UNIVERSIDADE FEDERAL FLUMINENSE

Programa de Mestrado e Doutorado em Engenharia de Produção

Multivariate Data Analysis

Multidimensional Scaling

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Outline

- 1. Definition
- 2. Distances
- 3. Metric MDS
- 4. Non-Metric MDS
 - 5. Bibliography

The MDS (Multidimensional Scaling) or PCoA (Principal Coordinates Analysis), can be considered as an alternative to EFA, and it is a technique of interdependence that allows mapping distances between points (objects or events) in a multidimensional space. It reduces large amounts of data in structures easy to understand, the points that are close together represent similar objects as different objects are represented by points that are distant, so clusters can be interpreted. Characteristics:

- **Data Type**: The input data may be from similarities or dissimilarities (differences or distances). Many authors encourage the use of dissimilarities because their relationship with distances is direct and positive, that is, greater the inequality, greater the distance.
- Analysis Type: Metric MDS (when order between objects does not matter) and Non-Metric MDS (when order between objects does matter).
- **Typical Applications**: Consumer perception of a brand, Perception evaluation of process implementation (before and after effect), Object clustering, etc.

Metric MDS produces a set of uncorrelated (orthogonal) axes to summarize the variability of a dataset, each axis has an eigenvalue that indicates a certain amount of variation. Each object has a coordinate along each axis.

Non-Metric MDS produces an ordination based on a dissimilarity matrix, representing as closely as possible, the pairwise dissimilarity between objects in a low-dimensional space. The original dissimilarity is substituted with ranks, therefore, the information about the magnitude of distances is lost but not its relative position, indicating that clusters possess similar objects.

Limitations:

- The dimensions change with time.
- An object may have bias.
- The interpretation of clusters can vary from researcher to researcher.
- The number of dimensions (between 1 and 3) may not be sufficient to identify the structure of relationship between objects.
- It is an **EXPLORATORY** technique.

Assumptions

ASSUMPTIONS

- **Number of respondents**: At least 1 respondent must evaluate the objects. To achieve an inferential power, sample size must be calculated statistically.
- Number of objects: When the number of objects increases, the more accurate is the output in statistical terms. However, data interpretation can be difficult to achieve. Rule of thumb For a MDS with 2 dimensions is advisable to have at least 10 objects. For a MDS with 3 dimensions it is advisable to have at least 15 objects.
- Number of dimensions: It is recommended that the number of dimensions could range between 1 and 3. For more than 3 dimensions interpretability is questionable and difficult to do.

Distances and Dissimilarities

For similarities, the following equation transforms similarities in dissimilarities.

$$DIS_{ij} = 1 - SIM_{ij}$$

Where:

 DIS_{ij} = degree of dissimilarity between two objects;

 SIM_{ij} = degree of similarity between two objects;

For dissimilarities, the following equation transforms dissimilarities in similarities.

$$SIM_{ij} = 1 - DIS_{ij}$$

For correlations, the following equation transforms correlations in disparity (estimated dissimilarity).

$$\delta_{ij} = \sqrt{2(1-\rho_{ij})}$$

$$or$$

$$\delta_{ij} = 1 - \rho_{ij}$$

Where:

 $\rho_{ij} = \text{Correlation between two objects;}$

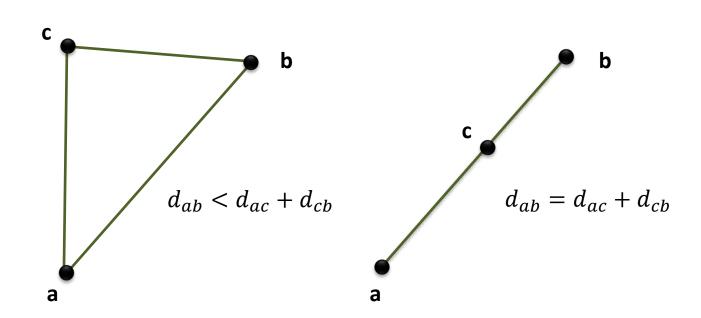
 $\delta_{ii}=$ Disparity.

