

# Statistics for the Sciences

## Multidimensional Scaling

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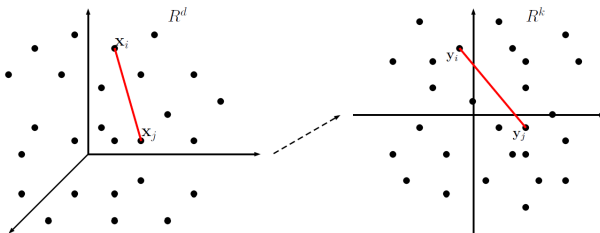
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# Outline

- Purpose of multidimensional scaling
- Multidimensional Scaling (MDS)
  - ▶ metric MDS
  - ▶ non-metric MDS
- Example
- Lab

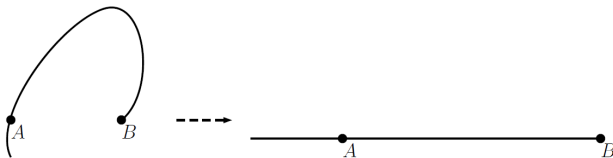
# Purpose of multidimensional scaling

- Multidimensional scaling (MDS) is another technique on dimensionality reduction.
- Unlike PCA, MDS maps the original high dimensional space to a lower dimensional space, but does so in an attempt to **preserve pairwise distances**.
  - ▶ The space is usually two-dimensional, sometimes may be three-dimensional.



# Purpose of multidimensional scaling

- The purpose of MDS is often just visualizing the data so it becomes easier for the user to explore and to understand their **structure**.
- The MDS algorithm starts with a distance matrix and uses it to reduce the data to two dimensions while preserving the pairwise distances.
  - ▶ PCA starts with the correlation matrix of data.



# Multidimensional Scaling

- MDS methods include
  - ▶ Classical MDS (Metric MDS) or Principal Coordinates Analysis (PCoA),
  - ▶ Non-metric MDS which focuses on preserving the rank order of the dissimilarities rather than the actual distances.

# Multidimensional Scaling

- Given  $p$ -dimensional data, how do we construct a **distance** matrix?

$$\mathbf{X} = \begin{pmatrix} x_{11} & x_{12} & \cdots & x_{1p} \\ x_{21} & x_{22} & \cdots & x_{2p} \\ \vdots & \vdots & \vdots & \vdots \\ x_{n1} & x_{n2} & \cdots & x_{np} \end{pmatrix}$$

# Multidimensional Scaling

- **Distance, dissimilarity** and **similarity** (or **proximity**) are defined for any pair of objects in any space.
  - ▶ Each objects or subject is just a row (record) of observations.
- Let  $x, y, z$  be objects. In mathematics, a distance function (that gives a distance between two objects) is also called **metric**, satisfying
  - ▶  $d(x, y) \geq 0$
  - ▶  $d(x, y) = 0$  if and only if  $x = y$
  - ▶  $d(x, y) = d(y, x)$
  - ▶  $d(x, z) \leq d(x, y) + d(y, z)$
- Given a set of dissimilarities, one can ask whether these values are distances.

# Multidimensional Scaling

- Dissimilarity Matrix:  $n \times n$  with the  $ij$ -th element  $d_{ij}$  measuring the distance (similarity or proximity) between the  $i$ th and the  $j$ th objects (or observations).  $D$  is symmetric.
  - ▶ Here  $p$  is the dimension of the data or the number of variables.
- Dissimilarity between points  $i$  and  $i'$ :  $\mathbf{x}_i$  and  $\mathbf{x}_{i'}$  ( $i, i' = 1, \dots, n$ ):

$$d(\mathbf{x}_i, \mathbf{x}_{i'}) = \sum_{j=1}^p d_j(x_{ij}, x_{i'j}).$$



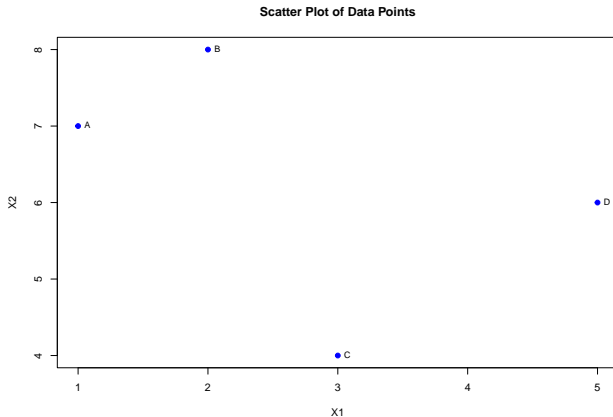
# Multidimensional Scaling

- An example

##		x1	x2
##	A	1	7
##	B	2	8
##	C	3	4
##	D	5	6

# Multidimensional Scaling

- A scatter plot of these data



# Multidimensional Scaling

- Construct a dissimilarity matrix

```
## [1] "Distance matrix of the data:"
```

```
##           A           B           C           D
## A 0.000000 1.414214 3.605551 4.123106
## B 1.414214 0.000000 4.123106 3.605551
## C 3.605551 4.123106 0.000000 2.828427
## D 4.123106 3.605551 2.828427 0.000000
```

