

Package ‘explor’

May 2, 2020

Type Package

Title Interactive Interfaces for Results Exploration

Version 0.3.6

Date 2020-05-02

Maintainer Julien Barnier <julien.barnier@cnrs.fr>

Description Shiny interfaces and graphical functions for multivariate analysis results exploration.

License GPL (>= 3)

VignetteBuilder knitr

URL <https://juba.github.io/explor/>

BugReports <https://github.com/juba/explor/issues>

Encoding UTF-8

Imports shiny (>= 1.0), DT, dplyr (>= 0.7), tidyr, ggplot2, highr,
formatR, scatterD3 (>= 0.9.1), RColorBrewer

Suggests FactoMineR, ade4 (>= 1.7-13), GDAtools, MASS, quanteda,
quanteda.textmodels, testthat, knitr, rmarkdown

RoxygenNote 7.1.0

NeedsCompilation no

Author Julien Barnier [aut, cre]

Repository CRAN

Date/Publication 2020-05-02 14:30:09 UTC

R topics documented:

CA_var_plot	2
explor	3
ggind	5
ggvar	6
MCA_biplot	7
MCA_ind_plot	8

MCA_var_plot	9
PCA_ind_plot	10
PCA_var_plot	11
prepare_results	12

Index**14**

CA_var_plot	<i>Interactive CA variables plot</i>
--------------------	--------------------------------------

Description

This function generates an HTML widget displaying the variables plot of a CA result.

Usage

```
CA_var_plot(
  res,
  xax = 1,
  yax = 2,
  lev_sup = TRUE,
  var_sup = TRUE,
  var_sup_choice = NULL,
  var_hide = "None",
  var_lab_min_contrib = 0,
  point_size = 64,
  col_var = NULL,
  symbol_var = NULL,
  size_var = NULL,
  size_range = c(10, 300),
  zoom_callback = NULL,
  in_explor = FALSE,
  ...
)
```

Arguments

res	Result of prepare_results() call
xax	Horizontal axis number
yax	Vertical axis number
lev_sup	TRUE to display supplementary levels
var_sup	TRUE to display supplementary variables
var_sup_choice	list of supplementary variables to display
var_hide	elements to hide (rows or columns)
var_lab_min_contrib	Contribution threshold to display points labels

point_size	base point size
col_var	name of the variable for points color
symbol_var	name of the variable for points symbol
size_var	name of the variable for points size
size_range	points size range with format c(minimum, maximum)
zoom_callback	scatterD3 zoom callback JavaScript body
in_explor	wether the plot is to be displayed in the explor interface
...	Other arguments passed to scatterD3

explor*Interface for analysis results exploration*

Description

This function launches a shiny app in a web browser in order to do interactive visualisation and exploration of an analysis results.

Usage

```
explor(obj)

## S3 method for class 'CA'
explor(obj)

## S3 method for class 'textmodel_ca'
explor(obj)

## S3 method for class 'coa'
explor(obj)

## S3 method for class 'MCA'
explor(obj)

## S3 method for class 'speMCA'
explor(obj)

## S3 method for class 'mca'
explor(obj)

## S3 method for class 'acm'
explor(obj)

## S3 method for class 'PCA'
explor(obj)
```

```
## S3 method for class 'princomp'
explor(obj)

## S3 method for class 'prcomp'
explor(obj)

## S3 method for class 'pca'
explor(obj)
```

Arguments

`obj` object containing analysis results

Details

If you want to display supplementary individuals or variables and you're using the `dudi.coa` function, you can add the coordinates of `suprow` and/or `supcol` to as `supr` and/or `supr` elements added to your `dudi.coa` result (See example).

If you want to display supplementary individuals or variables and you're using the `dudi.acm` function, you can add the coordinates of `suprow` and/or `supcol` to as `supi` and/or `supv` elements added to your `dudi.acm` result (See example).

If you want to display supplementary individuals or variables and you're using the `dudi.pca` function, you can add the coordinates of `suprow` and/or `supcol` to as `supi` and/or `supv` elements added to your `dudi.pca` result (See example).

Value

The function launches a shiny app in the system web browser.

