

Package ‘Factoshiny’

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Type Package

Title Perform Factorial Analysis from 'FactoMineR' with a Shiny Application

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Depends FactoMineR,colourpicker,shiny

Description Perform factorial analysis with a menu and draw graphs interactively thanks to 'FactoMineR' and a Shiny application.

License GPL (>= 2)

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Factoshiny-package *Perform classical factorial analysis from FactoMineR within a Shiny app* ~~ Factoshiny ~~

Description

Factoshiny allows to perform CA, PCA, MFA, HCPC and MFA (classical functions from FactoMineR) within a Shiny app. The user can easily change the function parameters and the graphs parameters and can automatically see the results of the change on the plot. All graphs can be downloaded in png, jpg and pdf.

Details

Package: Factoshiny
Type: Package
Version: 1.0
Date: 2015-01-20
License: GPL (>= 2)

Factoshiny have been created to be as easy to use as possible. Thus, only one parameter need to be informed in the several functions. CAshiny, PCAshiny, MCAshiny and MFAshiny need a dataset to work. HCPCshiny need the result of a factorial analysis or a dataframe.

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson

Maintainer: Who to complain to <Magalie.Houee@agrocampus-ouest.fr>

References

~~ Literature or other references for background information ~~

See Also

~~ Optional links to other man pages, e.g. ~~

Description

Performs Correspondance Analysis (CA) including supplementary row and/or column points on a Shiny application.

Columns or rows with NA are considered as supplementary in the analysis.

Allows to change CA parameters and graphical parameters.

Graphics can be downloaded in png, jpg and pdf.

Usage

```
CAshiny(X)
```

Arguments

`X` a data frame or a table with n rows and p columns, i.e. a contingency table or a CAshiny result or a CA result

Value

Returns several tabs:

`Graphs` a tab containing the row and column points factor map (with supplementary columns and supplementary rows)

`Values` a tab containing the summary of the CA performed, the eigenvalues, the results for the columns, for the rows, for the supplementary columns and for the supplementary rows.

`Summary of dataset`

a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.

`Data`

a tab containing the dataset with a nice display.

The left part of the application allows to change all the elements of the CA and the graphs (axes, variables, colors...)

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[CA, plot.CA](#)
[shiny website](#)

Examples

```
## Not run:  
require(FactoMineR)  
data(children)  
# Correspondance Analysis with Factoshiny:  
res.shiny=CAshiny(children)  
  
# Find your app the way you left it (by clicking on the "Quit the app" button)
```

```

res.shiny2=CAshiny(res.shiny)

#CAshiny on a result of a CA
data(children)
res.ca <- CA (children, row.sup = 15:18, col.sup = 6:8)
res.shiny=CAshiny(res.ca)

## End(Not run)

```

FAMDshiny

Factor Analysis for Mixed Data with Factoshiny

Description

Performs Factor Analysis for Mixed Data (FAMD) with supplementary individuals, supplementary quantitative variables and supplementary categorical variables on a Shiny application. Allows to change FAMD parameters and graphical parameters. Graphics can be downloaded in png, jpg and pdf.

Usage

```
FAMDshiny(X)
```

Arguments

X a data frame with n rows (individuals) and p columns (numeric variables and categorical variables), or a FAMDshiny result or a FAMD result

Value

Returns several tabs:

Graphs	a tab containing the individuals factor map, the variables factor (quantitative and categorical) map and the correlation circle.
Values	a tab containing the summary of the FAMD performed, the eigenvalues, the results for the variables, the results for the individuals, the results for the supplementary variables and the results for the supplementary individuals.
Summary of dataset	a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.
Data	a tab containing the dataset with a nice display.

The left part of the application allows to change all the elements of the FAMDA and the graphs (axes, variables, colors...)

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[FAMD,plot.FAMD](#)
[shiny website](#)

Examples

```
## Not run:
require(FactoMineR)
data(geomorphology)
# FAMD with Factoshiny:
res.shiny=FAMDshiny(geomorphology)

# Find your app the way you left it (by clicking on the "Quit the app" button)
res.shiny2=FAMDshiny(res.shiny)

## End(Not run)
```

HCPCshiny	<i>Hierarchical Clustering on Principal Components (HCPC) with Factoshiny</i>
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Description

Performs Hierarchical Clustering on Principal Components (HCPC) o results from a factor analysis on a Shiny application.
 Allows to change HCPC parameters and graphical parmeters.
 Graphics can be downloaded in png, jpg and pdf.

