

Package ‘StudyDataTools’

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Type Package

Title X

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Description Data

Remotes tidyverse/magrittr

Depends RODBC,

ggplot2,

sqldf,

lattice,

printr,

knitr,

reshape2,

rlist,

devtools,

haven,

sas7bdat

License GPL (>= 2)

LazyLoad yes

LazyData true

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automaticdatafConnect *get data about a file on the server.*

Description

get data about a file on the server.

Usage

```
automaticdatafConnect(tablename, folder = getwd(), schema = NULL,
  dicoT = NULL, splitvar = NULL, Connect = NULL, Connectf = NULL,
  alwaysexclude = NULL)
```

Value

a list

drop_last	<i>Drop last marginpos (cuts everything after last after "_")</i>
-----------	---

Description

Drop last marginpos (cuts everything after last after "_")

Usage

```
drop_last(x)
```

Arguments

x a vector of character strings

Details

if x is "aa.x_a_1_f_1" returns "a_1_f_1"

Value

a vector of character strings

Examples

```
drop_last("AA.char1_La_Ld_Lrn1")
```

exportRpackagedata_to_csvlibrary	<i>export all tables from an installed R data package to sas7bdat data files in a given library</i>
----------------------------------	---

Description

export all tables from an installed R data package to sas7bdat data files in a given library

Usage

```
exportRpackagedata_to_csvlibrary(package, path_to_export_to = getwd(),
  tables = NULL, zip = TRUE)
```

Arguments

package	package to be transformed
path_to_export_to	path to export the data
tables	tables to extract, if NULL, all tables.
zip	(should the files be zipped (requires utils package))

Value

nothing

Examples

```
exportRpackagedata_to_csvlibrary("datasets")
```

```
exportRpackagedata_to_sas7bdatlibrary
```

export all tables from an installed R data package to sas7bdat data files in a given library

Description

export all tables from an installed R data package to sas7bdat data files in a given library

Usage

```
exportRpackagedata_to_sas7bdatlibrary(package, path_to_export_to = getwd(),  
  tables = NULL)
```

Arguments

package	package to be transformed
SAS_library_path	path to export the data

Value

nothing

Examples

```
exportRpackagedata_to_sas7bdatlibrary("datasets")
```

```
exportRpackagedata_to_savlibrary
```

export all tables from an installed R data package to sav data files in a given library

Description

export all tables from an installed R data package to sav data files in a given library

Usage

```
exportRpackagedata_to_savlibrary(package, path_to_export_to = getwd(),
  tables = NULL, zip = TRUE)
```

Arguments

package	package to be transformed
path_to_export_to	path to export the data
tables	tables to extract, if NULL, all tables.
zip	(should the files be zipped (requires utils package))

Value

nothing

Examples

```
exportRpackagedata_to_csvlibrary("datasets")
```

```
GeneralReversetransposefunction
```

General Reverse Transpose function

Description

General Reverse Transpose function

Usage

```
GeneralReversetransposefunction(TtableA, key)
```

Arguments

key	A list of variables (columns of the transposed table)
table	A dataframe

Value

A list: first element of the list is a dataframe, the transposed version of the orioginal table. Second element is a key to allow back transposition

Examples

```
data(tableA);data(TtableA);data(XKA);key<-XKA$key
RtableA=GeneralReversettransposefunction(TtableA,key)
ordertableA <-do.call(order,tableA[c(id1,id2)])
orderRtableA<-do.call(order,RtableA[c(id1,id2)])
identical(nrow(tableA),nrow(RtableA))
identical(lapply(tableA,class),lapply(RtableA,class))
identical(tableA[ordertableA,],RtableA[orderRtableA,])
identical(names(tableA),names(RtableA))
all (lapply(names(tableA),function(x){identical(tableA[ordertableA,x],RtableA[orderRtableA,x])}))
```

Generaltransposefunction
<i>General Transpose function</i>

Description

General Transpose function

Usage

```
Generaltransposefunction(tableA, id1, id2,
  origin = deparse(substitute(tableA)))
```

