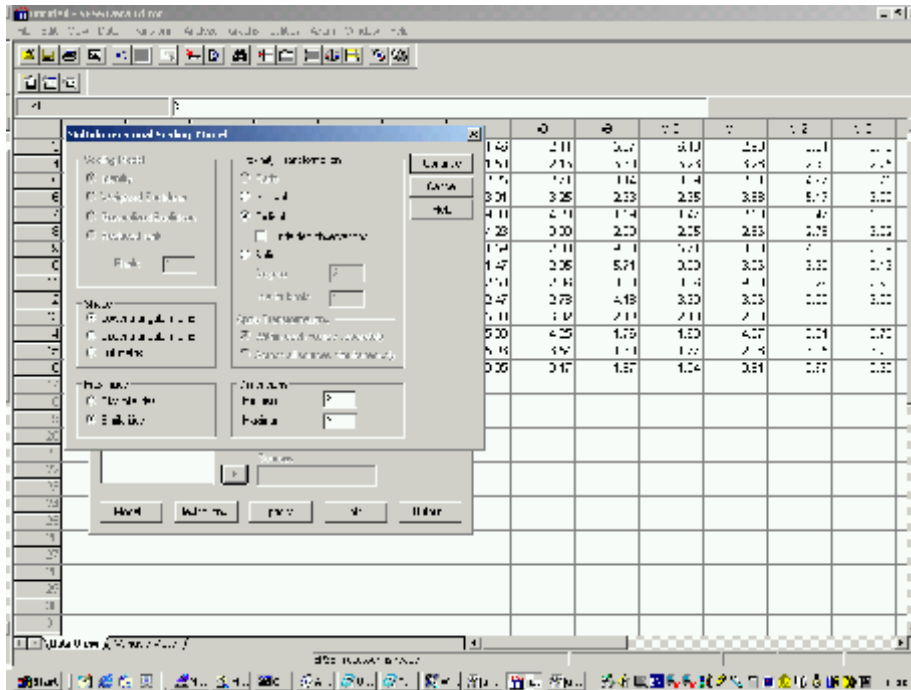
Second Window:



- Transfer Matrix (v1 - v16) across.
(Best to keep variables name short at this stage, to avoid over-printing at graphing stage; they can be selectively lengthened later in plotting routine)
- Note the crucial buttons at the foot of the Transfer Window:

MODEL- RESTRICTIONS-OPTIONS-PLOTS-OUTPUTS

these are the ones which set the detail of the analysis and run parameters.