# Multidimensional Scaling

- There are several types of distances. Consider two points  $A(x_1, y_1)$  and  $B(x_2, y_2)$

- ▶ Euclidean distance  $d(A, B) = \sqrt{(x_1 - x_2)^2 + (y_1 - y_2)^2}$
- ▶ Maximum Distance  $d(A, B) = \max(|x_1 - x_2|, |y_1 - y_2|)$
- ▶ Manhattan Distance  $d(A, B) = |x_1 - x_2| + |y_1 - y_2|$
- ▶ Bray Distance  $d(A, B) = \frac{|x_1 - x_2|}{|x_1| + |x_2|} + \frac{|y_1 - y_2|}{|y_1| + |y_2|}$
- ▶ Minkowski Distance  $d(A, B) = [(x_1 - x_2)^p + (y_1 - y_2)^p]^{1/p}, p > 0$
- ▶ Binary Distance (Dissimilarity for binary data)  
 $d(A, B) = (x_1 \text{ XOR } x_2) + (y_1 \text{ XOR } y_2)$

# Multidimensional Scaling

## The MDS problem:

- Assume a collection of  $p$ -dimensional  $n$  objects with Dissimilarity Matrix  $\mathbf{L} = \{l_{ij}\}, i, j = 1, 2, \dots, n$ .
- Our objective is to represent them as points in 2-d Euclidean space,  $\mathbf{y}_1, \dots, \mathbf{y}_n \in \mathbb{R}^2$ , such that

$$d_{ij} = d_{\mathbf{y}_i, \mathbf{y}_j} = l_{ij} \text{ or as close as possible for any } i, j.$$

- MDS
  - ▶ Called **metric MDS** if dissimilarities  $l_{ij}$  are quantitative.
  - ▶ Called **non-metric MDS** if dissimilarities  $l_{ij}$  are qualitative (e.g. ordinal).

# Multidimensional Scaling

- The mathematical tool for MDS often involves eigen decomposition of the Dissimilarity matrix to transform the distance information into a lower-dimensional space while preserving the distances between data points.
- Solution to MDS is not unique.
  - ▶ If  $\mathbf{Y}$  is a solution, then  $\mathbf{Y}^* = \mathbf{Y} + \mathbf{c}$ ,  $\mathbf{c} \in \mathbb{R}^2$  is also a solution.
  - ▶  $\mathbf{c}$  is just a location transformation.

# Non-metric MDS

- In many applications of MDS, dissimilarities are known only by their **rank order**, and the spacing between successively ranked dissimilarities is of no interest or is unavailable.
  - ▶ Non-metric MDS focuses on preserving the rank order of the dissimilarities.
- Non-metric MDS: non-metric MDS seeks to find an optimal configuration  $\mathbf{Y} \subset \mathbb{R}^2$  such that

$$f(l_{ij}) \approx d_{\mathbf{y}_i, \mathbf{y}_j} = d_{ij} \text{ for any } i, j.$$

where  $f$  is a general monotonic function implicitly defined by the rank order of the original dissimilarities  $l_{ij}$ .

- ▶  $f(l_{ij})$  generates **disparities**, which only preserve the order of the original dissimilarities  $l_{ij}$ . That is

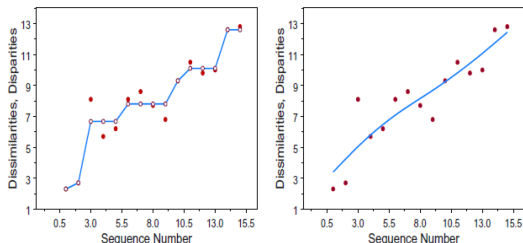
$$l_{ij} < l_{i'j'} \Leftrightarrow f(l_{ij}) < f(l_{i'j'})$$

# Non-metric MDS

- Kruskal's non-metric MDS minimizes the stress-1

$$\text{stress-1}(f(l_{ij}), d_{ij}) = \left( \frac{\sum_{i < j} (f(l_{ij}) - d_{ij})^2}{\sum d_{ij}^2} \right)^{1/2}$$

- It quantifies the difference between the disparities  $f(l_{ij})$  and the distances  $d_{ij}$  in  $\mathbb{R}^2$ .
- The function  $f$  works as if it were a regression curve
  - ▶ approximated dissimilarities  $d_{ij}$  treated as response,
  - ▶ disparities  $f(l_{ij})$  serves as estimate average of the response, and
  - ▶ the order of dissimilarities as explanatory
  - ▶ In the following figure, Left panel: Isotonic regression. right panel: Monotone spline. For the red points, the vertical axis is the dissimilarity  $l_{ij}$ , whereas for the fitted blue points, the vertical axis is the disparity  $f(l_{ij})$ .