Arguments

- id1 A list of variables (rows)
- id2 A list of variables (columns of the transposed table), id2 can contain as a last element the strint "rn", if the variable rn is an index for the cells formed by the variables listed first in id2
- table A dataframe

Value

A list: first element of the list is a dataframe, the transposed version of the orioginal table. Second element is a key to allow back transposition

Examples

```
tableA<-sampledata(TRUE)
id1=c("id1a","id1b")
id2=c("id2a","id2b")
TtableA<-Generaltransposefunction(tableA,id1,id2)
```

Generaltransposefunctionsimple
<i>Simple General Transpose function</i>

Description

Simple General Transpose function

Usage

Generaltransposefunctionsimple(tableA, id1, id2)

Arguments

- tableA A dataframe
- id1 A list of variables (rows)
- id2 A list of variables (columns of the transposed table)

Value

A data frame

Examples

```
tableA<-sampledata(TRUE)
id1=c("id1a","id1b")
id2=c("id2a","id2b")
TtableA<-Generaltransposefunctionsimple(tableA,id1,id2)
```

get_cell	<i>get cell without the row number</i>
----------	--

Description

get cell without the row number

Usage

get_cell(x, iscellrn = FALSE, iscell = FALSE)

Arguments

- x a vector of character strings

Details

if x is "aa.xoijj_a_1_f_1_" returns "a_1_f"

Value

a vector of character strings

Examples

```
get_cell("aa.x_1_2_3_4")#default
get_cell("1_2_3", TRUE)
get_cell("1_2_3", FALSE, TRUE)
unique(Tsampledata(TRUE)$variables))
unique(get_cell(Tsampledata(FALSE)$variables))
```

get_cellrn	<i>Get cell and row number</i>
------------	--------------------------------

Description

Get cell and row number

Usage

```
get_cellrn(x)
```

Arguments

x a vector of character strings

Details

if x is "aa.x_a_1_f_1" returns "a_1_f_1"

Value

a vector of character strings

Examples

```
get_cellrn("AA.char1_La_Ld_Lrn1")
data(TtableA);
unique(get_cellrn(names(TtableA)))
#Second example: no transposing variables
data(TtableB);data(XKB)
unique(get_cellrn(names(XKB)))
```

get_cellXXgroup	<i>Get cell group</i>
-----------------	-----------------------

Description

Get cell group

Usage

```
get_cellXXgroup(x, marginpos, iscellXX = TRUE)
```

Arguments

x	a vector of character strings
marginpos	a vector of integer

Details

if x is "a_1_f_2_aa.xoijj",marginpos=2 returns "1" if x is "a_1_f_2_aa.xoijj",marginpos=-2 returns "a_f_2" if x is "a_1_f_2_aa.xoijj",marginpos=c(1:2) returns "a_1"

Value

a vector of character strings

Examples

```
get_cellXXgroup(c("aa.x_1_2_3_4","bb.x_1_2_3_4"),2,iscellXX=FALSE)
get_cellXXgroup(c("1_2_3_4","1_2_3_4"),2:3,iscellXX=TRUE)
variables<-Tsampledata(TRUE)$variables
unique(get_cellXXgroup(variables,2,iscellXX=FALSE))
unique(get_cellXXgroup(variables,-2,iscellXX=FALSE))
get_cellXXgroup(variables[50],2,iscellXX=FALSE)
get_cellXXgroup(variables[50],-2,iscellXX=FALSE)

#Second example: no transposing variables
TK<-Tsampledata(FALSE)
unique(get_cellXXgroup(TK$variable,1,iscell=FALSE))
```

```
get_cellXXmarginscount
```

Get the number of margins for a cell

Description

Get the number of margins for a cell

Usage

```
get_cellXXmarginscount(x, iscellXX = FALSE)
```

Arguments

`x` a vector of character strings
`iscell` a boolean indicating if `x` is a variable name or a cell name.

Details

if `x` is "aa.xoijj_a_1_f_1", `cell=FALSE` returns 4" if `x` is "a_1_f_1", `cell=TRUE` returns 4"

Value

a vector of integers.

