

# Package ‘MFAg’

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**Type** Package

**Title** Multiple Factor Analysis (MFA)

**Version** 2.3

**Date** 2026-04-13

**Description** Performs Multiple Factor Analysis method for quantitative, categorical, frequency and mixed data, in addition to generating a lot of graphics, also has other useful functions.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**NeedsCompilation** no

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MFAg-package

*Multiple Factor Analysis (MFA)*

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## Description

Performs multiple factor analysis method for quantitative, categorical, frequency and mixed data.

## Details

Package: MFAg  
Type: Package  
Version: 2.3  
Date: 2026-04-13  
License: GPL (>=2)  
LazyLoad: yes

## Author(s)

Paulo Cesar Ossani,

Marcelo Angelo Cirillo

Maintainer: Paulo Cesar Ossani <ossanipc@hotmail.com>

## References

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DataMix

*Mixed data set.*

---

### **Description**

Simulated set of mixed data on consumption of coffee.

### **Usage**

`data(DataMix)`

**Format**

Data set with 10 rows and 7 columns. Being 10 observations described by 7 variables: Cooperatives/Tasters, Average grades given to analyzed coffees, Years of work as a taster, Taster with technical training, Taster exclusively dedicated, Average frequency of the coffees Classified as special, Average frequency of the coffees as commercial.

**Author(s)**

Paulo Cesar Ossani

Marcelo Angelo Cirillo

**Examples**

```
data(DataMix)
DataMix
```

---

DataQuali

*Qualitative data set*

---

**Description**

Set simulated of qualitative data on consumption of coffee.

**Usage**

```
data(DataQuali)
```

**Format**

Data set simulated with 12 rows and 6 columns. Being 12 observations described by 6 variables: Sex, Age, Smoker, Marital status, Sportsman, Study.

**Author(s)**

Paulo Cesar Ossani

Marcelo Angelo Cirillo

**Examples**

```
data(DataQuali)
DataQuali
```

---

DataQuan	<i>Quantitative data set</i>
----------	------------------------------

---

**Description**

Set simulated of quantitative data on grades given to some sensory characteristics of coffees.

**Usage**

```
data(DataQuan)
```

**Format**

Data set with 6 rows and 11 columns. Being 6 observations described by 11 variables: Coffee, Chocolate, Caramelised, Ripe, Sweet, Delicate, Nutty, Caramelised, Chocolate, Spicy, Caramelised.

**Author(s)**

Paulo Cesar Ossani  
Marcelo Angelo Cirillo

**Examples**

```
data(DataQuan)  
DataQuan
```

---

GSVD	<i>Generalized Singular Value Decomposition (GSVD).</i>
------	---

---

**Description**

Given the matrix  $A$  of order  $n \times m$ , the generalized singular value decomposition (GSVD) involves the use of two sets of positive square matrices of order  $n \times n$  and  $m \times m$  respectively. These two matrices express constraints imposed, respectively, on the lines and columns of  $A$ .

**Usage**

```
GSVD(data, plin = NULL, pcol = NULL)
```

**Arguments**

data	Matrix used for decomposition.
plin	Weight for rows.
pcol	Weight for columns

**Details**

If `plin` or `pcol` is not used, it will be calculated as the usual singular value decomposition.

**Value**

`d` Eigenvalues, that is, line vector with singular values of the decomposition.  
`u` Eigenvectors referring rows.  
`v` Eigenvectors referring columns.

**Author(s)**

Paulo Cesar Ossani  
 Marcelo Angelo Cirillo

**References**

Abdi, H. Singular Value Decomposition (SVD) and Generalized Singular Value Decomposition (GSVD). In: SALKIND, N. J. (Ed.). *Encyclopedia of measurement and statistics*. Thousand Oaks: Sage, 2007. p. 907-912.

**Examples**

```
data <- matrix(c(1,2,3,4,5,6,7,8,9,10,11,12), nrow = 4, ncol = 3)

svd(data) # Usual Singular Value Decomposition

GSVD(data) # GSVD with the same previous results

# GSVD with weights for rows and columns
GSVD(data, plin = c(0.1,0.5,2,1.5), pcol = c(1.3,2,0.8))
```

---

IM *Indicator matrix.*

---

**Description**

In the indicator matrix the elements are arranged in the form of *dummy* variables, in other words, 1 for a category chosen as a response variable and 0 for the other categories of the same variable.