# Non-metric MDS

- Process of Non-metric MDS

- ▶ Initialization: Start with an initial configuration of points in the lower-dimensional space like  $\mathbb{R}^2$ .
- ▶ Compute Distances: Calculate the Euclidean distances  $d_{ij}$  between points in the current configuration.
- ▶ Monotonic Regression: Find the disparities  $f(l_{ij})$  that best match the distances  $d_{ij}$  while preserving the rank order of  $l_{ij}$ .
- ▶ Optimization: Adjust the configuration of points (the coordinates of  $\mathbf{y}$ ) to minimize the stress function in the last slide. The goal is to find the configuration where the distances  $d_{ij}$  are as close as possible to the disparities  $f(l_{ij})$ .

## Example 1: Metric MDS

- `lemminvert.csv`: Lemmens et al. (2015) did a detailed study of various biotic communities in artificial ponds in Belgium. They sampled 28 ponds that represented different types of management, a combination of fish farming strategies (no fish, farming young fish, low intensity management, no management), and drainage frequencies ( $> 10$  years ago, occasional, annual). They also quantified taxon abundances for fish, zooplankton, and macro-invertebrates (different families and species within some groups) and covers of submerged, floating, and emergent vegetation. The macroinvertebrate dataset only included 23 ponds.
- See analysis with R here:  
<https://mjkeough.github.io/examples/lemmpcoa.nb.html>

```
## [1] "1J1" "1J2" "1J3" "1J5" "1J6" "1J7" "BK1" "BK2" "BK5" "BK6"  
## [13] "02" "03" "04" "05" "06" "07" "V1" "V2" "V3" "V4"
```

# Example 1: Metric MDS

- The first 3 variables are pond classification variables

```
## 'data.frame': 23 obs. of 32 variables:
## $ site : Factor w/ 23 levels "1J1","1J2","1J3",...: 1 2 3 4 5 6 7 8 9 10 ...
## $ manag : Factor w/ 4 levels "li","nf","nm",...: 4 4 4 4 4 4 2 2 2 2 ...
## $ managsymb: int 0 0 0 0 0 0 1 1 1 1 ...
## $ caen : int 0 6 0 5 0 3 31 36 0 16 ...
## $ baet : int 158 13 20 98 3 605 1571 494 73 90 ...
## $ acrol : int 0 0 0 0 0 0 0 0 1 0 ...
## $ lymn : int 3 0 3 10 9 7 763 109 46 15 ...
## $ plan : int 5 50 16 206 14 19 1164 1035 64 421 ...
## $ sphaer : int 0 0 0 2 0 0 80 147 0 0 ...
## $ vivip : int 0 0 0 0 0 0 0 0 0 0 ...
## $ corix : int 671 95 8 98 23 9 730 389 239 25 ...
## $ nauc : int 0 0 0 0 0 0 0 2 2 0 ...
## $ nepid : int 0 0 0 0 0 0 0 0 0 0 ...
## $ noto : int 0 0 0 0 0 0 3 1 0 0 ...
## $ pleid : int 0 0 0 0 0 0 20 141 5 0 ...
## $ gamm : int 0 0 1 2 1 0 4 0 4 1 ...
## $ asell : int 0 0 0 4 0 0 0 0 0 1 ...
## $ cerat : int 4 7 9 5 0 7 88 3 6 20 ...
## $ chaob : int 23 37 60 20 3 5 68 0 204 22 ...
## $ chiro : int 222 322 240 26 60 394 1236 62 569 69 ...
## $ culic : int 19 17 12 4 3 19 12 0 24 21 ...
## $ cylind : int 0 0 0 0 0 0 0 0 0 0 ...
## $ dixid : int 40 52 30 2 5 0 4 6 47 38 ...
## $ empid : int 0 0 0 0 1 0 0 0 0 5 ...
## $ ephyd : int 0 0 0 0 1 0 0 0 0 0 ...
## $ limon : int 8 1 25 9 2 0 8 0 0 10 ...
## $ parob : int 0 0 0 0 1 0 0 0 0 0 ...
```

