Package 'exstat'

December 13, 2022

Title Designed for the statistical description of data
Version 1.0.0
Description It is designed for the statistical description of data. This module provides functions for creating the basic descriptive statistics table, testing normality distribution of variable, screening extreme values, comparing the size between/among groups, and calculating correlation coefficient between variables. The visualization of the statistical results are also provided.
License GPL (>= 3)
Encoding UTF-8
Roxygen list(markdown = TRUE)
RoxygenNote 7.2.2
Imports httr,vroom,ggplot2,readxl,gridExtra,gt
NeedsCompilation no
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InitStat

Initialize ExpoStat Module

Description

The first step to start ExpoStat Module

Usage

InitStat()

Value

An R6 class object.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
```

LoadStat

Upload data file for ExpoStat Module

Description

Upload data file for ExpoStat Module

Usage

```
LoadStat(PID, UseExample = "default", DataPath = NULL, VocaPath = NULL)
```

Arguments

PID	chr. Program ID. It must be the same with the PI	D generated by InitStat.
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UseExample chr. Whether uses example data for analyses, available option include "exam-

ple#1" for using example data1 and "default" for using data.

DataPath chr. Input file directory, e.g. "D:/test/expostat_data.xlsx". It should be noted

that the slash symbol is "/", not "\".

VocaPath chr. Input file directory, e.g. "D:/test/expostat_voca.xlsx". It should be noted

that the slash symbol is "/", not "\".

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Value

A list object containing imported data.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
```

StatComp

Size comparison between groups

Description

Size comparison between groups

"nejm".

Usage

```
StatComp(PID, OutPath="default", Group, Task = "mean",
  Vars, VarsBy, Method = "wilcox", Layout = "density", Brightness = "dark" ,
  Palette = "default1")
```

PID	chr. Program ID. It must be the same with the PID generated by InitStat.
OutPath	chr. Output file directory, e.g. "D:/ExpoStat/StatComp". It should be noted that the slash symbol is "/", not "\".If "default", the current working directory will be set.
Group	lgl. Whether to separate dataset into train and test data for normality test. The default is "TRUE".
Task	chr. Comparison task. At present, only the mean comparison is available.
Vars	chr. Target variables used for modelling. It should be noted that there is fixed format for the entering characters separated with "," and without space. The default values is "all" (all variables are included).
VarsBy	chr. Variable used to group the observation for size comparison.
Method	chr. Comparison method. At present, only "wilcox" (Wilcoxon rank sum test) is available.
Layout	chr. Visualization layout. Available values include "column.points", "density".
Brightness	chr. Visualization brightness. Available values include "light" and "dark".
Palette	chr. Visualization palette. Available values include "default1", "default2", "default3" and 5 journal option including "cell", "nature", "science", "lancet",

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Value

A list object containing the results of size comparisons between groups for variables and visualization of the results.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatComp(PID=res$PID, Group = T, Task = "mean", Vars = "X5,X6,X7,X8,X9",
  VarsBy = "Y1", Method = "wilcox", Layout = "density", Brightness = "dark",
  Palette = "default1")
```

StatCorr

Correlation analysis between variables

Description

Correlation analysis between variables

Usage

```
StatCorr(PID, OutPath="default", Group, VarsX, VarsY, VarsBy,
   Method = "spearman", Layout= "bubble", Brightness = "dark", Palette = "default1")
```

PID	chr. Program ID. It must be the same with the PID generated by InitStat.
OutPath	chr. Output file directory, e.g. "D:/ExpoStat/StatCorr". It should be noted that the slash symbol is "/", not "\".If "default", the current working directory will be set.
Group	lgl. Whether to separate dataset into train and test data for normality test. The default is "TRUE".
VarsX	chr. Target variables used for modelling. It should be noted that there is fixed format for the entering characters separated with "," and without space. The default values is "all.x" (all variables are included).
VarsY	chr. Target outcome variables used for correlation analysis.
VarsBy	chr. Variable used to group the observation for correlation analysis.
Method	chr. Method for orrelation analysis. Available values include "spearman" (Spearman's rank correlation analysis) and "pearson" (Pearson correlation analysis).
Layout	chr. Visualization layout. Available values include "heatmap", "bubble", "matrix".

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Brightness chr. Visualization brightness. Available values include "light" and "dark".

Palette chr. Visualization palette. Available values include "default1", "default2", "default3" and 5 journal option including "cell", "nature", "science", "lancet", "neim".

Value

A list object containing the results of correlation analysis between variables and visualization of the results.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatCorr(PID = res$PID, Group = T, VarsX = "X5,X6,X7,X8,X9", VarsY = "Y1",
  VarsBy = "Y1", Method = "pearson", Layout = "bubble", Brightness = "dark",
  Palette = "nature")
```

StatDesc

Variable description

Description

Variable description

Usage

```
StatDesc(PID, OutPath="default", Group, Vars, VarsBy, Layout="box",
    Brightness="light", Palette="default1")
```

PID	chr. Program ID. It must be the same with the PID generated by InitStat.
OutPath	chr. Output file directory, e.g. "D:/ExpoStat/StatDesc". It should be noted that the slash symbol is "/", not "\".If "default", the current working directory will be set.
Group	lgl. Whether to separate dataset into train and test data for normality test. The default is "TRUE".
Vars	chr. Target variables used for modelling. It should be noted that there is fixed format for the entering characters separated with "," and without space. The default values is "all" (all variables are included).
VarsBy	chr. Variable used to group the observation for size description.

