Package 'doudpackage'

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Description
Creates the ``table one" of biomedical papers. Fill it with your data and the name of the vari
able wich you'll make the group(s) out of and it will make univariate bivariate analy-

able wich you'll make the group(s) out of and it will make univariate, bivariate analy sis and parse it into HTML.

It also allows you to visualise all your data with graphic representation. It relies on kableExtra.

License GPL (>= 3) **Encoding** UTF-8 **Roxygen** list(markdown = TRUE) RoxygenNote 7.2.3 Imports dplyr, kableExtra, methods, purrr, stats, stringi, tibble, tidyr NeedsCompilation no

Title Create elegant table 1 in HTML/LaTeX for biostatistics

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R topics documented:

Version 2.0

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anaBiv

anaBiv generic function

Description

Generic function of anaBiv which gives bivariate analysis according to group anaBiv data.frame function which gives bivariate analysis according to group anaBiv data.frame function which gives bivariate analysis according to group

Usage

```
anaBiv(var, group, ...)
## S4 method for signature 'listVar,character'
anaBiv(var, group, ...)
## S4 method for signature 'data.frame,character'
anaBiv(var, group, ...)
```

Arguments

var listVar object or data.frame
group Variable to make subgroups with
... digits.p can be specified as descTab

Value

A list of VarGroup object or data.frame

descTab

Generic function to create a table of descriptive analysis of a dataset

Description

This function allows you to display all together all univariate analysis (median/mean; IQR/SD; proportions) and bivariates analysis (wilcoxon, chisq or fisher). The univariate analysis can be sub-grouped by a variable of interest of n levels. Approriate statistics test will be applied

descTab 3

Usage

```
descTab(
  data,
  group = NULL,
  quanti = TRUE,
  quali = TRUE,
  na.print = FALSE,
  pvalue = TRUE,
  digits.p = 3L,
  digits.qt = 1L,
  digits.ql = 1L,
  normality = "normal"
)
```

Arguments

data	A datasaset. Needs to be a data.frame/tibble object			
group	Optional. The name of the variable to make sub-groups comparisons.			
quanti, quali, na.print, pvalue				
	Bool. If false, won't display quantitative/qualitative/Missing values/pvalues variable results			
digits.p	Integer. Significant digits for p value			
digits.qt	Integer. Significant digits for mean/median, SD/IQR			
digits.ql	Integer. Significant digits for proportions			
normality	One of "assess", "normal", "manual", "not normal". See details			

Value

A S4 objects containing the main table accessible by ["table"] subscript.

Examples

```
data(iris)
library(stringi)
iris$fact_1<-as.factor(as.character(sample(1:5, 150, replace = TRUE)))
n_na<-sample(1:150, 30)
iris[n_na, "fact_1"]<-NA
iris$fact_2<-as.factor(as.character(sample(1:2, 150, replace = TRUE)))
n_na<-sample(1:150, 10)
iris[n_na, "fact_2"]<-NA
iris$fact_3<-as.factor(as.character(stri_rand_strings(150, 1, '[A-B]')))
iris$num<-runif(150, min = 0, max = 100)
n_na<-sample(1:150, 5)
iris[n_na, "num"]<-NA
iris_test<-descTab(iris, group = "Species", na.print = TRUE)</pre>
```

ft_parse

ft_ana_biv

This function is depreciated, please use anaBiv(). anaBiv()

Description

This function is depreciated, please use anaBiv(). anaBiv()

Usage

```
ft_ana_biv(...)
```

Arguments

... None

ft_desc_tab

This function is depreciated, please use anaBiv(). descTab()

Description

This function is depreciated, please use anaBiv(). descTab()

Usage

```
ft_desc_tab(...)
```

Arguments

... None

ft_parse

This function is depreciated, please use anaBiv(). parseClassFun()

Description

This function is depreciated, please use anaBiv(). parseClassFun()

Usage

```
ft_parse(...)
```

Arguments

... None

```
initialize,parseClass-method

S4 class initialization function
```

Description

Initialization function for parseClass object initialize, parseClass-method()

Usage

```
## S4 method for signature 'parseClass'
initialize(
   .Object,
   table,
   group,
   pvalue,
   na.print,
   quanti,
   quali,
   var_list,
   data,
   digits.qt,
   digits.ql
)
```

Arguments

Value

```
parseClass object
```

initialize, Var-method S4 class initialization function

Description

Initialization function for Var initialize, Var-method()

Usage

```
## S4 method for signature 'Var'
initialize(.Object, name, type, normal)
```

Arguments

.0bject Object to be initialized

name A character taking name of the variable

type A character taking name of the variable type normal Logical, if variable, is numeric; is it normal

Value

Var Object

```
initialize, {\tt VarGroup-method}\\ S4\ class\ initialization\ function
```

Description

Initialization function for VarGroup initialize, VarGroup-method()

Usage

```
## S4 method for signature 'VarGroup'
initialize(
   .Object,
   x,
   group_var,
   pvalue,
   parsed_name,
   value,
   missing.value,
   missing.value.name
```

listVar-class 7

Arguments

.0bject Object to be initialized

x A Var object

group_var The subgroup for whichproportions, mean/sd were calculated and missing val-

ues

pvalue The calculated pvalue

parsed_name The name of the variable parsed with the n (%), mean (SD)

value The values calculated parsed

missing.value Missing values numbers and proportions n (%)

missing.value.name

Missing values concatenate with the leve of the variable if it factor

Value

VarGroup object

listVar-class

S4 class

Description

A class of list of Var object

Slots

List a list of Var

parseClass

S4 class initialization function

Description

Initialization function for parseClass object initialize, parseClass-method()

Usage

```
parseClass(
  table,
  group,
  pvalue,
  na.print,
  quanti,
  quali,
  var_list,
  data,
  digits.qt,
  digits.qt,
)
```