Usage

```
HCPCshiny(res)
```

Arguments

res	Either the result of a factor analysis or a dataframe, the result of a HCPCshiny app or an HCPC result
-----	--

Value

Returns several tabs:

Graphs	a tab containing the factor map,individuals colores by cluster, the 3D map (same graph but with tree above) and the tree map
Values	a tab containing the description of classes (by variables and by axes) and the parangons of each cluster

Summary of dataset

a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.

Data

a tab containing the dataset with a nice display.

The left part of the application allows to change the number of cluster very easily. Some parameters from the graphs can also be changed. By default, the numeric value of the number of clusters is set to the value advised by the HCPC function.

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[HCPC](#), [plot.HCPC](#)
[shiny website](#)

Examples

```
## Not run:
require(FactoMineR)
data(decathlon)
# HCPC with Factoshiny on a result of a factor analysis:
res.pca <- PCA(decathlon, quanti.sup = 11:12, quali.sup=13)
res.shiny=HCPCshiny(res.pca)

# HCPC with Factoshiny on a dataframe:
data(wine)
res.shiny2=HCPCshiny(wine)

# Find your app the way you left it (by clicking on the "Quit the app" button)
res.shiny3=HCPCshiny(res.shiny2)

## End(Not run)
```

MCAshiny

Multiple Correspondence Analysis (MCA) with Factoshiny

Description

Performs Multiple Correspondence Analysis (MCA) with supplementary individuals, supplementary quantitative variables and supplementary categorical variables on a Shiny application.

Allows to change MCA parameters and graphical parameters.

Graphics can be downloaded in png, jpg and pdf.

Usage

MCAshiny(X)

Arguments

X a data frame with n rows (individuals) and p columns (numeric variables), or a MCAshiny result or a MCA result

Value

Returns several tabs:

Graphs a tab containing the individuals factor map and the variables factor map.

Values a tab containing the summary of the MCA performed, the eigenvalues, the results for the variables, the results for the individuals, the results for the supplementary variables and the results for the numerical variables.

Automatic description of axes

a tab containing the output of the dimdesc function. This function is designed to point out the variables and the categories that are the most characteristic according to each dimension obtained by a Factor Analysis.

Summary of dataset

a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.

Data a tab containing the dataset with a nice display.

The left part of the application allows to change all the elements of the MCA and the graphs (axes, variables, colors...)

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[MCA,dimdesc,plot.MCA](#)
[shiny website](#)

Examples

```
## Not run:
require(FactoMineR)
data(poison)
# Multiple Correspondance Analysis with Factoshiny:
res.shiny=MCAshiny(poison)

# Find your app the way you left it (by clicking on the "Quit the app" button)
res.shiny2=MCAshiny(res.shiny)

#MCAshiny on a result of a MCA
data(hobbies)
res.mca <- MCA(hobbies,quali.sup=19:22,quanti.sup=23)
res.shiny=MCAshiny(res.mca)

## End(Not run)
```

Description

Performs Multiple Factor Analysis (MFA) with supplementary individuals and supplementary groups of variables on a Shiny application.

Groups of variables can be quantitative, categorical or contingency tables.

Allows to change MFA parameters and graphical parameters. A maximum of 10 groups can be created

Graphics can be downloaded in png, jpg, pdf and emf.

Usage

```
MFAshiny(X)
```

Arguments

X a data frame with n rows (individuals) and p columns or a result of the MFA function

Value

Returns several tabs:

Graphs a tab containing the individuals factor map, the variables factor map, the groups factor map, the axes factor map and the frequencies factor map (if there is at least one frequency group)

Values a tab containing the summary of the MFA performed, the eigenvalues, the results for the variables, the results for the individuals and the results of the groups

Summary of dataset a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.

Data a tab containing the dataset with a nice display.

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <Francois.Husson@agrocampus-ouest.fr>

See Also

[PCA](#), [plot.PCA](#)
[shiny website](#)

Examples

```
## Not run:
require(FactoMineR)
data(wine)
# Multiple Factor Analysis with Factoshiny:
res.shiny=MFAshiny(wine)

#If you want to create your groups first and customize your graphs
res.mfa= MFA(wine, group=c(2,5,3,10,9,2), type=c("n",rep("s",5)),
  ncp=5, name.group=c("orig","olf","vis","olfag","gust","ens"),
  num.group.sup=c(1,6))
res.shiny=MFAshiny(res.mfa)

## End(Not run)
```

PCAshiny

*Principal Component Analysis (PCA) with FactoShiny***Description**

Performs Principal Component Analysis (PCA) with supplementary individuals, supplementary quantitative variables and supplementary categorical variables on a Shiny application. Allows to change PCA parameters and graphical parameters. Graphics can be downloaded in png, jpg, pdf and emf.