# Example 1: Metric MDS

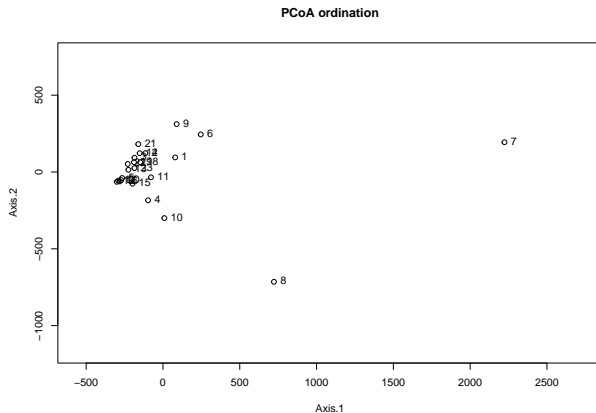
## ● PCoA Analysis

```
## $correction
## [1] "none" "1"
##
## $note
## [1] "There were no negative eigenvalues. No correction was applied"
##
## $values
##      Eigenvalues Relative_eig Broken_stick Cumul_eig Cumul_br_stick
## 1  6.284770e+06  7.820467e-01  0.167764239  0.7820467    0.1677642
## 2  9.438104e+05  1.174433e-01  0.122309693  0.8994900    0.2900739
## 3  4.249575e+05  5.287968e-02  0.099582420  0.9523696    0.3896564
## 4  2.622459e+05  3.263263e-02  0.084430905  0.9850023    0.4740873
## 5  7.088309e+04  8.820352e-03  0.073067269  0.9938226    0.5471545
## 6  2.045955e+04  2.545889e-03  0.063976360  0.9963685    0.6111309
## 7  1.124622e+04  1.399426e-03  0.056400602  0.9977679    0.6675315
## 8  5.798579e+03  7.215475e-04  0.049907096  0.9984895    0.7174386
## 9  5.194204e+03  6.463420e-04  0.044225278  0.9991358    0.7616639
## 10 2.820873e+03  3.510159e-04  0.039174773  0.9994868    0.8008386
## 11 1.844753e+03  2.295523e-04  0.034629318  0.9997164    0.8354680
## 12 1.287003e+03  1.601485e-04  0.030497087  0.9998765    0.8659650
## 13 5.023922e+02  6.251528e-05  0.026709208  0.9999391    0.8926742
## 14 2.379086e+02  2.960421e-05  0.023212704  0.9999687    0.9158870
## 15 1.646356e+02  2.048647e-05  0.019965951  0.9999891    0.9358529
## 16 4.038504e+01  5.025321e-06  0.016935648  0.9999942    0.9527886
## 17 2.848683e+01  3.544765e-06  0.014094739  0.9999977    0.9668833
## 18 8.453043e+00  1.051856e-06  0.011420942  0.9999988    0.9783042
## 19 4.215181e+00  5.245170e-07  0.008895690  0.9999993    0.9871999
## 20 3.346910e+00  4.164734e-07  0.006503345  0.9999997    0.9937033
## 21 1.664062e+00  2.070680e-07  0.004230618  0.9999999    0.9979339
## 22 6.604191e-01  8.217940e-08  0.002066116  1.0000000    1.0000000
##
## $vectors
##      Axis.1      Axis.2      Axis.3      Axis.4      Axis.5      Axis.6
## 1   79.740869   95.10472  498.99343  215.307907 -38.041790 -21.18016513
## 2  -114.437233  122.79086   36.06851 -115.990258 -23.898179 -39.23757341
## 3  -185.375472   92.69808  -47.57754  -84.558400 -12.730992  2.97584395
## 4  -96.499521 -184.16382 -11.40450  17.971440  8.590388 -0.03897887
```

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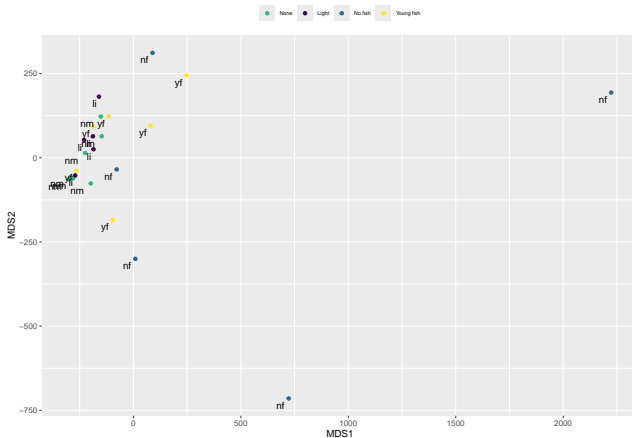
# Example 1: Metric MDS

