Package 'mcvis'

October 13, 2022

Type Package		
Title Multi-Collinearity Visualization		
Version 1.0.8		
Description Visualize the relationship between linear regression variables and causes of multi-collinearity. Implements the method in Lin et. al. (2020) <doi:10.1080 10618600.2020.1779729="">.</doi:10.1080>		
Encoding UTF-8		
Imports assertthat, igraph, ggplot2, purrr, magrittr, reshape2, shiny, dplyr, psych, rlang		
RoxygenNote 7.1.1.9001		
License GPL-3		
Suggests testthat (>= 2.1.0), covr, knitr, rmarkdown		
VignetteBuilder knitr		
NeedsCompilation no		
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Repository CRAN		
Date/Publication 2021-07-30 08:20:05 UTC		
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alt_mcvis

Multi-collinearity Visualization plots

Description

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Usage

```
alt_mcvis(mcvis_result, eig_max = 1L, var_max = ncol(mcvis_result$MC))
ggplot_mcvis(
 mcvis_result,
 eig_max = 1L,
 var_max = ncol(mcvis_result$MC),
  label_dodge = FALSE
)
igraph_mcvis(mcvis_result, eig_max = 1L, var_max = ncol(mcvis_result$MC))
## S3 method for class 'mcvis'
plot(
  х,
  type = c("ggplot", "igraph", "alt"),
  eig_max = 1L,
  var_max = ncol(x$MC),
  label_dodge = FALSE,
)
```

Arguments

mcvis_result	Output of the mcvis function
eig_max	The maximum number of eigenvalues to be displayed on the plot.
var_max	The maximum number of variables to be displayed on the plot.
label_dodge	If variable names are too long, it might be helpful to dodge the labelling. Default to FALSE.
Х	Output of the mcvis function
type	Plotting mcvis result using "igraph" or "ggplot". Default to "ggplot".
	additional arguments (currently unused)

Value

A mcvis visualization plot

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Author(s)

Chen Lin, Kevin Wang, Samuel Mueller

Examples

```
set.seed(1)
p = 10
n = 100
X = matrix(rnorm(n*p), ncol = p)
X[,1] = X[,2] + rnorm(n, 0, 0.1)
mcvis_result = mcvis(X)
plot(mcvis_result)
plot(mcvis_result, type = "igraph")
plot(mcvis_result, type = "alt")
```

mcvis

Multi-collinearity Visualization

Description

Multi-collinearity Visualization

Usage

```
mcvis(
   X,
   sampling_method = "bootstrap",
   standardise_method = "studentise",
   times = 1000L,
   k = 10L
)
```

Arguments

X A matrix of regressors (without intercept terms).

sampling_method

The resampling method for the data. Currently supports 'bootstrap' or 'cv' (cross-validation).

standardise_method

The standardisation method for the data. Currently supports 'euclidean' (default, centered by mean and divide by Euclidean length) and 'studentise' (centred by mean and divide by standard deviation)

times Number of resampling runs we perform. Default is set to 1000.

k Number of partitions in averaging the MC-index. Default is set to 10.

shiny_mcvis

Value

A list of outputs:

- t_square: The t^2 statistics for the regression between the VIFs and the tau's.
- MC:The MC-indices
- col_names:Column names (export for plotting purposes)

Author(s)

Chen Lin, Kevin Wang, Samuel Mueller

Examples

```
set.seed(1)
p = 10
n = 100
X = matrix(rnorm(n*p), ncol = p)
X[,1] = X[,2] + rnorm(n, 0, 0.1)
mcvis_result = mcvis(X = X)
mcvis_result
```

shiny_mcvis

Shiny app for mcvis exploration

Description

Shiny app for mcvis exploration

Usage

```
shiny_mcvis(mcvis_result, X)
```

Arguments

```
mcvis_result Output of the mcvis function
X The original X matrix
```

Value

A shiny app allowing for interactive exploration of mcvis results

Author(s)

Chen Lin, Kevin Wang, Samuel Mueller

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Examples

```
if(interactive()){
set.seed(1)
p = 10
n = 100
X = matrix(rnorm(n*p), ncol = p)
mcvis_result = mcvis(X)
shiny_mcvis(mcvis_result = mcvis_result, X = X)
}
```

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