

Package ‘rstatsToolkit’

January 9, 2015

Title Bundles up all my most used functions for doing statistical analysis

Version 0.1

Description This package is mainly my personal collection of code that I commonly use in my research and statistical analysis. Eventually I would like to develop it into a toolkit that other graduate students (I'm a graduate student right now btw) and in the future for my own graduate students to use and develop. Until that point, I am slowly developing this package into toolkit for analyzing and exploring data.

Imports ggplot2 (>= 1.0.0),
plyr (>= 1.8.1),
grid (>= 3.1.1),
visreg (>= 2.0.5),
reshape2 (>= 1.4)

Depends R (>= 3.1.1)

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LazyData true

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bivarPlot

Bivariate plot

Description

In development ..

Usage

```
bivarPlot(x, y, data, ...)
```

Arguments

x

y

data

...

Details

.. content for details ..

Value

Outputs a plot

Author(s)

Luke Johnston

diagnosticPlots*Regression diagnostic plots and tests*

Description

Generate regression diagnostic plots and tests for linear regression models.

Usage

```
diagnosticPlots(data, y, x, covar)
```

Arguments

data	The dataset with the variables of interest
y	The dependent or outcome variable (that is, the y in the regression equation)
x	The independent, exposure, or predictor variable (that is, the x in the regression equation)
covar	The variables selected as to condition or adjust for the y and x relationship, also known as the confounding variables

Details

This function runs a linear regression on the specified variables and generates diagnostics based on the regression. Basic diagnostics include checking the normality of the residuals, assessing outliers, influence and Cook's D, and multicollinearity. Several tests have been commented out, though they can be uncommented if desired (edit the function to output these if desired). Some of the tests I don't fully understand how to interpret them, but as I learn more I will probably know. This function relies on **MASS** and **gplots**.

Value

Outputs multiple plots and textplots with diagnostic information

Author(s)

Luke Johnston

heatmapCorr	<i>Correlation heatmap</i>
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Description

Generate a matrix or non-matrix style heatmap of correlation coefficients (i.e. the number of columns and rows can be different).

Usage

```
heatmapCorr(data, x, y, leg.range = c(-1, 1), levels.xlab = NULL,  
  levels.ylab = NULL, xlab = "", ylab = "", lo.color = "darkorange2",  
  hi.color = "skyblue4", rm.legend = FALSE, matrix.sty = FALSE,  
  corr.values = TRUE, leg.title.size = 8, leg.number.size = 7)
```

Arguments

<code>data</code>	Dataset that contains the variables of interest
<code>x</code>	Vector of variables that will run along the x-axis
<code>y</code>	Same as the <code>x</code> arg, but for the y-axis
<code>leg.range</code>	Range in values for the legend, between -1 and 1
<code>levels.xlab</code>	Specifies custom variable names for the x-axis (it needs to be a list object)
<code>levels.ylab</code>	Same as the <code>levels.xlab</code> arg, but for the y-axis
<code>xlab</code>	Sets the label for the x-axis
<code>ylab</code>	Same as <code>xlab</code> , but for the y-axis
<code>lo.color</code>	Color of negative correlation coefficients
<code>hi.color</code>	Color of positive correlation coefficients
<code>rm.legend</code>	In development. Goal is it will remove the legend
<code>matrix.sty</code>	Select whether the heatmap will be a matrix style (the same variable on the x- and y- axis) or non-matrix (different variables on both axes)
<code>corr.values</code>	Set whether to have the correlation values on the heatmap, or if it will be blank. Value is either TRUE or FALSE
<code>leg.title.size</code>	Size of the font in the legend title
<code>leg.number.size</code>	Size of the numbers in or near the legend

Details

This function takes two arguments, the `x` variables and the `y` variables, and generates a heatmap from the variables. A correlation matrix is computed from the data, melted (reshape package), and input into ggplot2 to generate a heatmap. The output is the correlations and the plot object.

Dependencies are: **reshape2** and **ggplot2**

Value

Outputs a plot object

Author(s)

Luke Johnston

Examples

```
xlabs <- list("X Name" = "x1", "X Name Two!" = "x2" ... )
ylabs <- list("Y Name" = "y1", "Y Name Two!" = "y2" ... )
df <- data.frame(replicate(10, sample(0:1, 1000, rep=TRUE)))
plot <- heatmap.corr(data = df, x = 1:3, y = 4:10,
                     levels.xlab=xlabs, levels.ylab=ylabs)
print(plot)
```

jitterBoxplot	<i>Univariate jittered boxplot</i>
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Description

Generates a boxplot of variables on one axis with raw values "jittered" as dots underneath. The variables need to represent a similar concept or have the same units for the plot to make sense.

Usage

```
jitterBoxplot(subset.ds, dot.size = 2, dot.colour = "grey50",
  custom.var.names = NULL, xlab = NULL, ylab = NULL)
```

Arguments

subset.ds	The dataset that only contains the series of variables that will be plotted
dot.size	Size of the dot for <code>geom_jitter</code>
dot.colour	Color of the dots for <code>geom_jitter</code>
custom.var.names	List object that contains the custom (alternative) variable names for the variable (column) names you passed into the function
xlab	The x-axis label
ylab	The y-axis label

Details

This function is useful for exploring the distribution of a series of variables that share a common unit, such as kilogram. The values for each variable are plotted as jittered dots with a boxplot of the distribution layered on top of the dots. The function takes a subsetting dataset that contains only the series of variables that share a common unit. The output object is the plot. This function depends on **ggplot2** and **reshape2**.