A true measure of distance, follows three properties:

$$1. d_{ab} = d_{ba}$$

2.
$$d_{ab} \ge 0$$
 e $d_{ab} = 0 \leftrightarrow a = b$

$$3. d_{ab} \le d_{ac} + d_{cb}$$



Distance - Minkowski:

$$d_{mw} = \left(\sum |x_k - y_k|^p\right)^{\frac{1}{p}}$$

Depending on the value p, different distances metrics are obtained. The value of p can not be less than 1, otherwise the triangle inequality is not obeyed.

Distance - Manhattan:

$$d_{mh} = \sum |x_k - y_k|$$

When p = 1; we have the Manhattan distance.

Distance - Euclidian:

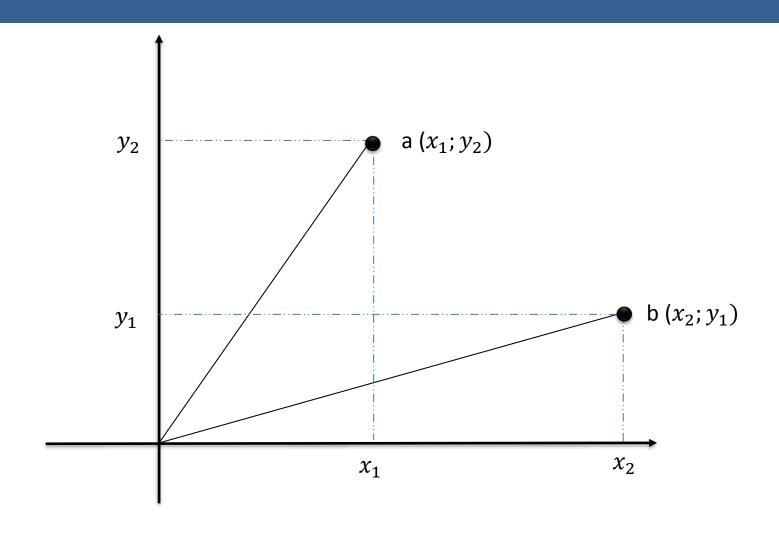
$$d_{ec} = \left(\sum |x_k - y_k|^2\right)^{\frac{1}{2}}$$

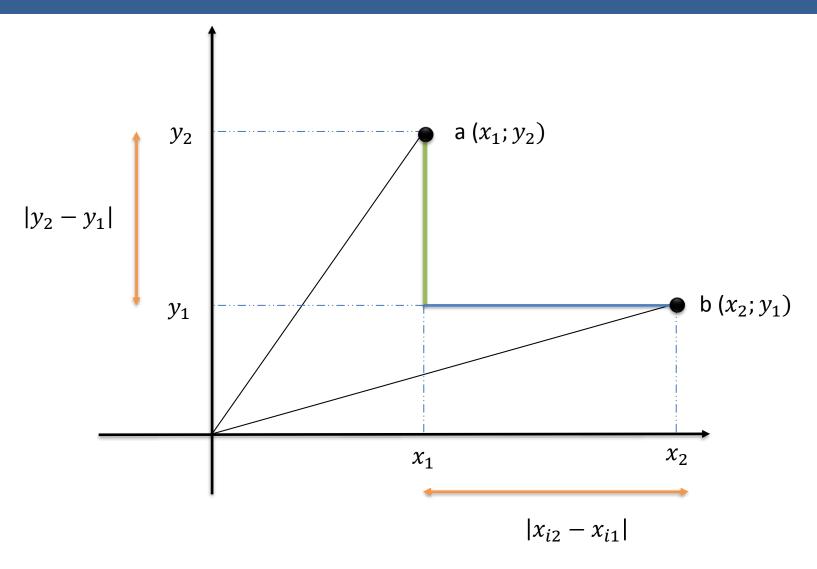
When p = 2; we have the Euclidean distance.