- biplot



# Example 1: Metric MDS

- biplot



## Example 2: Non-Metric MDS

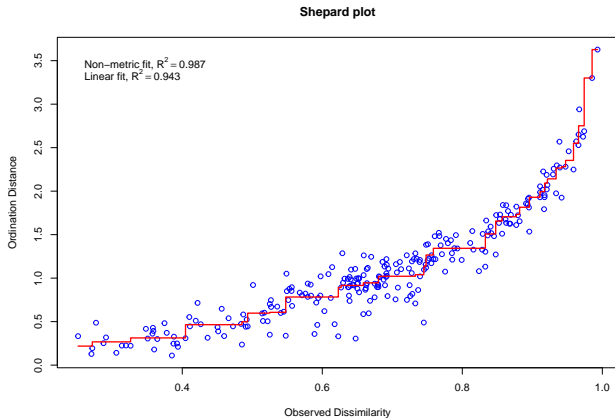
- NDMS using bray distance

<https://mjkeough.github.io/examples/lemmmds.nb.html>

```
## Run 0 stress 0.1156702
## Run 1 stress 0.1153229
## ... New best solution
## ... Procrustes: rmse 0.01893663  max resid 0.07573146
## Run 2 stress 0.115323
## ... Procrustes: rmse 0.0002871722  max resid 0.001111882
## ... Similar to previous best
## Run 3 stress 0.1153229
## ... New best solution
## ... Procrustes: rmse 0.0001334567  max resid 0.0005117371
## ... Similar to previous best
## Run 4 stress 0.115323
## ... Procrustes: rmse 0.0002058442  max resid 0.0008060438
## ... Similar to previous best
## Run 5 stress 0.1153229
## ... Procrustes: rmse 0.0001425949  max resid 0.0005563484
## ... Similar to previous best
## Run 6 stress 0.1172332
## Run 7 stress 0.1156703
## ... Procrustes: rmse 0.01890827  max resid 0.07510586
## Run 8 stress 0.11567
## ... Procrustes: rmse 0.01893865  max resid 0.07545109
## Run 9 stress 0.1156703
## ... Procrustes: rmse 0.0189112  max resid 0.07523154
## Run 10 stress 0.1172331
## Run 11 stress 0.1156701
```