Value

Outputs the plot object

Author(s)

Luke Johnston

Examples

```
varnames <- list("Name of X1" = "X1", "Name of X2" = "X2", ...)
df <- subset(ds, VN == 0, select=nefa)
names(df)
jitter.boxplot(df, xlab="Concentration (nmol/mL)",
  ylab="Fatty acid", custom.var.names=varnames)
```

multiPlot	<i>Multiple plots on page</i>
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Description

Lay out multiple ggplots on one frame or page.

Usage

```
multiPlot(..., plotlist = NULL, file, cols = 1, layout = NULL)
```

Arguments

...	Where the ggplot objects are placed to be laid out on the graph grid
plotlist	Can be used in place of the ... argument by specifying the ggplot objects as a list object
file	Not sure what this is used for
cols	Number of columns for the layout. For example cols=2 provides two columns and with four ggplot objects, the resulting output would be a 2 by 2 graphic
layout	A matrix that indicates the plot grid layout. For example, if layout = matrix(c(1, 2, 3, 3), nrow = 2, byrow = TRUE) the result would have plot 1 in the upper left, plot 2 in the upper right, and plot 3 would be go across the bottom

Details

This function, which was from <http://www.cookbook-r.com/Graphs>, is used to lay out several ggplot objects onto one frame or pdf page. For instance, you can have 3 plots on a page, one going vertically across the top, the other two in each corner on the bottom. This function makes up for the difficulty **ggplot2** has with outputting multiple plots on one grid. This function depends on **grid**.

Author(s)

Cookbook R

plotVisreg	<i>Visualizing adjusted linear regression models</i>
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Description

Generates plots of a linear regression model which includes confounding variables.

Usage

```
plotVisreg(data, y, x, covar, ylabel = x, xlabel = y, ...)
```

Arguments

data	Dataset with the variables of interest
y	The dependent or outcome variable in the regression equation
x	The independent or exposure variable in the regression equation
covar	The confounding variables, that is the variables being adjusted for
ylabel	The y-axis label
xlabel	The x-axis label
...	Other options. In development

Details

This function runs a linear regression on the specified variables and plots the partial residuals. This allows for visualizing the relationship between the outcome and the exposure, after adjusting for confounders. A linear slope is plotted through the partial residuals, with a confidence interval band around it. The output is a plot. This function depends on **visreg**.

Value

Outputs a plot of the regression model

Author(s)

Luke Johnston

rstatsToolkit	<i>rstatsToolkit</i> .
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Description

rstatsToolkit.

smoothPlot	<i>Smooth plot</i>
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Description

In development ..

Usage

```
smoothPlot(x, y, data, ...)
```

Arguments

x
y
data
...

Details

.. content for details ..

Value

Outputs a plot

Author(s)

Luke Johnston

smoothTimePlot

Smooth time plots

Description

Generates a plot of two variables, particularly for longitudinal data. The x-axis is typically the time variable.

Usage

```
smoothTimePlot(dsn, x, y, id, smooth.type = "loess", ylab = y, xlab = x)
```

Arguments

dsn	The dataset name.
x	Specify the variable for the x-axis. Must be either in quotes or as a number.
y	As with x, but for the y-axis.
id	Specify the ID variable for longitudinal or repeated measures type data.
smooth.type	Set the smoothing method, which includes "loess", "lm", "rlm".
ylab	Optionally set the y-axis label.
xlab	Optionally set the x-axis label.

Details

This function creates a plot with smooth lines, typically using the loess method, along with the means of a discrete time variable (eg. time 0, 1, 2, 3 etc).

Dependencies are: **ggplot2**

Value

Outputs a ggplot object

Author(s)

Luke Johnston

summarySE

Summarize means and standard errors of the mean

Description

Calculates the sample size, mean, standard deviation, standard error of the mean, and the confidence interval of specified variables.

Usage

```
summarySE(data = NULL, measurevar, groupvars = NULL, na.rm = FALSE,  
  conf.interval = 0.95, .drop = TRUE)
```

Arguments

data	A dataset (dataframe) that contains the values to be summarized
measurevar	The name of a column that contains the variable to be summarized
groupvars	A vector containing names of columns that contain grouping variables
na.rm	A binary (boolean) response that indicates whether to ignore missing (NA) data
conf.interval	Percent range of the confidence interval

Details

I took this function on 2014-01-21 from the website <http://www.cookbook-r.com/Graphs>. It basically summarizes the provided data by giving count, mean, standard deviation, standard error of the mean, and confidence interval (default 95). The dependencies are **plyr**

Value

Outputs a dataframe that contains the summarized statistics (means, etc.)

Author(s)

Cookbook R

themeWhite	<i>Custom white ggplot theme</i>
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Description

Creates a white, simple theme for **ggplot2** objects

Usage

```
themeWhite()
```

Details

The default **ggplot2** theme is decent for most purposes, but is visually unappealing. This function aims to correct that by setting the theme to something more similar to the default in the base R plot package. The function depends on **ggplot2**.

Author(s)

Luke Johnston

Examples

```
## This creates a white theme  
themeWhite()
```

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