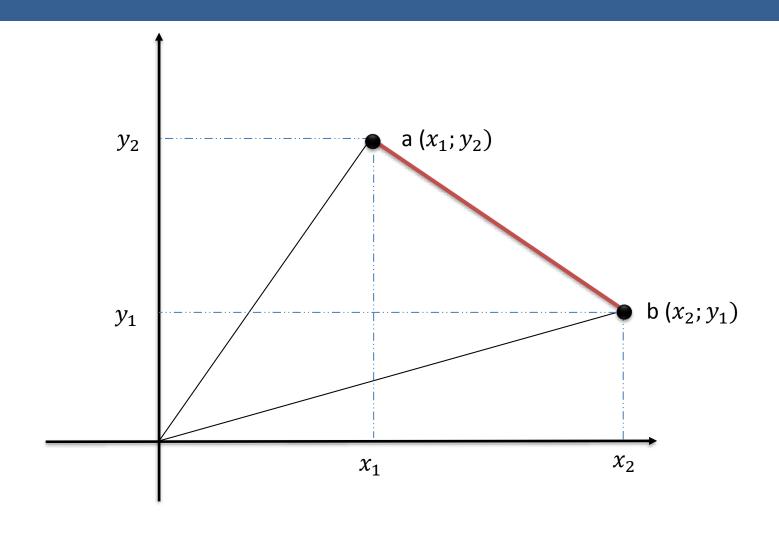
Distance - Chebychev:

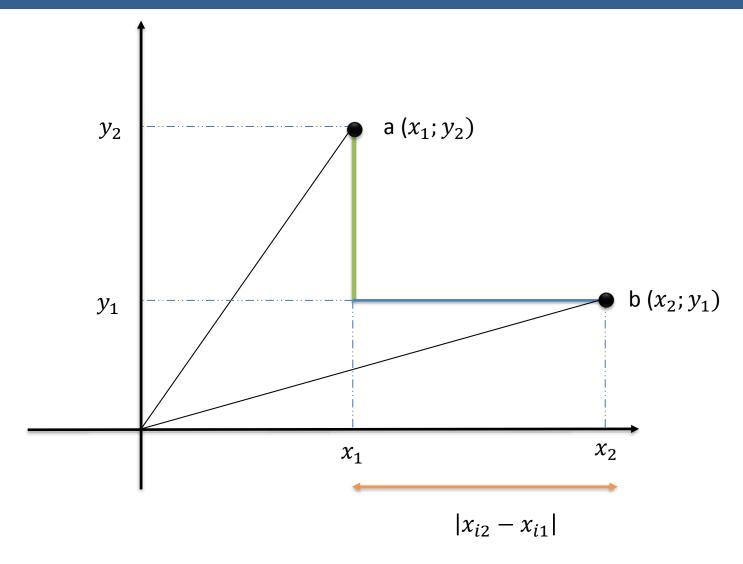
$$d_{cb} = \left(\sum |x_k - y_k|^{\infty}\right)^{\frac{1}{\infty}} = \max|x_k - y_k|$$

When p = 0; we have the Chebychev distance.









Counting Dissimilarity - χ^2 (Chi-square measure: This measure is based on the chi-square test of equality for two sets of frequencies):

$$d_{x2} = \sqrt{\sum_{k} \frac{(x_k - E(x_k))^2}{E(x_k)} + \sum_{k} \frac{(y_k - E(y_k))^2}{E(y_k)}}$$

Counting Dissimilarity - φ^2 (**Phi-square measure**: This measure is equal to the chi-square measure normalized by the square root of the combined frequency):

$$d_{\varphi^{2}} = \sqrt{\frac{\sum_{k} \frac{(x_{k} - E(x_{k}))^{2}}{E(x_{k})} + \sum_{k} \frac{(y_{k} - E(y_{k}))^{2}}{E(y_{k})}}{N}}$$

Counting Dissimilarity - Bray-Curtis:

$$d_{bc} = \frac{\sum |x_k - y_k|}{\sum x_k + y_k}$$

Bray-Curtis is a rank order dissimilarity (used in Non-Metric MDS) between two objects. A value of 0 that the two objects are identical. The similarity can be calculated as:

$$s_{bc} = 1 - d_{bc}$$

A value of 1 means that the two objects are identical.