## Example 2: Non-Metric MDS

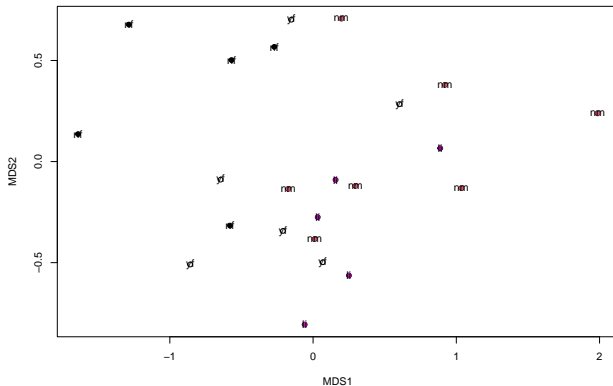
- Stress plot





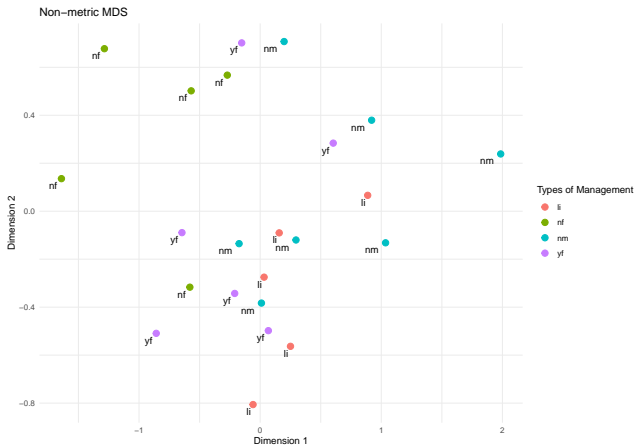
## Example 2: Non-Metric MDS

- biplot



## Example 2: Non-Metric MDS

- biplot

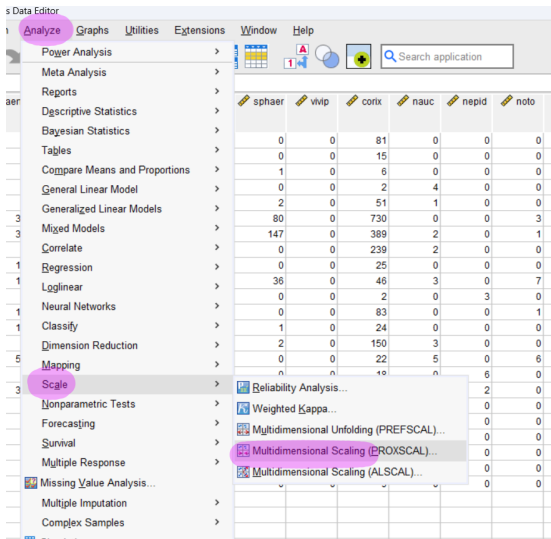


- After importing data, change the measures of all numerical variables to Scale

File	Edit	View	Data	Transform	Analyze	Graphs	Utilities	Extensions	Window	Help					
												Search application			
	Name	Type	Width	Decimals	Label	Values	Missing	Columns	Align	Measure	Role				
1	sife	String	3	0		None	None	7	Left	Nominal	Input				
2	manag	String	2	0		None	None	4	Left	Nominal	Input				
3	managsymb	Numeric	1	0		None	None	8	Right	Nominal	Input				
4	caen	Numeric	2	0		None	None	8	Right	Scale	Input				
5	baet	Numeric	4	0		None	None	8	Right	Scale	Input				
6	acrol	Numeric	2	0		None	None	8	Right	Scale	Input				
7	lymn	Numeric	3	0		None	None	8	Right	Scale	Input				
8	plan	Numeric	4	0		None	None	8	Right	Scale	Input				
9	sphaer	Numeric	3	0		None	None	8	Right	Scale	Input				
10	vlvp	Numeric	1	0		None	None	8	Right	Scale	Input				
11	cortx	Numeric	3	0		None	None	8	Right	Scale	Input				
12	nauc	Numeric	1	0		None	None	8	Right	Scale	Input				
13	nepid	Numeric	1	0		None	None	8	Right	Scale	Input				
14	noto	Numeric	1	0		None	None	8	Right	Scale	Input				
15	pleid	Numeric	3	0		None	None	8	Right	Scale	Input				
16	gamm	Numeric	2	0		None	None	8	Right	Scale	Input				
17	asell	Numeric	1	0		None	None	8	Right	Scale	Input				
18	cerat	Numeric	2	0		None	None	8	Right	Scale	Input				
19	chaob	Numeric	3	0		None	None	8	Right	Scale	Input				
20	chiro	Numeric	4	0		None	None	8	Right	Scale	Input				
21	culic	Numeric	2	0		None	None	8	Right	Scale	Input				
22	cylind	Numeric	1	0		None	None	8	Right	Scale	Input				
23	diuid	Numeric	2	0		None	None	8	Right	Scale	Input				
24	empid	Numeric	1	0		None	None	8	Right	Scale	Input				
25	ephyd	Numeric	1	0		None	None	8	Right	Scale	Input				
26	limon	Numeric	2	0		None	None	8	Right	Scale	Input				
27	psych	Numeric	1	0		None	None	8	Right	Scale	Input				
28	ptych	Numeric	1	0		None	None	8	Right	Scale	Input				
29	sciom	Numeric	1	0		None	None	8	Right	Scale	Input				
30	strat	Numeric	2	0		None	None	8	Right	Scale	Input				

# Lab

- Go to Analyze → Scale → Multidimensional Scaling (ProxSCAL)....
  - PROXSCAL is for metric and non-metric MDS



# Lab

- We have raw data only, so we need to construct a distance matrix

Multidimensional Scaling: Data Format

**Data Format**

☐ The data are proximities


☒ Create proximities from data


**Number of Sources**

☒ One matrix source


☐ Multiple matrix sources


**One Source**


 The proximities are in a matrix across columns.

 The proximities are in a single column.

**Multiple Sources**

 The proximities are in stacked matrices across columns.

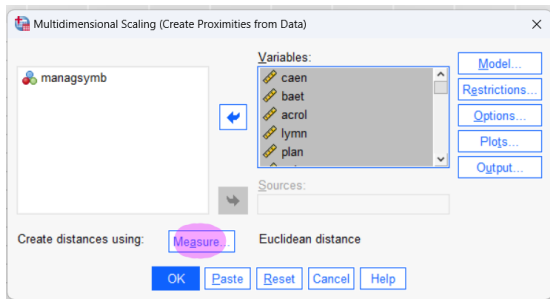
 The proximities are in columns, one source per column.

 The proximities are stacked in a single column.

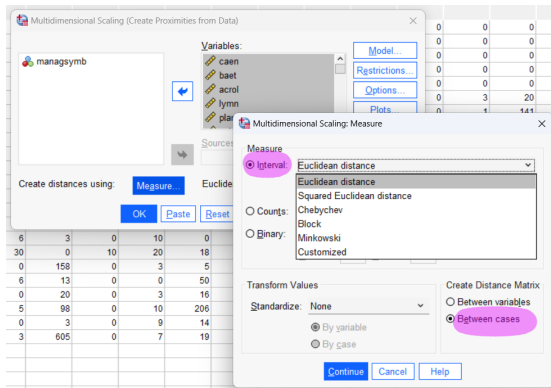
**Define** **Cancel** **Help**

# Lab

- Move all numerical variables to the Variables box



- Suppose we use Euclidean distance
  - ▶ Make sure we create distance matrix **Between cases**