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Layout chr. Visualization layout. Available values include "box", "violin".

Brightness chr. Visualization brightness. Available values include "light" and "dark".

Palette chr. Visualization palette. Available values include "default1", "default2" and 5 journal option including "cell", "nature", "science", "lancet", "nejm".

Value

A list object containing the results of variable description for continuous and discrete variables respectively and visualization of the continuous variables.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatDesc(PID = res$PID, Group = T, Vars = "C1,C2,X5,X6,X7,X8,X9",
  VarsBy = NULL, Layout = "box", Brightness = "dark", Palette = "default2")
```

StatExtre

Extreme value calculation

Description

Extreme value calculation

Usage

```
StatExtre(PID, OutPath="default", Group, Vars, LimitLow = 0.025,
   LimitUpper = 0.975, Layout = "column.points", Brightness = "light",
   Palette = "default2")
```

PID	chr. Program ID. It must be the same with the PID generated by InitStat.
OutPath	chr. Output file directory, e.g. "D:/ExpoStat/StatExtre". It should be noted that the slash symbol is "/", not "\".If "default", the current working directory will be set.
Group	lgl. Whether to separate dataset into train and test data for normality test. The default is "TRUE".
Vars	chr. Target variables used for modelling. It should be noted that there is fixed format for the entering characters separated with "," and without space. The default values is "all" (all variables are included).
LimitLow	num. Lower limit ratio to screen the small extreme values located from 0 to this lower limit of the target variables.

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LimitUpper num. Upper limit ratio to screen the large extreme values located from this lower limit to 1 of the target variables.

Layout chr. Visualization layout . Available values include "column.points", "heatmap".

Brightness chr. Visualization brightness . Available values include "light" and "dark".

Palette chr. Visualization palette . Available values include "default1", "default2",

"default3" and 5 journal option including "cell", "nature", "science", "lancet",

"nejm".

Value

A list object containing the results of extremum for variables and visualization of the results.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatExtre(PID = res$PID, Group = T, Vars = "X5,X6,X7,X8,X9",
  LimitLow = 0.025, LimitUpper = 0.975, Layout = "column.points",
  Brightness = "dark", Palette = "default2")
```

StatNorm

Normality test for numeric variables

Description

Normality test for numeric variables

Usage

```
StatNorm(PID, OutPath="default", Group, Vars, Method = "shapiro.test", Layout = "rose.chart",
Brightness = "light", Palette = "default1")
```

Arguments

PID chr. Program ID. It must be the same with the PID generated by InitStat.

OutPath chr. Output file directory, e.g. "D:/ExpoStat/StatNorm". It should be noted that

the slash symbol is "/", not "\".If "default", the current working directory will be

set.

Group lgl. Whether to separate dataset into train and test data for normality test. The

default is "TRUE".

Vars chr. Target variables used for modelling. It should be noted that there is fixed

format for the entering characters separated with "," and without space. The

default values is "all" (all variables are included).

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Method	chr. Normality test method. Only "shapiro.test" method is available at present.
Layout	chr. Visualization layout. Available values include "column", "column.points", "rose.chart", and "density".
Brightness	chr. Visualization brightness. Available values include "light" and "dark".
Palette	chr. Visualization palette. Available values include "default1", "default2", "default3" and 5 journal option including "cell", "nature", "science", "lancet", "nejm".

Value

A list object containing the results of normality test for variables and visualization of the results.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatNorm(PID=res$PID, Group = T, Vars = 'X5,X6,X7,X8,X9',
  Method = "shapiro.test",Layout = "rose.chart" , Brightness = "dark",
  Palette = "default3")
```

StatTable1

Create Table 1 for for different epidemilogical study designs

Description

Create Table 1 for different epidemilogical study designs

Usage

```
StatTable1(PID, OutPath="default", EpiDesign = "cohort",
    Group, VarsY, VarsC, Missing = "ifany")
```

PID	chr. Program ID. It must be the same with the PID generated by InitStat.
OutPath	chr. Output file directory, e.g. "D:/ExpoStat/StatTable1". It should be noted that the slash symbol is "/", not "\".If "default", the current working directory will be set.
EpiDesign	chr. Research types provided for users, include "cohort", "case-control", "cross-section".
Group	lgl. Whether to separate dataset into train and test data for creating Table 1. The default is "TRUE".
VarsY	chr. Outcome variable used for modelling. Only one variable can be entered.

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VarsC chr. Covariate variables needing further statistical test. It should be noted that

there is fixed format for the entering characters separated with "," and without space. The defaults value is all covariate variables listed in the data file, which

can be entered with "all.c".

Missing chr. Counts of missing values in the table, available options include are "no"

(never display missing values), "ifany" (only display if any missing values), and

"always" (includes missing count row for all variables). Default is "ifany".

Value

A list object containing standardized table 1.

Author(s)

Yanqiu Feng, Bin Wang (corresponding author)

Examples

```
res = InitStat()
  res1 = LoadStat(PID = res$PID, UseExample = "example#1")
  res2 = StatTable1(PID = res$PID, EpiDesign = "cohort" ,
  Group = 'T', VarsY = "Y1", VarsC = "C1,C2,C3,C4,C5,C6",
  Missing = "ifany")
```

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