Examples

```
get_cellXXmarginscount("1_2_3_4", iscellXX=TRUE)
get_cellXXmarginscount("aa.x_1_2_3_4", iscellXX=FALSE)
data(TtableA)
unique(get_cellXXmarginscount(names(TtableA), iscellXX=FALSE))
#Second example: no transposing variables
TK<-Tsampledata(FALSE)
unique(get_cellXXmarginscount(TK$variables))
```

```
get_cellXXsplit
```

split a cell

Description

split a cell

Usage

```
get_cellXXsplit(x, marginpos = NULL, iscellXX = FALSE)
```

Arguments

x a vector of character strings
 iscell x a boolean indicating if x is a cell

Details

if x is "aa.xoiij_a_1_f_1" returns c("a","1","f","1")

Value

a vector of character strings

Examples

```
get_cellXXsplit("aa.x_1_2_3_4", iscellXX=FALSE)
get_cellXXsplit("1_2_3_4", iscellXX=TRUE)
get_cellXXsplit("1_2_3_4", 2:3, iscellXX=TRUE)
get_cellXXsplit("1_2_3_4", -(2:3), iscellXX=TRUE)
variables<-Tsampledata(TRUE)$variables
unique(get_cellXXsplit(variables, iscell=FALSE))
get_cellXXsplit(variables[50], iscell=FALSE)
get_cellXXsplit(variables[50], -(2:3), iscell=FALSE)
unique(get_cellXXsplit(variables, 2, iscell=FALSE))
#Second example: no transposing variables
TK<-Tsampledata(FALSE)
unique(get_cellXXsplit(TK$variables, iscell=FALSE))
```

get_missingind	<i>Get missing indicator for a cell or variable</i>
----------------	---

Description

Get missing indicator for a cell or variable

Usage

```
get_missingind(x, variables)
```

Arguments

x a vector of character strings

Details

if x is "a_1_f_1_aa.xoiij" returns c("a","1","f","1")

Value

a vector of character strings

Examples

```
variables<-Tsampledata(TRUE)$variables
unlist(unique(get_missingind(variables,variables)))
variables<-Tsampledata(FALSE)$variables
unlist(unique(get_missingind(variables,variables)))
```

```
get_natural.predictors
```

get variable predecessors at margin

Description

get variable predecessors at margin

Usage

```
get_natural.predictors(x, variables = x, predictors = NULL)
```

Arguments

x	a vector of character strings
variables	a vector of character strings
cells	a vector of character strings containing the potential predecessors
marginpos	a vector of integers
x	a vector of character strings

Details

if x is "a_1_f_1_aa.xoijj" returns c("a","1","f","1")

if x is "a_1_f_1_aa.xoijj" returns c("a","1","f","1")

Value

a vector of character strings

a vector of character strings

Examples

```
get_XXpredecessoratmargin(cellXXs="aa.x_1_2_3_4", refcellXXs=c("bb.x_1_2_2_4", "aa.x_1_2_2_4", "aa.x_1_1_3_4"), 2,
get_XXpredecessoratmargin(cellXXs=c("1_2_2_4", "1_2_2_4", "1_1_3_4", "1_1_3_3"), iscellXX=FALSE)
data(XKA)
cells<-unique(get_cellrn(XKA$variables))
get_XXpredecessoratmargin(cells,marginpos=1,iscellXX=TRUE)
get_XXpredecessoratmargin(cells[10],cells,1,iscellXX=TRUE)
Get natural predictors

TK<-TtableA
get_natural.predictors(x=sample(names(TtableA),5),variables=names(TtableA))
```

get_origin	<i>Get origin table</i>
------------	-------------------------

Description

Get origin table

Usage

```
get_origin(x)
```

Arguments

x a vector of character strings

Details

if x is "aa.xoijj_a_1_f_1_" returns c("aa")

Value

a vector of character strings

Examples

```
variables<-Tsampledata(TRUE)$variables
unlist(unique(get_origin(variables,variables)))
variables<-Tsampledata(FALSE)$variables
unlist(unique(get_origin(variables,variables)))
```

get_presentind	<i>get the present indicator for a cell</i>
----------------	---

Description

get the present indicator for a cell

Usage

```
get_presentind(variables, refvariables = variables,
  rns = unlist(unique(get_cellrn(refvariables))))
```