Binary Dissimilarity:

Coso 1	Case 2			
Case 1	0	1		
0	a	b		
1	С	d		

Binary Dissimilarity – Binary Euclidean:

$$d_{\{0;1\}ec} = \sqrt{b+c}$$

Binary Dissimilarity – Squared Binary Euclidean:

$$d_{\{0;1\}eq} = b + c$$

Binary Dissimilarity – Matching Coefficient:

$$d_{\{0;1\}mc} = \frac{b+c}{(a+b+c+d)}$$

Binary Dissimilarity – Jaccard Index:

$$d_{\{0;1\}jc} = \frac{b+c}{(a+b+c)}$$

Binary Dissimilarity – Pattern Difference:

$$d_{\{0;1\}pd} = \frac{bc}{(a+b+c+d)^2}$$

Binary Dissimilarity – Size Difference:

$$d_{\{0;1\}sd} = \frac{(b+c)^2}{(a+b+c+d)^2}$$

Binary Dissimilarity – Variance:

$$d_{\{0;1\}var} = \frac{b+c}{4(a+b+c+d)}$$

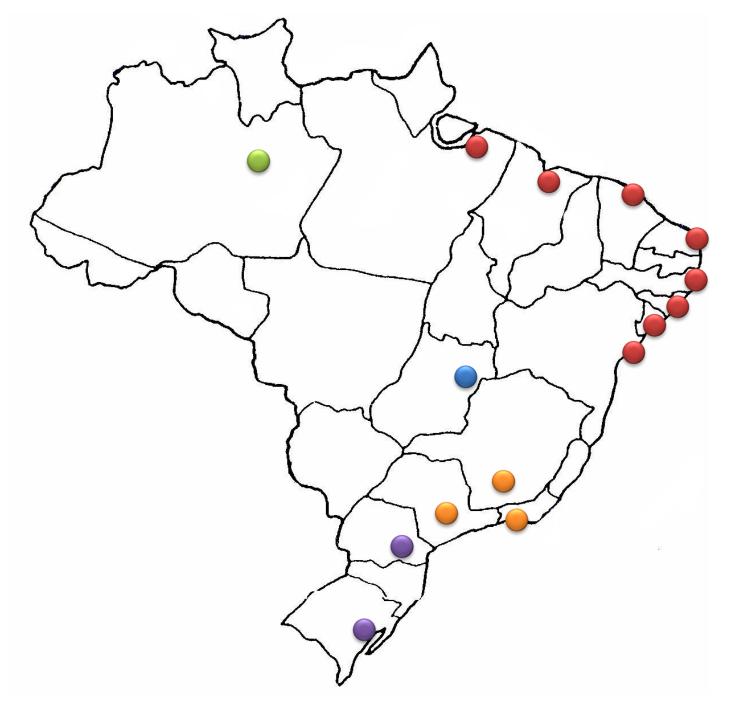
Binary Dissimilarity – Lance Williams:

$$d_{\{0;1\}lw} = \frac{b+c}{2a+b+c}$$

Metric MDS

In order to explain a Metric MDS approach, the following dataset will be used: The collected data represents the Euclidean distance between 15 Brazilian capitals.

	Manaus	Belem	Sao_Luis	Fortaleza	Natal	Maceio	Recife	Aracaju	Salvador	Brasilia	ВН	RJ	SP	Curitiba	Porto_Al
Manaus	0														
Belem	1288	0													
Sao_Luis	1752	493	0												
Fortaleza	2295	1138	640	0											
Natal	2658	1550	1035	444	0										
Maceio	2670	1632	1200	717	435	0									
Recife	2823	1680	1197	640	252	191	0								
Aracaju	2574	1590	1237	810	606	202	386	0							
Salvador	2617	1695	1290	1018	870	464	654	267	0						
Brasilia	1967	1627	1530	1682	1775	1566	1632	1271	1053	0					
BH	2569	2123	1848	1860	1800	1416	1632	1350	980	589	0				
RJ	2854	2460	2271	2190	2122	1680	1865	1485	1220	900	340	0			
SP	3100	2490	2360	2238	2486	1940	2135	1740	1486	865	500	364	0		
Curitiba	2634	2574	2514	2598	2580	2205	2400	2010	1734	1087	823	669	330	0	
Porto_A1	3987	3084	3042	3126	3069	2712	3083	2520	2241	1617	1370	1133	844	547	0



```
# MDS
mds <- cmdscale(my_data, 2, eig = TRUE)
mds$points
                       [,1]
                                      [,2]
Aracaju
                    291.3074
                                  948.45993
Belem
                    981.8100
                                  -381.28651
Belo_Horizonte
                    -754.9364
                                  363.38085
Brasilia
                    -433.8077
                                   -54.72129
Curitiba
                   -1413.0517
                                  -232.91621
                                  559.84226
Fortaleza
                    985.5966
Maceio
                    460.9759
                                  940.98162
Manaus
                    1042.2848
                                  -1871.25345
Natal
                    936.4713
                                  753.20862
Porto Alegre
                   -2034.4776
                                   36.75666
Recife
                    884.6164
                                  878.77954
Rio Janeiro
                    -933.8447
                                  216.05265
Sao Luis
                   1189.6683
                                  -159.71171
Sao Paulo
                   -1135.0267
                                   185.68239
Salvador
                    274.0372
                                  652.88174
```