Usage

```
PCAshiny(X)
```

Arguments

X a data frame with n rows (individuals) and p columns (numeric variables) or a PCAshiny result, or a PCA result

Value

Returns several tabs:

Graphs a tab containing the individuals factor map and the variables factor map.

Values a tab containing the summary of the PCA performed, the eigenvalues, the results for the variables, the results for the individuals, the results for the supplementary variables and the results for the categorical variables.

Automatic description of axes a tab containing the output of the dimdesc function. This function is designed to point out the variables and the categories that are the most characteristic according to each dimension obtained by a Factor Analysis.

Summary of dataset

a tab containing the summary of the dataset and a boxplot and histogram for quantitative variables.

Data

a tab containing the dataset with a nice display.

The left part of the application allows to change all the elements of the PCA and the graphs (axes, variables, colors...). The "Get your code" button print the PCA and the plots code in your RStudio session.

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[PCA](#), [plot.PCA](#)
[shiny website](#)

Examples

```
## Not run:
require(FactoMineR)
data(decathlon)
# Principal Component Analysis with Factoshiny:
res.shiny=PCAshiny(decathlon)

# Find your app the way you left it (by clicking on the "Quit the app" button)
res.shiny2=PCAshiny(res.shiny)

#PCAshiny on a result of a PCA
res.pca <- PCA(decathlon, quanti.sup = 11:12, quali.sup=13)
res.shiny=PCAshiny(res.pca)

## End(Not run)
```

print.CAshiny

Print the CAshiny results

Description

Print the CAshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'CAshiny'
print(x,...)
```

Arguments

x an object of class CAshiny
... further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paolinevaissie@gmail.com>

See Also

[CAshiny](#)

Examples

```
## Not run:  
data(children)  
res.shiny = CAshiny(children)  
#click on the "Quit the app" button  
print(res.shiny)  
  
## End(Not run)
```

`print.FAMDshiny` *Print the FAMDshiny results*

Description

Print the FAMDshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'FAMDshiny'  
print(x,...)
```

Arguments

x an object of class FAMDshiny
... further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paolinevaissie@gmail.com>

See Also

[FAMDshiny](#)

Examples

```
## Not run:
data(geomorphology)
res.shiny = FAMDshiny(geomorphology)
#click on the "Quit the app" button
print(res.shiny)

## End(Not run)
```

<code>print.HCPCshiny</code>	<i>Print the HCPCshiny results</i>
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Description

Print the HCPCshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'HCPCshiny'
print(x,...)
```

Arguments

<code>x</code>	an object of class HCPCshiny
<code>...</code>	further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[HCPCshiny](#)

Examples

```
## Not run:
data(decathlon)
res.shiny = HCPCshiny(decathlon)
#click on the "Quit the app" button
print(res.shiny)

## End(Not run)
```

print.MCAshiny	<i>Print the MCAshiny results</i>
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Description

Print the MCAshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'MCAshiny'  
print(x,...)
```

Arguments

x	an object of class MCAshiny
...	further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paolinevaissie@gmail.com>

See Also

[MCAshiny](#)

Examples

```
## Not run:  
data(poison)  
res.shiny = MCAshiny(poison)  
#click on the "Quit the app" button  
print(res.shiny)  
  
## End(Not run)
```

print.MFAshiny	<i>Print the MFAshiny results</i>
----------------	-----------------------------------

Description

Print the MFAshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'MFAshiny'  
print(x,...)
```

Arguments

x an object of class MFAshiny
... further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[MFAshiny](#)

Examples

```
## Not run:  
data(wine)  
res<- MFA(wine, group=c(2,5,3,10,9,2), type=c("n",rep("s",5)),  
          ncp=5, name.group=c("orig","olf","vis","olfag","gust","ens"),  
          num.group.sup=c(1,6),graph=FALSE)  
res.shiny=MFAshiny(res)  
  
#click on the "Quit the app" button  
print(res.shiny)  
  
## End(Not run)
```

`print.PCAshiny` *Print the PCAshiny results*

Description

Print the PCAshiny results i.e the R corresponding script

Usage

```
## S3 method for class 'PCAshiny'  
print(x,...)
```

Arguments

x an object of class PCAshiny
... further arguments passed to or from other methods

Author(s)

Pauline Vaissie, Astrid Monge, Francois Husson <paulinevaissie@gmail.com>

See Also

[PCAshiny](#)

Examples

```
## Not run:  
data(decathlon)  
res.shiny = PCAshiny(decathlon)  
#click on the "Quit the app" button  
print(res.shiny)  
  
## End(Not run)
```

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