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Arguments

Value

parseClass object

parseClass-class S4 class

Description

A S4 class containing all the information needed for parsClassFun the missing values and the group for which it was calculated

Slots

```
table The result of descTab
group The variable from which to make subgroups
pvalue,na.print,quanti,quali Values from descTab descTab()
var_list An object of listVar listVar-class()
data The dataset provided in descTab
digits.qt,digits.ql As provided in descTab
```

parseClassFun

Make the LaTeX/HTML table. Generic function

Description

Make the LaTeX/HTML table. Generic function

Usage

```
parseClassFun(
  table,
  col.order = NULL,
  levels_to_keep = NULL,
  group_rows_labels = NULL)
```

Arguments

table The output of descTab, an S4 object.

col.order Optional. A vector containing the column order. If set, must contains at least all

levels of group. Three columns created are "var", "Total", and "pvalue" which

can be present in the vector

levels_to_keep Optional, named list. If the variable is binary, which level to keep. Default is the

last level of levels(variable). Must be as: list("variable name" = "level to keep").

group_rows_labels

Optional, named list. Create row labels in order to regroup them. Must be as

list("label" = c("var1", "var2), "label2" = c("var3", "var4")).

Value

An HTML/LaTex file which can be used directly in Rmarkdown and copy paste

parseClassFun,parseClass-method

Make the LaTeX/HTML table

Description

This functions takes the S4 output of descTab to create an HTML parsed table

Usage

```
## S4 method for signature 'parseClass'
parseClassFun(
  table,
  col.order = NULL,
  levels_to_keep = NULL,
  group_rows_labels = NULL
)
```

Arguments

table The output of descTab, an S4 object.

col.order Optional. A vector containing the column order. If set, must contains at least all

levels of group. Three columns created are "var", "Total", and "pvalue" which

can be present in the vector

levels_to_keep Optional, named list. If the variable is binary, which level to keep. Default is the

last level of levels(variable). Must be as: list("variable name" = "level to keep").

group_rows_labels

Optional, named list. Create row labels in order to regroup them. Must be as

list("label" = c("var1", "var2), "label2" = c("var3", "var4")).

Value

An HTML/LaTex file which can be used directly in Rmarkdown and copy paste

Var-class

Examples

```
data(iris)
library(stringi)
iris$fact_1<-as.factor(as.character(sample(1:5, 150, replace = TRUE)))
n_na<-sample(1:150, 30)
iris[n_na, "fact_1"]<-NA
iris$fact_2<-as.factor(as.character(stri_rand_strings(150, 1, '[A-B]')))
iris$num<-runif(150, min = 0, max = 100)
n_na<-sample(1:150, 5)
iris[n_na, "num"]<-NA
iris_test<-descTab(iris, group = "Species", na.print = TRUE)
testParse<-parseClassFun(iris_test, levels_to_keep = list("fact_2" = "A"),
group_rows_labels = list("Size" = c("Petal.Length", "Petal.Width"),
"My_f" = c("num", "fact_2")))</pre>
```

Var

S4 class initialization function

Description

Initialization function for Var initialize, Var-method()

Usage

```
Var(name, type = "", normal = TRUE)
```

Arguments

name A character taking name of the variable
type A character taking name of the variable type
normal Logical, if variable, is numeric; is it normal

Value

Var Object

Var-class

S4 class

Description

A S4 class containing name, type and normality assessment of variable

Slots

```
name A character taking name of the variable
type A character taking name of the variable type
normal Logical, if variable, is numeric; is it normal
```

VarGroup-class 11

VarGroup-class

S4 class

Description

A S4 class containing Var initialize, Var-method() It also contains the pvalue, the parsed value the missing values and the group for which it was calculated

Slots

```
group_var The subgroup for whichproportions, mean/sd were calculated and missing values pvalue The calculated pvalue parsed_name The name of the variable parsed with the n (%), mean (SD) value The values calculated parsed missing.value Missing values numbers and proportions n (%) missing.value.name Missing values concatenate with the level of the variable if it factor
```

[,parseClass-method

Method to access S4 Var elements

Description

Method to acess parseClass initialize, parseClass-method() elements by name

Usage

```
## S4 method for signature 'parseClass' x[i]
```

Arguments

x : Object

i : Element name

Value

object

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[,Var-method

Method to access S4 Var elements

Description

Method to access Var elements by name

Usage

```
## S4 method for signature 'Var' x[i]
```

Arguments

x : object i : value

Value

object of Var

[,VarGroup-method

Method to access S4 Var elements

Description

Mathod to access VarGroup initialize, VarGroup-method() elements by name

Usage

```
## S4 method for signature 'VarGroup' x[i]
```

Arguments

x : object i : value

Value

object element

```
[<-,parseClass,character-method
```

Method to modify S4 Var elements

Description

 $Method \ to \ modify \ parseClass \ initialize, parseClass-method() \ elements \ by \ name$

Usage

```
## S4 replacement method for signature 'parseClass,character' x[i] \leftarrow value
```

Arguments

x : Object

i : Element name value : Value to be added

Value

object

[<-,Var,ANY-method</pre>

Method to access S4 Var elements

Description

Method to modify Var elements by name

Usage

```
## S4 replacement method for signature 'Var,ANY' x[i] \leftarrow value
```

Arguments

x : object

i : Element name value : Value to be added

Value

object

```
[<-,VarGroup,ANY-method</pre>
```

Method to access S4 Var elements

Description

Method to modify VarGroup initialize, VarGroup-method() elements by name

Usage

```
## S4 replacement method for signature 'VarGroup, ANY' x[i] \leftarrow value
```

Arguments

x Object

i Element name value Value to be added

Value

object

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