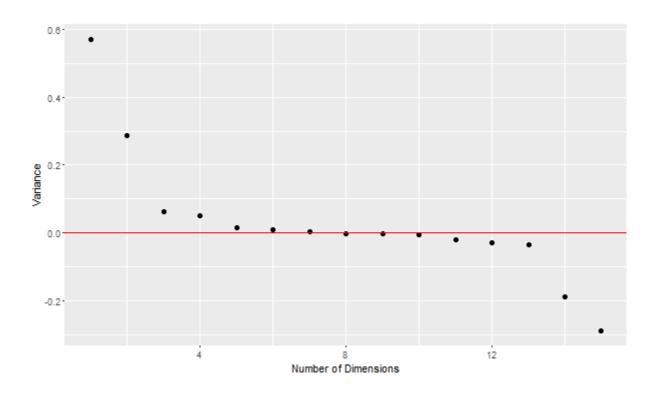
```
# MDS – Extracting 2 Dimensions
mds <- cmdscale(my_data, 2, eig = TRUE)
mds$eig

[1] 15523283.41 7808551.77 1715852.55 1361090.96 459742.14 230271.54
[7] 88970.33 -14663.68 -84346.22 -170203.71 -512328.51 -763414.61
[13] -929346.59 -5096607.32 -7819836.34

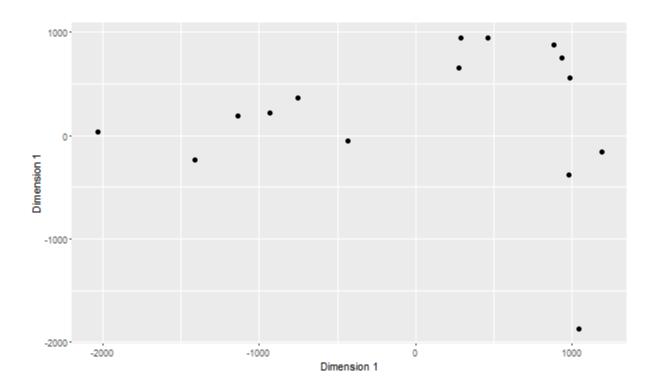
mds$GOF[2] # Percentage of variance explained by 2 dimensions

[1] 0.8581742
```

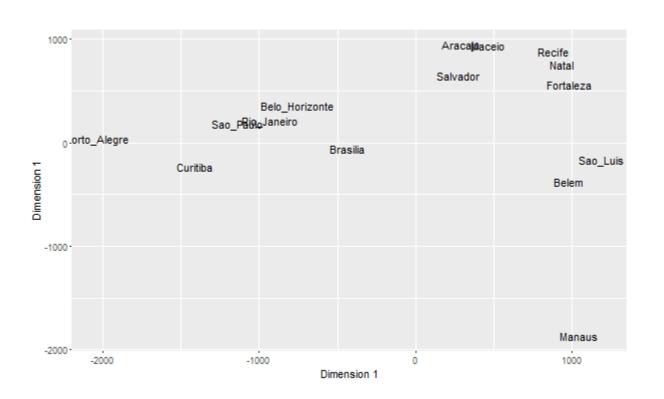
```
# Variance Plot library(ggplot2) ggplot(data = my_data, aes(x = 1:15, y = mds$eig/sum(mds$eig[which(mds$eig > 0)]))) + geom_point(size = 2) + labs(x = "Number of Dimensions", y = "Variance") + geom_hline(yintercept = 0, color = "red")
```