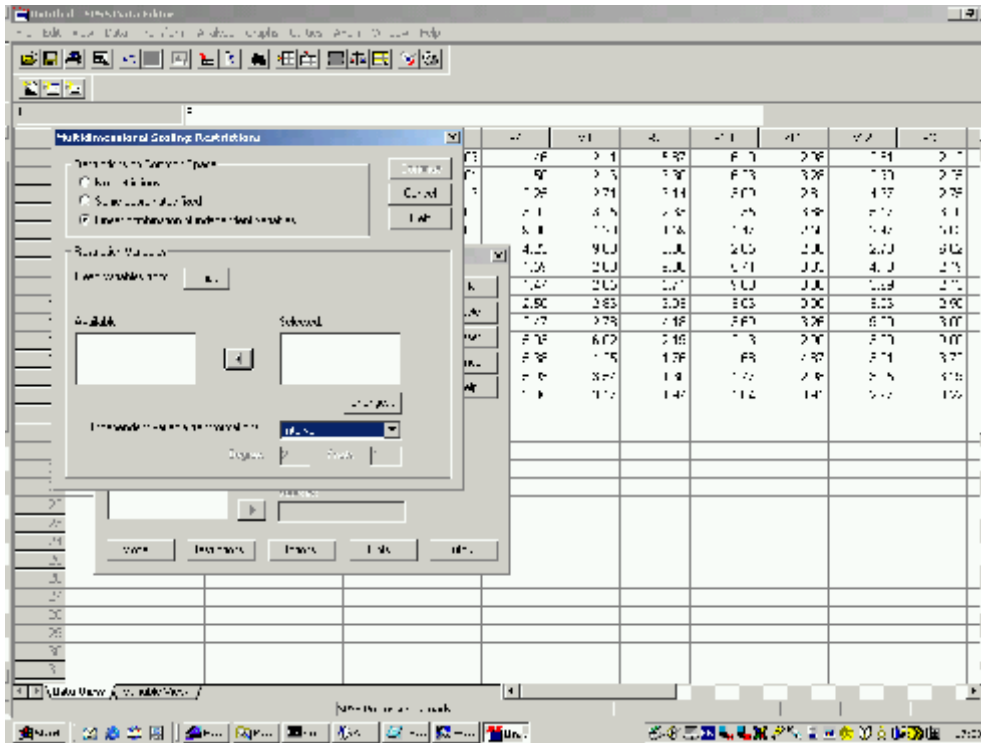


- **SCALING MODEL:**
 - < Identity means simple Euclidean
 - < Weighted Euclidean means INDSCAL
 - < Generalised Euclidean means IDIOSCAL (individual rotation and then weighting)
 - < Reduced Rank means IDIOSCAL with minimal rank of matrix
- **SHAPE:** *does not refer to input matrix ...which must be SSM*
 - < Lower triangular means "only the lower triangular data are analysed" i.e. symmetric
 - < Upper triangular: ditto, upper
 - < Full matrix: data may be asymmetric, but are symmetrised
- **PROXIMITIES (data)**
 - < Similarity data (hi means more similar)
 - < Dissimilarity (hi means more dissimilar)

n.b. this option is default: beware!
- **DIMENSIONS**
MDS solutions proceed from max (-1) min

PROXSCAL MODEL PARAMETERS (cont.)

- **Proximity TRANSFORMATIONS**
 - < Ratio (LoM) – implies metric analysis
 - < **Interval** (LoM) – metric
 - < **Ordinal** (LoM) – non-metric. Default is secondary approach to ties, unless ...
 - T Untie tied values (=primary)
 - < **Spline** (cf Ramsay & MULTISCALE): piece-wise polynomial transformation of the original data. “Pieces” and shape of transformation are specified by:
 - T degree (1=linear; 2= quadratic ...), and
 - T Number of internal knots.
- **APPLY TRANSFORMATIONS** (applies only to INDSCAL and higher models; in effect local versus global application).



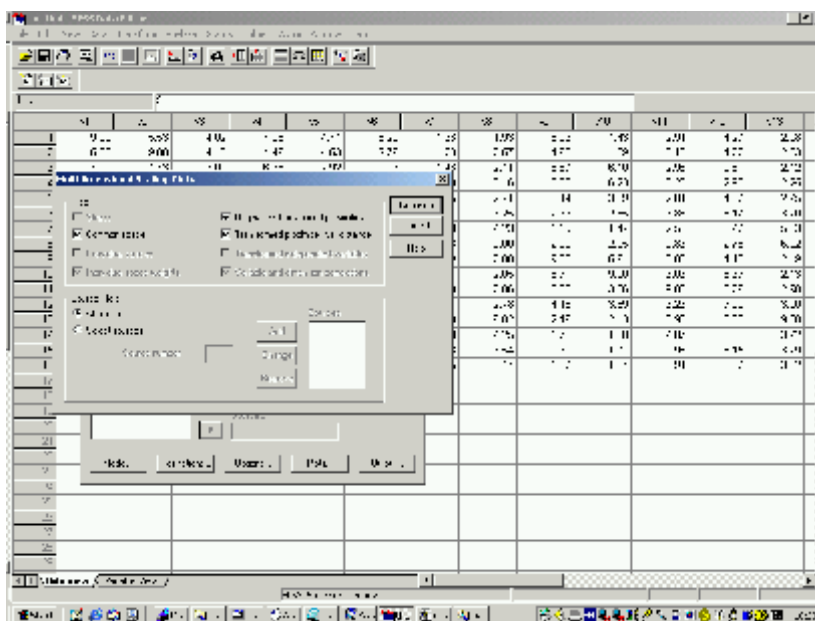
**RESTRICTIONS = EXTERNAL or
CONSTRAINED / CONFIRMATORY
ANALYSIS** (cf Borg and Groenen 1997, pp181-199).