Arguments

x a vector of character strings

Details

if x is "a_1_f_1_aa.xoiij" returns c("a","1","f","1")

Value

a vector of character strings

Examples

```
get_presentind("AA.x_1_2_3_4", "AA.present_1_2_3_4")
get_presentind("AA.present_1_2_3_4", c("AA.present_1_2_3_3", "AA.present_1_2_3_4"))
variables<-Tsampledata(TRUE)$variables
variable<-"AA.present_La_La_Lrn1"
get_presentind(variable, variables)
unlist(unique(get_presentind(variables)))
variables<-Tsampledata(FALSE)$variables
unlist(unique(get_presentind(variables, variables)))
```

get_var

Get variable name

Description

Get variable name

Usage

```
get_var(x)
```

Arguments

x a vector of character strings

Details

if x is "aa.xoiij_a_1_f_1" returns "aa.xoiij"

Value

a vector of character strings

Examples

```
get_var("aa.x_1_2_3_4")
data(TtableA)
unique(get_var(names(TtableA)))
#Second example: no transposing variables
TK<-Tsampledata(FALSE)
unique(get_var(TK$variables))
```

```
get_XXpredecessoratmargin
      get cell predecessors at margin
```

Description

get cell predecessors at margin

Usage

```
get_XXpredecessoratmargin(XXs, refXXs = XXs, marginpos = NULL,
  iscellXX = FALSE)
```

Arguments

XXs	a vector of character strings
refXXs	a vector of character strings containing the potential predecessors
marginpos	a vector of integers

Details

if XXs is "aa.xoiij_a_1_f_1" and refXXs contains "aa.xoiij_a_1_e_1" and marginpos=3 returns "aa.xoiij_a_1_e_1" if XXs is "aa.xoiij_a_1_f_2" and refXXs contains "aa.xoiij_a_1_f_1" and marginpos=NULL returns "aa.xoiij_a_1_f_1" if XXs is "id1" and iscellXX=FALSE whatever refXXs returns character(0) if XXs is "" and iscellXX=FALSE whatever refXXs returns character(0) if XXs is "b_1_f_1" and iscellXX=TRUE and refXXs contains "a_1_f_1" returns "a_1_f_1"

Value

a vector of character strings

Examples

```
get_XXpredecessoratmargin(XXs="aa.x_1_2_3_4", refXXs=c("bb.x_1_2_2_4","aa.x_1_2_2_4","aa.x_1_1_3_4"),2,iscellXX=FALSE)
get_XXpredecessoratmargin(XXs=c("1_2_2_4","1_2_2_4","1_1_3_4","1_1_3_3"),iscellXX=TRUE)
get_XXpredecessoratmargin(XXs="1_1_3_4",refXXs=c("1_2_2_4","1_2_2_4","1_1_3_4","1_1_3_3"),iscellXX=TRUE)
data(XKA)
cells<-unique(get_cellrn(XKA$variables))
get_XXpredecessoratmargin(cells,marginpos=1,iscellXX=TRUE)
get_XXpredecessoratmargin(cells[10],cells,1,iscellXX=TRUE)
```

ggplot_missing	Create missing chart
----------------	----------------------

Description

Create missing chart

Usage

```
ggplot_missing(x, reordonne = FALSE)
```

Arguments

x	a dataframe
reordonne	a boolean

Value

a ggplot graph

Examples

```
library(reshape2)
library(ggplot2)
library(plyr)
library(magrittr)
X=cars
for(i in 1:40){
  X[sample(1:50,1,replace=TRUE),sample(1:2,1,replace=TRUE)]<-NA}
ggplot_missing(X,reordonne=TRUE)
ggplot_missing(X,reordonne=FALSE)createallautomaticRMD(schema="SDP")
```

ggplot_missing2	Create missing chart
-----------------	----------------------

Description

Create missing chart

Usage

```
ggplot_missing2(X, reordonne = TRUE, keep = NULL)
```


Arguments

X	a dataframe
reordonne	a boolean
keep	a boolean

Value

a ggplot graph

Examples

```
library(reshape2)
library(ggplot2)
library(plyr)
X=cars
X$year=sample(2012:2017,nrow(cars),replace=TRUE)
for(i in 1:40){
  X[sample(1:50,1,replace=TRUE),sample(1:2,1,replace=TRUE)]<-NA}
ggplot_missing2(X,keep="year")
```

missing.summary	<i>Percentage of missing for each variable</i>
-----------------	--