**Usage**

```
IM(data, names = TRUE)
```

**Arguments**

`data` Categorical data.  
`names` Include the names of the variables in the levels of the Indicator Matrix (default = TRUE).

**Value**

mtxIndc Returns converted data in the indicator matrix.

**Author(s)**

Paulo Cesar Ossani  
Marcelo Angelo Cirillo

**References**

Rencher, A. C. *Methods of multivariate analysis*. 2th. ed. New York: J.Wiley, 2002. 708 p.

**Examples**

```
data <- matrix(c("S","S","N","N",1,2,3,4,"N","S","T","N"), nrow = 4, ncol = 3)
IM(data, names = FALSE)
data(DataQuali) # qualitative data set
IM(DataQuali, names = TRUE)
```

---

 LocLab

---

*Function for better position of the labels in the graphs.*


---

**Description**

Function for better position of the labels in the graphs.

**Usage**

```
LocLab(x, y = NULL, labels = seq(along = x), cex = 1,
       method = c("SANN", "GA"), allowSmallOverlap = FALSE,
       trace = FALSE, shadotext = FALSE,
       doPlot = TRUE, ...)
```

**Arguments**

x	Coordinate x
y	Coordinate y
labels	The labels
cex	cex
method	Not used
allowSmallOverlap	Boolean
trace	Boolean

shadotext	Boolean
doPlot	Boolean
...	Other arguments passed to or from other methods

**Value**

See the text of the function.

---

MFA *Multiple Factor Analysis (MFA).*

---

**Description**

Perform Multiple Factor Analysis (MFA) on groups of variables. The groups of variables can be quantitative, qualitative, frequency (MFACT) data, or mixed data.

**Usage**

```
MFA(data, groups, typegroups = rep("n",length(groups)), namegroups = NULL)
```

**Arguments**

data	Data to be analyzed.
groups	Number of columns for each group in order following the order of data in 'data'.
typegroups	Type of group: "n" for numerical data (default), "c" for categorical data, "f" for frequency data.
namegroups	Names for each group.

**Value**

vtrG	Vector with the sizes of each group.
vtrNG	Vector with the names of each group.
vtrplin	Vector with the values used to balance the lines of the Z matrix.
vtrpcol	Vector with the values used to balance the columns of the Z matrix.
mtxZ	Matrix concatenated and balanced.
mtxA	Matrix of the eigenvalues (variances) with the proportions and proportions accumulated.
mtxU	Matrix U of the singular decomposition of the matrix Z.
mtxV	Matrix V of the singular decomposition of the matrix Z.
mtxF	Matrix global factor scores where the lines are the observations and the columns the components.
mtxEFG	Matrix of the factor scores by group.
mtxCCP	Matrix of the correlation of the principal components with original variables.
mtxEV	Matrix of the partial inertias / scores of the variables

**Author(s)**

Paulo Cesar Ossani  
 Marcelo Angelo Cirillo

**References**

- Abdessemed, L.; Escofier, B. Analyse factorielle multiple de tableaux de frequences: comparaison avec l'analyse canonique des correspondences. *Journal de la Societe de Statistique de Paris*, Paris, v. 137, n. 2, p. 3-18, 1996..
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- Ossani, P. C.; Cirillo, M. A.; Borem, F. M.; Ribeiro, D. E.; Cortez, R. M. Quality of specialty coffees: a sensory evaluation by consumers using the MFACT technique. *Revista Ciencia Agronomica (UFC. Online)*, v. 48, p. 92-100, 2017.

Pages, J. Analyse factorielle multiple appliquee aux variables qualitatives et aux donnees mixtes. *Revue de Statistique Appliquee*, Paris, v. 50, n. 4, p. 5-37, 2002.

Pages, J.. Multiple factor analysis: main features and application to sensory data. *Revista Colombiana de Estadistica*, Bogota, v. 27, n. 1, p. 1-26, 2004.

### See Also

[Plot.MFA](#)

### Examples

```
data(DataMix) # mixed dataset

data <- DataMix[,2:ncol(DataMix)]

rownames(data) <- DataMix[1:nrow(DataMix),1]

group.names = c("Grade Cafes/Work", "Formation/Dedication", "Coffees")

mf <- MFA(data = data, c(2,2,2), typegroups = c("n","c","f"), group.names) # performs MFA

print("Principal Component Variances:"); round(mf$mtxA,2)

print("Matrix of the Partial Inertia / Score of the Variables:"); round(mf$mtxEV,2)
```

---

NormData

*Normalizes the data.*

---

### Description

Function that normalizes the data globally, or by column.