Examples

```
## Not run:

require(FactoMineR)

## FactoMineR::MCA exploration
data(hobbies)
mca <- MCA(hobbies[1:1000,c(1:8,21:23)], quali.sup = 9:10,
            quanti.sup = 11, ind.sup = 1:100, graph = FALSE)
explor(mca)

## FactoMineR::PCA exploration
data(decathlon)
d <- decathlon[,1:12]
pca <- PCA(d, quanti.sup = 11:12, graph = FALSE)
explor(pca)

## End(Not run)
## Not run:
```

```

library(ade4)

data(bordeaux)
tab <- bordeaux
row_sup <- tab[5,-4]
col_sup <- tab[-5,4]
coa <- dudi.coa(tab[-5,-4], nf = 5, scannf = FALSE)
coa$supr <- suprow(coa, row_sup)
coa$supc <- supcol(coa, col_sup)
explor(coa)

## End(Not run)
## Not run:

library(ade4)
data(banque)
d <- banque[-(1:100), -(19:21)]
ind_sup <- banque[1:100, -(19:21)]
var_sup <- banque[-(1:100), 19:21]
acm <- dudi.acm(d, scannf = FALSE, nf = 5)
acm$supv <- supcol(acm, dudi.acm(var_sup, scannf = FALSE, nf = 5)$tab)
colw <- acm$cw*ncol(d)
X <- acm.disjonctif(ind_sup)
X <- data.frame(t(t(X)/colw) - 1)
acm$supi <- suprow(acm, X)
explor(acm)

## End(Not run)
## Not run:

library(ade4)
data(deug)
d <- deug$tab
sup_var <- d[(-(1:10), 8:9]
sup_ind <- d[1:10, -(8:9)]
pca <- dudi.pca(d[(-(1:10), -(8:9)], scale = TRUE, scannf = FALSE, nf = 5)
supi <- suprow(pca, sup_ind)
pca$supi <- supi
supv <- supcol(pca, dudi.pca(sup_var, scale = TRUE, scannf = FALSE)$tab)
pca$supv <- supv
explor(pca)

## End(Not run)

```

Description

This function displays a graphical representation of the individuals (rows) of a multivariate analysis.

This function displays a graphical representation of the individuals (rows) of a multiple correspondence analysis generated by the MCA function of the FactoMineR package.

Usage

```
gginde(obj, ...)

## S3 method for class 'MCA'
gginde(
  obj,
  xax = 1,
  yax = 2,
  fac = NA,
  label = NULL,
  alpha = 0.5,
  palette = "Set1",
  ...
)
```

Arguments

<code>obj</code>	a multivariate analysis results object. Currently only MCA is supported
<code>...</code>	arguments passed to other methods
<code>xax</code>	number of the x axis
<code>yax</code>	number of the y axis
<code>fac</code>	an optional factor by which points are colored, and confidence ellipses drawn
<code>label</code>	legend title
<code>alpha</code>	points opacity
<code>palette</code>	palette for points coloring, if <code>fac</code> is not <code>NULL</code>

<code>ggvar</code>	<i>Graphical representation of the variables (columnss) of a multivariate analysis</i>
--------------------	--

Description

This function displays a graphical representation of the variables (columns) of a multivariate analysis.

This function displays a graphical representation of the variables (columns) of a multiple correspondence analysis generated by the MCA function of the FactoMineR package.

Usage

```
ggvar(obj, ...)

## S3 method for class 'MCA'
ggvar(obj, xax = 1, yax = 2, size = 4, alpha = 0.5, palette = "Set1", ...)
```

Arguments

obj	a multivariate analysis results object. Currently only MCA is supported
...	arguments passed to other methods
xax	number of the x axis
yax	number of the y axis
size	text size
alpha	points opacity
palette	palette for variables coloring

See Also

[MCA](#)

MCA_biplot

Interactive MCA biplot

Description

This function generates an HTML widget displaying the variables plot of an MCA result.