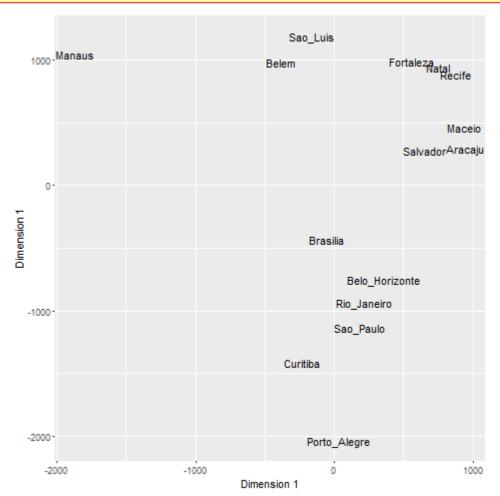
```
# MDS Plot
dimension_x <- mds$points[, 1]
dimension_y <- mds$points[, 2]
library(ggplot2)
ggplot(data = my_data, aes(x = dimension_x, y = dimension_y, label = row.names(my_data))) + geom_point(size = 2) + labs(x = "Dimension 1", y = "Dimension 1")</pre>
```

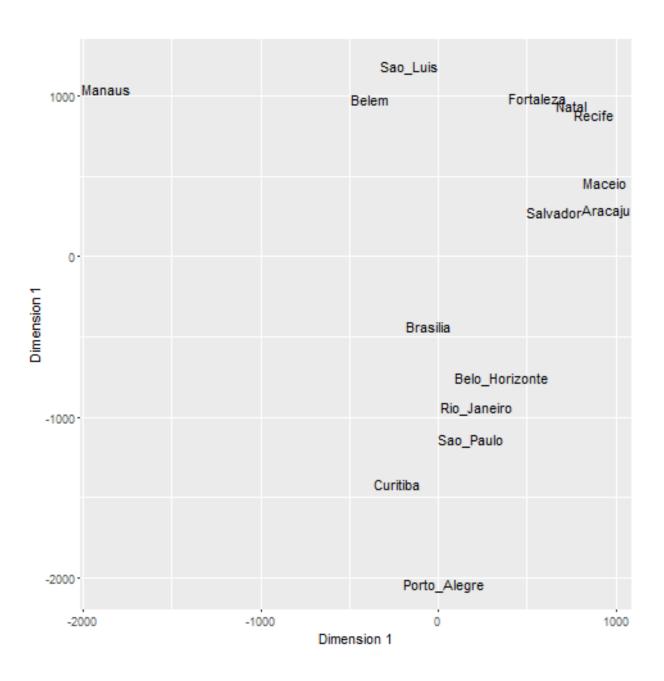


```
# MDS Plot library(ggplot2) ggplot(data = my_{data}, aes(x = dimension_x, y = dimension_y, label = <math>row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data}))
```



```
# MDS Plot library(ggplot2) ggplot(data = my_{data}, aes(x = dimension_y, y = dimension_x, label = <math>row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data})) + row.names(my_{data}))
```

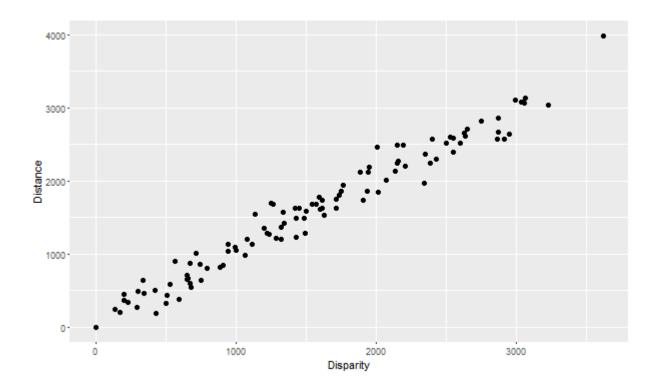




Goodness of Fit

```
# Diagnosis
d_hat <- dist(mds$points, method = "euclidean", diag = TRUE, upper = TRUE, p = 2)
cor (c(as.matrix(d_hat)),c(as.matrix(my_data)))
[1] 0.9843927

# An angle of 45° indicates a good fit.
qplot(x = c(as.matrix(d_hat)), y = c(as.matrix(my_data))) + geom_point(size = 2) + labs(x = "Disparity", y = "Distance")</pre>
```



The Kruskal's *Stress* (Standardized Residual Sum of Squares) is a dimensionless measure that indicates the model's goodness of fit (lower value = best model).

```
p = Number of dimensions;

d_{ij} = Distance between objects;

\delta_{ij} = Estimated distance between objects (disparity).
```

Kruskal's STRESS 2 =
$$\sqrt{\frac{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (d_{ij} - \delta_{ij})^{2}}{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (\delta_{ij} - d_{..})^{2}}}$$

The Young's *Stress* is also dimensionless measure that indicates the model's goodness of fit (lower value = best model).