### Usage

```
NormData(data, type = 1)
```

### Arguments

data	Data to be analyzed.
type	1 normalizes overall (default), 2 normalizes per column.

### Value

dataNorm	Normalized data.
----------	------------------

**Author(s)**

Paulo Cesar Ossani  
Marcelo Angelo Cirillo

**Examples**

```
data(DataQuan) # set of quantitative data

data <- DataQuan[,2:8]

res <- NormData(data, type = 1) # normalizes the data globally

res # Globally standardized data

sd(res) # overall standard deviation

mean(res) # overall mean

res <- NormData(data, type = 2) # normalizes the data per column

res # standardized data per column

apply(res, 2, sd) # standard deviation per column

colMeans(res) # column averages
```

---

Plot.MFA

*Graphics of the Multiple Factor Analysis (MFA).*

---

**Description**

Graphics of the Multiple Factor Analysis (MFA).

**Usage**

```
Plot.MFA(MFA, titles = NA, xlabel = NA, ylabel = NA,
         posleg = 2, boxleg = TRUE, size = 1.1, grid = TRUE,
         color = TRUE, groupcolor = NA, namarr = FALSE,
         linlab = NA, savptc = FALSE, width = 3236,
         height = 2000, res = 300, casc = TRUE)
```

**Arguments**

MFA	Data of the MFA function.
titles	Titles of the graphics, if not set, assumes the default text.
xlabel	Names the X axis, if not set, assumes the default text.

<code>ylabel</code>	Names the Y axis, if not set, assumes the default text.
<code>posleg</code>	1 for caption in the left upper corner, 2 for caption in the right upper corner (default), 3 for caption in the right lower corner, 4 for caption in the left lower corner.
<code>boxleg</code>	Puts frame in legend (default = TRUE).
<code>size</code>	Size of the points in the graphs.
<code>grid</code>	Put grid on graphs (default = TRUE).
<code>color</code>	Colored graphics (default = TRUE).
<code>groupscolor</code>	Vector with the colors of the groups.
<code>namarr</code>	Puts the points names in the cloud around the centroid in the graph corresponding to the global analysis of the Individuals and Variables (default = FALSE).
<code>linlab</code>	Vector with the labels for the observations, if not set, assumes the default text.
<code>savptc</code>	Saves graphics images to files (default = FALSE).
<code>width</code>	Graphics images width when <code>savptc = TRUE</code> (default = 3236).
<code>height</code>	Graphics images height when <code>savptc = TRUE</code> (default = 2000).
<code>res</code>	Nominal resolution in ppi of the graphics images when <code>savptc = TRUE</code> (default = 300).
<code>cas</code>	Cascade effect in the presentation of the graphics (default = TRUE).

**Value**

Returns several graphs.

**Author(s)**

Paulo Cesar Ossani  
Marcelo Angelo Cirillo

**See Also**

[MFA](#)

**Examples**

```
data(DataMix) # set of mixed data

data <- DataMix[,2:ncol(DataMix)]

rownames(data) <- DataMix[1:nrow(DataMix),1]

group.names = c("Grade Cafes/Work", "Formation/Dedication", "Coffees")

mf <- MFA(data, c(2,2,2), typegroups = c("n","c","f"), group.names) # performs MFA

tit <- c("Scree-Plot","Observations","Observations/Variables",
```

```
"Correlation Circle", "Inertia of the Variable Groups")
```

```
Plot.MFA(MFA = mf, titles = tit, xlabel = NA, ylabel = NA,  
  posleg = 2, boxleg = FALSE, color = TRUE,  
  groupcolor = c("blue3", "red", "goldenrod3"),  
  namarr = FALSE, linlab = NA, savptc = FALSE,  
  width = 3236, height = 2000, res = 300,  
  casc = TRUE) # plotting several graphs on the screen
```

```
Plot.MFA(MFA = mf, titles = tit, xlabel = NA, ylabel = NA,  
  posleg = 2, boxleg = FALSE, color = TRUE,  
  namarr = FALSE, linlab = rep("A?", 10),  
  savptc = FALSE, width = 3236, height = 2000,  
  res = 300, casc = TRUE) # plotting several graphs on the screen
```

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