## • Output options

The screenshot displays the 'Multidimensional Scaling: Output' dialog box, which is used to configure the output of the MDS analysis. The dialog is divided into several sections:

- Display:** This section contains checkboxes for the types of output to display. The 'Distances' checkbox is highlighted with a pink box. Other options include 'Common space coordinates', 'Individual space coordinates', 'Individual space weights', 'Multiple stress measures', 'Stress decomposition', 'Transformed proximities', 'Transformed independent variables', and 'Variable and dimension correlations'.
- Save to New File:** This section allows the user to save the output to a new file. It is divided into three sub-sections:
  - Common space coordinates:** Options to 'Create a new dataset' or 'Write a new data file'.
  - Individual space weights:** Options to 'Create a new dataset' or 'Write a new data file'.
  - Transformed proximities:** Options to 'Create a new dataset' or 'Write a new data file'.
- Transformed independent variables:** Options to 'Create a new dataset' or 'Write a new data file'.

In the background, the main MDS window is visible, showing a list of variables (caen, baet, acrol, lyrm, plan) and a table of distances between objects (nepid, noto, pleid, gamm, asell, cerat, chaob, chiro).



- Stress Value: This measures the goodness-of-fit of the MDS solution. Lower values indicate a better fit.
  - ▶ Typically, stress values below 0.1 indicate a good fit.

*Stress and Fit Measures*

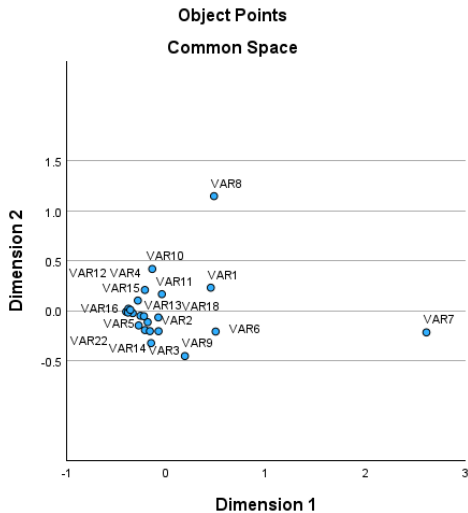
Normalized Raw Stress	.00753
Stress-I	.08678 <sup>a</sup>
Stress-II	.11662 <sup>a</sup>
S-Stress	.01063 <sup>b</sup>
Dispersion Accounted For (D.A.F.)	.99247
Tucker's Coefficient of Congruence	.99623
PROXSCAL minimizes Normalized Raw Stress.	

a. Optimal scaling factor = 1.008.

b. Optimal scaling factor = .994.

*Final Coordinates*

	Dimension	
	1	2
VAR1 Case 1	.445	.231
VAR2 Case 2	-.079	-.206
VAR3 Case 3	-.215	-.194
VAR4 Case 4	-.216	.208
VAR5 Case 5	-.338	-.025
VAR6 Case 6	.494	-.209
VAR7 Case 7	2.610	-.218
VAR8 Case 8	.478	1.148
VAR9 Case 9	.185	-.455
VAR10 Case 10	-.142	.418
VAR11 Case 11	-.046	.166
VAR12 Case 12	-.404	-.010
VAR13 Case 13	-.261	-.051
VAR14 Case 14	-.166	-.205
VAR15 Case 15	-.287	.102
VAR16 Case 16	-.382	.020
VAR17 Case 17	-.386	-.019
VAR18 Case 18	-.081	-.067
VAR19 Case 19	-.188	-.115
VAR20 Case 20	-.362	.008
VAR21 Case 21	-.154	-.325
VAR22 Case 22	-.279	-.148
VAR23 Case 23	-.226	-.055