Young's STRESS 2 =
$$\sqrt{\frac{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (d_{ij}^{2} - \delta_{ij}^{2})^{2}}{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (\delta_{ij}^{2} - d_{..}^{2})^{2}}}$$

Young's S_STRESS =
$$\frac{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (d_{ij}^{2} - \delta_{ij}^{2})^{2}}{\sum_{i=1}^{p-1} \sum_{j=i+1}^{p} (\delta_{ij}^{4})^{4}}$$

$$Raw \ STRESS = \sum (d_{ij} - \delta_{ij})^2$$

Normalized Raw STRESS =
$$\sqrt{\frac{STRESS\ Bruto}{\sum {d_{ij}}^2}}$$

ou

Normalized STRESS =
$$\sqrt{\frac{STRESS\ Bruto}{\sum (d_{ij} - d_{..})^2}}$$

The following interpretation is suggested:

Stress	Goodness of Fit
20.0%	Poor
10.0%	Fair
05.0%	Good
02.5%	Excelent
00.0%	Perfect

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Stress	Goodness of Fit
20.0%	Poor
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00.0%	Perfect

Rule of Thumb: Excelent $\leq 10\%$; Acceptable $> 10\% \, \& < 15\%$; Inacceptable $\geq 15\%$

Other adjustment measures can also be checked.

RSQ (r-squared) = Values close to 1, indicates a good fit.

$$RSQ = \frac{(\sum_{i} \sum_{j} (\delta_{ij} - \delta_{..}) \cdot (d_{ij} - d_{..}))^{2}}{(\sum_{i} \sum_{j} (\delta_{ij} - \delta_{..})^{2}) \cdot (\sum_{i} \sum_{j} ((d_{ij} - d_{..}))^{2})}$$

```
# Diagnosis
d_hat <- as.matrix(d_hat)</pre>
K_STRESS_2 <-((sum((my_data - d_hat)^2))/sum(((d_hat - sum(my_data))^2)))^(1/2)
[1] 0.0004934325
Y_STRESS_2 <- ((sum(((my_data)^2 - d_hat^2)^2))/sum(((d_hat^2 - sum(my_data)^2)^4)))^(1/2)
[1] 4.776563e-17
Y_STRESS_S <- (sum(((my_data)^2 - d_hat^2)^2))/sum(((d_hat^2 - sum(my_data)^2)^4))
[1] 2.281556e-33
```

```
# Diagnosis
d_hat <- as.matrix(d_hat)</pre>
R_STRESS <- sum((my_data - d_hat)^2)
[1] 6345153
N_R_STRESS < - (sum((my_data - d_hat)^2)/sum((my_data)^2))^(1/2)
[1] 0.09468024
N_{STRESS} <- (sum((my_data - d_hat)^2)/sum((my_data - sum(my_data))^2))^(1/2)
[1] 0.0004934924
              ((sum((my_data - sum(my_data))*(d_hat - sum(d_hat))))^2)/(sum((my_data
RSQ <-
sum(my_data))^2)*sum((d_hat - sum(d_hat))^2))
[1] 0.9999998
```

Non-Metric MDS

In order to explain a **Non-Metric MDS** approach, the following dataset will be used: A questionnaire consisting of 12 questions that were applied to 185 students in a college. Each question was answered by a 4-point Likert Scale (1 = Very Bad, 2 = Poor, 3 = Good, 4 = Very Good). The questions were:

Code	Questions (Variables)
PR01	How do you evaluate your commitment to the course?
PR02	Attendance to classes?
PR03	Punctuality?
PR04	Frequency to Library?
PR05	How do you evaluate your general reading habit (literature, magazines, newspapers, internet, etc.)?
PR06	How do you evaluate your reading habit focused on the course?
PR07	Involvement and participation in institutional events (plenary lecture, seminars, lectures, etc.)?
PR08	Involvement and participation in the events of your course (seminars, lectures)?
PR09	Involvement and participation in classroom discussions?
PR10	Involvement and participation in student representation in the institution?
PR11	Level of learning about the content taught by teachers?
PR12	You commitment in doing the activities recommended by the teacher?

```
my_data2 <- R.MVDA.nMDS
library(vegan)
n_mds <- metaMDS(my_data2, k = 2, trymax = 100)
Call:
metaMDS(comm = my_data2, k = 2, trymax = 100)
global Multidimensional Scaling using monoMDS
       my_data2
Data:
Distance: bray
Dimensions: 2
Stress: 0.2306176
Stress type 1, weak ties
No convergent solutions - best solution after 100 tries
Scaling: centring, PC rotation, halfchange scaling
```

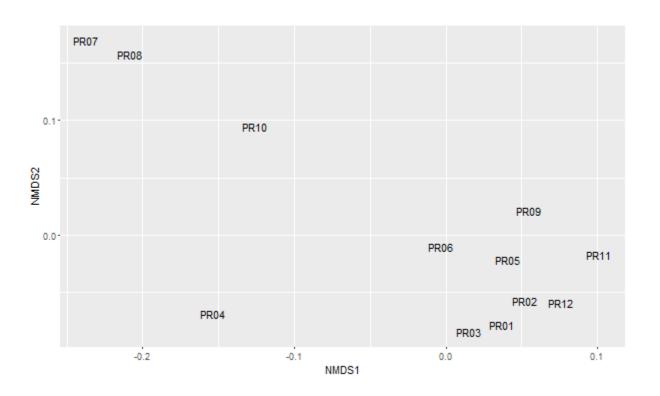
Species: expanded scores based on 'my_data2'

nMDS

The following interpretation is suggested:

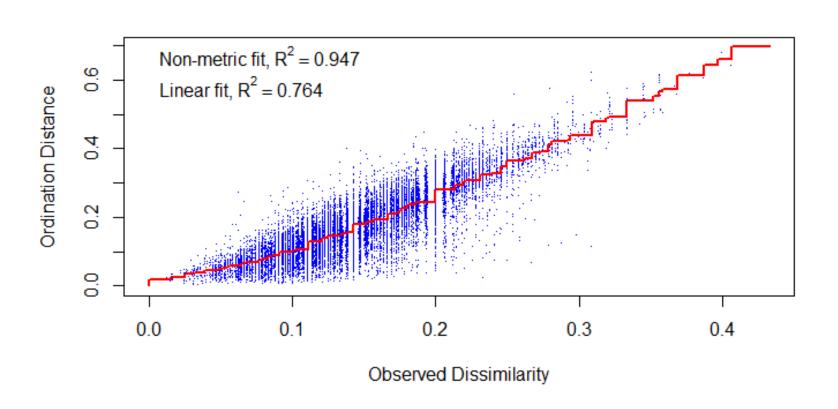
Stress	Goodness of Fit
> 30.0%	Poor
> 20.0% & \le 30.0%	Fair
> 10.0% & \le 20.0%	Good
> 05.0% & \le 10.0%	Great
≤ 05.0%	Excelent
00.0%	Perfect

```
# Plot nMDS Variables
library(ggplot2)
n_mds_variables <- scores(n_mds, "spec")
n_mds_variables <- cbind.data.frame(n_mds_variables, label = rownames(n_mds_variables))
ggplot(data = n_mds_variables, aes(x = NMDS1, y = NMDS2)) + geom_text(aes(label = label))
```



Goodness of Fit

Shepard Plot stressplot(n_mds)



MVDA

https://github.com/Valdecy/Multivariate_Data_Analysis

```
# Created by: Prof. Valdecy Pereira, D.Sc.
```

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- # Course: Multivariate Data Analysis
- # Lesson: Exploratory Factor Analysis

Citation:

PEREIRA, V. (2016). Project: Multivariate Data Analysis, File: R-MVDA-04-MDS.pdf, GitHub

repository: https://github.com/Valdecy/Multivariate_Data_Analysis

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