These options allow for:

- Fixing some (known?) points in a configuration, and estimating the others (**T** Some co-ordinates fixed)
- Fitting (regressing) external properties (PRO-FIT) (**T** Linear combinations ...)

Additional information is contained in an separate (or integral) SPSS file

PLOTS

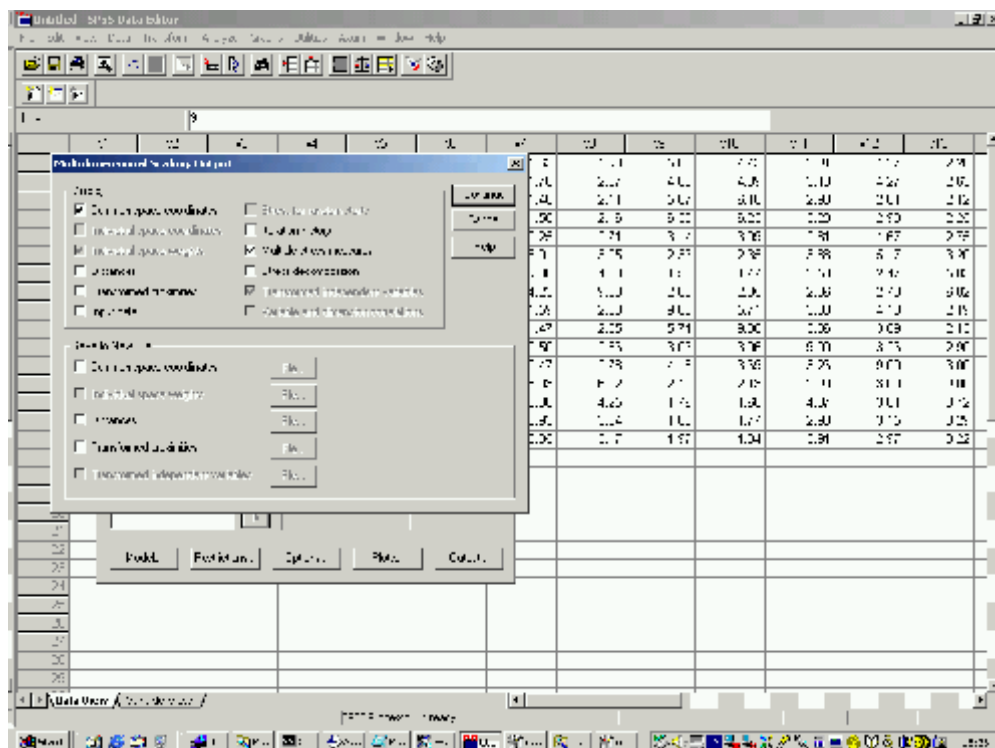


- “Common Space” = Group or Stimulus Configuration.

n.b. The Shepard Diagram is not immediately available in PROXSCAL; it combines (and can in principle be reconstructed from)

- original vs transformed proximities = disparities (d^* vs \hat{d} – Monotonic Fit)
- transformed proximities vs distances of solution (\hat{d} vs d – OLS fit)

OUTPUT



- **DISPLAY**
 - < Common Space (=Stimulus Configuration)
 - < Distances (of solution)
 - < Transformed proximities (=disparities)
 - < Input data (ALWAYS recommended, to ensure the program is working on the data YOU think it is ...)
 - < Iteration history (for diagnosis of stress minimization)
 - < Multiple stress measures (use Stress1 for comparison with other solutions; not S-STRESS. Note: *normalised* raw stress ... raw stress)
 - < Stress decomposition = point contribution to stress
 - < "Save to new file" is equivalent of MDSX's "PUNCH". Useful for graphic output.