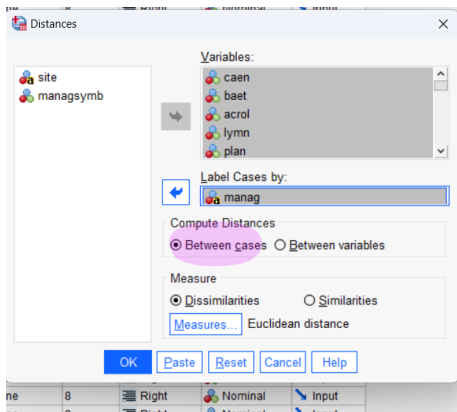
## Distance matrix

Distance

	VAR1 Case 1	VAR2 Case 2	VAR3 Case 3	VAR4 Case 4	VAR5 Case 5	VAR6 Case 6	VAR7 Case 7	VAR8 Case 8	VAR9 Case 9	VAR10 Case 10	VAR11 Case 11	VAR12 Case 12	VAR13 Case 13	VAR14 Case 14	VAR15 Case 15	VAR16 Case 16	VAR17 Case 17	VAR18 Case 18	VAR19 Case 19	VAR20 Case 20	VAR21 Case 21	VAR22 Case 22	VAR23 Case 23
VAR1 Case 1	000																						
VAR2 Case 2	063	000																					
VAR3 Case 3	798	137	800																				
VAR4 Case 4	062	405	402	000																			
VAR5 Case 5	834	566	710	363	000																		
VAR6 Case 6	442	573	710	824	833	800																	
VAR5 Case 7	2 311	2 089	2 826	2 938	2 953	2 116	000																
VAR6 Case 8	918	1 464	1 511	1 669	1 429	1 338	2 332	000															
VAR9 Case 9	733	363	478	775	676	395	2 436	1 630	800														
VAR10 Case 10	617	637	817	223	484	864	2 825	958	932	000													
VAR11 Case 11	467	375	395	178	349	837	2 883	1 113	867	330	000												
VAR12 Case 12	583	380	364	288	068	830	3 822	1 458	738	382	480	000											
VAR13 Case 13	788	239	153	262	082	771	2 876	1 408	802	464	385	546	808										
VAR14 Case 14	751	087	870	414	249	600	2 776	1 489	431	634	390	305	181	000									
VAR15 Case 15	744	371	305	127	137	841	2 915	1 296	730	348	230	367	155	338	000								
VAR16 Case 16	823	377	271	230	082	903	3 823	1 418	740	483	386	838	140	312	122	000							
VAR17 Case 17	868	359	345	263	048	901	3 883	1 452	719	581	387	830	139	388	154	039	000						
VAR18 Case 18	605	188	185	307	261	592	2 895	1 338	470	480	236	328	188	160	267	313	389	000					
VAR19 Case 19	723	142	884	324	175	689	2 830	1 428	385	535	215	540	897	483	238	286	233	117	000				
VAR20 Case 20	838	355	230	247	041	884	2 981	1 416	717	485	354	846	118	396	128	023	036	281	213	000			
VAR21 Case 21	817	141	144	158	352	858	2 708	1 083	383	745	382	402	204	125	447	413	384	268	215	393	000		
VAR22 Case 22	817	388	878	361	137	775	2 860	1 581	558	391	507	889	176	238	187	168	213	086	177	316	308	000	
VAR23 Case 23	738	310	139	263	117	736	2 841	1 384	573	481	285	384	835	160	169	173	164	145	070	151	239	197	000

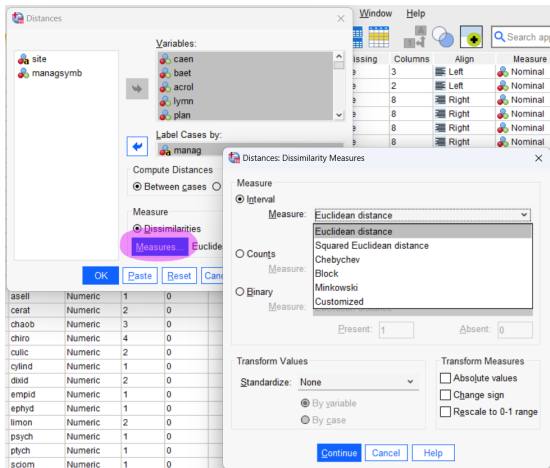
# Lab

- Another method is that we start with the input of a **Distance Matrix**
  - ▶ For example, in SPSS, go to Analyze → Correlate → Distances.
  - ▶ In the dialog box, choose the variables you want to calculate distances for (we select all except the first three variables).
  - ▶ We label the rows by manag



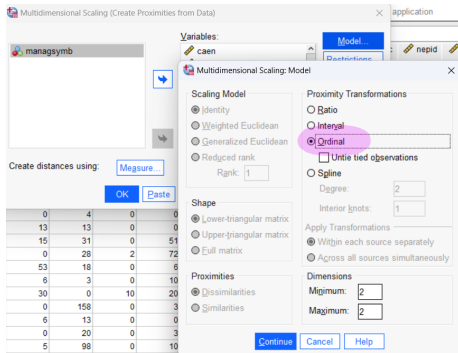
# Lab

- Select the type of distance measure you want to use (e.g., Euclidean).
  - ▶ Click OK to compute the distance matrix. And then export the matrix to an Excel file → Convert to a CSV file and then import this distance matrix



# Lab

- To conduct non-metric MDS with SPSS, we can still click Analyze → Scale → Multidimensional Scaling (ProxSCAL)....
- In the model, we choose Ordinal Proximity transformation



- It is better to have a distance matrix as input. SPSS does not support many distance functions, we will need to prepare our distance matrix using another software.
  - ▶ for example, function `vegdist` in R library `vegan`.

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