Usage

```
MCA_biplot(  
  res,  
  xax = 1,  
  yax = 2,  
  col_var,  
  ind_sup = TRUE,  
  var_sup = TRUE,  
  bi_lab_min_contrib = 0,  
  symbol_var = NULL,  
  ind_point_size = 16,  
  var_point_size = 96,  
  ind_opacity = 0.5,  
  ind_opacity_var = NULL,  
  ind_labels = FALSE,  
  zoom_callback = NULL,  
  in_explor = FALSE,  
  ...  
)
```

Arguments

<code>res</code>	Result of <code>prepare_results()</code> call
<code>xax</code>	Horizontal axis number
<code>yax</code>	Vertical axis number
<code>col_var</code>	name of the variable for points color
<code>ind_sup</code>	TRUE to display supplementary individuals
<code>var_sup</code>	TRUE to display supplementary variables
<code>bi_lab_min_contrib</code>	Contribution threshold to display points labels
<code>symbol_var</code>	name of the variable for points symbol
<code>ind_point_size</code>	base point size for individuals
<code>var_point_size</code>	base point size for variable levels
<code>ind_opacity</code>	individuals point opacity (constant)
<code>ind_opacity_var</code>	individuals point opacity (variable)
<code>ind_labels</code>	TRUE to display individuals labels
<code>zoom_callback</code>	scatterD3 zoom callback JavaScript body
<code>in_explor</code>	wether the plot is to be displayed in the explor interface
<code>...</code>	Other arguments passed to scatterD3

`MCA_ind_plot`

Interactive MCA individuals plot

Description

This function generates an HTML widget displaying the individuals plot of an MCA result.

Usage

```
MCA_ind_plot(
  res,
  xax = 1,
  yax = 2,
  ind_sup = TRUE,
  ind_lab_min_contrib = 0,
  lab_var = NULL,
  col_var = NULL,
  symbol_var = NULL,
  opacity_var = NULL,
  size_var = NULL,
  size_range = c(10, 300),
  zoom_callback = NULL,
  in_explor = FALSE,
  ...
)
```

Arguments

res	Result of prepare_results() call
xax	Horizontal axis number
yax	Vertical axis number
ind_sup	TRUE to display supplementary individuals
ind_lab_min_contrib	Contribution threshold to display points labels
lab_var	variable to be used for points names
col_var	variable to be used for points color
symbol_var	name of the variable for points symbol
opacity_var	name of the variable for points opacity
size_var	name of the variable for points size
size_range	points size range with format c(minimum, maximum)
zoom_callback	scatterD3 zoom callback JavaScript body
in_explor	wether the plot is to be displayed in the explor interface
...	Other arguments passed to scatterD3

Description

This function generates an HTML widget displaying the variables plot of an MCA result.

Usage

```
MCA_var_plot(
  res,
  xax = 1,
  yax = 2,
  var_sup = TRUE,
  var_sup_choice = NULL,
  var_lab_min_contrib = 0,
  point_size = 64,
  labels_prepend_var = FALSE,
  col_var = NULL,
  symbol_var = NULL,
  size_var = NULL,
  size_range = c(10, 300),
  zoom_callback = NULL,
  in_explor = FALSE,
  ...
)
```

Arguments

<code>res</code>	Result of <code>prepare_results()</code> call
<code>xax</code>	Horizontal axis number
<code>yax</code>	Vertical axis number
<code>var_sup</code>	TRUE to display supplementary variables
<code>var_sup_choice</code>	list of supplementary variables to display
<code>var_lab_min_contrib</code>	Contribution threshold to display points labels
<code>point_size</code>	base point size
<code>labels_prepend_var</code>	if TRUE, prepend variable names to labels
<code>col_var</code>	name of the variable for points color
<code>symbol_var</code>	name of the variable for points symbol
<code>size_var</code>	name of the variable for points size
<code>size_range</code>	points size range with format <code>c(minimum, maximum)</code>
<code>zoom_callback</code>	scatterD3 zoom callback JavaScript body
<code>in_explor</code>	wether the plot is to be displayed in the explor interface
<code>...</code>	Other arguments passed to scatterD3

`PCA_ind_plot`*Interactive PCA individuals plot***Description**

This function generates an HTML widget displaying the individuals plot of a PCA result.

Usage

```
PCA_ind_plot(
  res,
  xax = 1,
  yax = 2,
  ind_sup = TRUE,
  ind_lab_min_contrib = 0,
  col_var = NULL,
  symbol_var = NULL,
  opacity_var = NULL,
  size_var = NULL,
  size_range = c(10, 300),
  lab_var = NULL,
  zoom_callback = NULL,
  in_explor = FALSE,
  ...
)
```

Arguments

res	Result of prepare_results() call
xax	Horizontal axis number
yax	Vertical axis number
ind_sup	TRUE to display supplementary individuals
ind_lab_min_contrib	Contribution threshold to display points labels
col_var	variable to be used for points color
symbol_var	name of the variable for points symbol
opacity_var	name of the variable for points opacity
size_var	name of the variable for points size
size_range	points size range with format c(minimum, maximum)
lab_var	variable to be used for points names
zoom_callback	scatterD3 zoom callback JavaScript body
in_explor	wether the plot is to be displayed in the explor interface
...	Other arguments passed to scatterD3

Description

This function generates an HTML widget displaying the variables plot of a PCA result.

Usage

```
PCA_var_plot(
  res,
  xax = 1,
  yax = 2,
  var_sup = TRUE,
  var_sup_choice = NULL,
  var_lab_min_contrib = 0,
  scale_unit = FALSE,
  col_var = NULL,
  size_var = NULL,
  zoom_callback = NULL,
  in_explor = FALSE,
  xlim = NULL,
  ylim = NULL,
  ...
)
```

Arguments

<code>res</code>	Result of <code>prepare_results()</code> call
<code>xax</code>	Horizontal axis number
<code>yax</code>	Vertical axis number
<code>var_sup</code>	TRUE to display supplementary variables
<code>var_sup_choice</code>	list of supplementary variables to display
<code>var_lab_min_contrib</code>	Contribution threshold to display points labels
<code>scale_unit</code>	wether the PCA is scaled
<code>col_var</code>	name of the variable for points color
<code>size_var</code>	name of the variable for points size
<code>zoom_callback</code>	<code>scatterD3</code> zoom callback JavaScript body
<code>in_explor</code>	wether the plot is to be displayed in the <code>explor</code> interface
<code>xlim</code>	custom x axis limits
<code>ylim</code>	custom y axis limits
<code>...</code>	Other arguments passed to <code>scatterD3</code>

`prepare_results` *Analysis results preparation*

Description

This function prepares results to be used by `explor`. Not to be used directly.

Usage

```
prepare_results(obj)

## S3 method for class 'CA'
prepare_results(obj)

## S3 method for class 'mca'
prepare_results(obj)

## S3 method for class 'MCA'
prepare_results(obj)

## S3 method for class 'PCA'
prepare_results(obj)

## S3 method for class 'coa'
prepare_results(obj)
```

```
## S3 method for class 'acm'  
prepare_results(obj)  
  
## S3 method for class 'pca'  
prepare_results(obj)  
  
## S3 method for class 'prcomp'  
prepare_results(obj)  
  
## S3 method for class 'princomp'  
prepare_results(obj)  
  
## S3 method for class 'speMCA'  
prepare_results(obj)  
  
## S3 method for class 'textmodel_ca'  
prepare_results(obj)
```

Arguments

`obj` object containing analysis results

See Also

[CA](#)
[mca](#)
[MCA](#)
[PCA](#)
[CA](#)
[dudi.acm](#)
[dudi.pca](#)
[prcomp](#)
[princomp](#)
[speMCA](#)
[textmodel_ca](#)

Index

CA, [13](#)
CA_var_plot, [2](#)

dudi.acm, [4](#), [13](#)
dudi.coa, [4](#)
dudi.pca, [4](#), [13](#)

explor, [3](#)

ggind, [5](#)
ggvar, [6](#)

MCA, [7](#), [13](#)
mca, [13](#)
MCA_biplot, [7](#)
MCA_ind_plot, [8](#)
MCA_var_plot, [9](#)

PCA, [13](#)
PCA_ind_plot, [10](#)
PCA_var_plot, [11](#)
prcomp, [13](#)
prepare_results, [12](#)
princomp, [13](#)

speMCA, [13](#)
supcol, [4](#)
suprow, [4](#)

textmodel_ca, [13](#)