Description

Percentage of missing for each variable

Usage

```
missing.summary(X, info2 = NULL)
```

Arguments

X	a data frame
info2	a data frame with two variables named c("COLUMN_NAME","CONSTRAINT_TYPE")

Details

Percentage of missing for each variable of a data frame.

Value

a data frame

```
predictor.matrix.default
```

Define a default predictor matrix

Description

Define a default predictor matrix

Usage

```
predictor.matrix.default(variables)
```

Arguments

`variables` a vector of character strings

Details

Returns the lower diagonal matrix with ones.

Value

a matrix

Examples

```
variables<-Tsampledata(TRUE)$variables
predictor.matrix.default(TK$variables)
```

```
predictor.matrix.rate    predictor.matrix.rate
```

Description

`predictor.matrix.rate`

Usage

```
predictor.matrix.rate(variables, nopredictor = character(0),
  allpredictor = character(0), marginposs = integer(0))
```

Arguments

`x` a vector of character strings

Details

if x is "aa.xoiij_a_1_f_1_" returns c("a","1","f","1")

Value

a vector of character strings

runCompare	<i>Shiny App to visualize Data</i>
------------	------------------------------------

Description

Shiny App to visualize Data

Usage

```
runCompare(package1 = NULL, package2 = NULL)
```

Examples

```
package1<-NULL  
package2<-NULL  
runCompare()
```

sampladata	<i>Sample data for transposition</i>
------------	--------------------------------------

Description

Sample data for transposition

Usage

```
sampladata(transposingvariables = TRUE)
```

Arguments

transposingvariables
a boolean. If TRUE, transposing id variables are created.

Value

a data frame with id variables, numeric, factor and character variables.

SDPSYN

*General SDP function.***Description**

General SDP function.

Usage

```
SDPSYN(TtableA, asis = NULL, notpredictor = asis, replicate = 1,
       Sparameters = Sparameters.default.f(TtableA, asis, notpredictor),
       STtableA = plyr::rdply(replicate, TtableA[asis]))
```

Examples

```
TK<-Tsampledata(TRUE)
TtableA<-TK$TtableA;key=TK$key;
STtableA=as.data.frame(matrix(NA,nrow(TtableA),0))
Sparameters=Sparameters.default.f(ref.table=TtableA)
Sparameters_i<-Sparameters[[54]]
Split=Sparameters_i$split[[1]]
Split=Sparameters_i$split[[2]]
library(synthpop)
TtableA[sapply(TtableA,is.character)]<-lapply(TtableA[sapply(TtableA,is.character)],as.factor)
STtableA<-SDPSYN(TtableA)
StableA<-StudyDataTools::GeneralReversetransposefunction(STtableA,key)
```

Sparameters.default.f *Default synthetisation parameters based on variable names*

Description

Default synthetisation parameters based on variable names

Usage

```
Sparameters.default.f(ref.table, asis = NULL, notpredictor = NULL,
                      variables = Sparameters.variables.reorder.default(names(ref.table)),
                      predictors.matrix = predictor.matrix.default(variables)[!is.element(variables,
                      asis), !is.element(variables, notpredictor)], moresplits = NULL,
                      preferredmethod = "rf")
```

Arguments

a vector of character strings

Examples

```
data(TtableA)
ref.table<-TtableA
Sparameters.default.f(ref.table=TtableA)
```

```
Sparameters.variables.reorder.default
```

General Default ordering of variables for synthetisation based on name of the variable.

Description

General Default ordering of variables for synthetisation based on name of the variable.

Usage

```
Sparameters.variables.reorder.default(variables, orderwithinorigin = NULL,
  id = NULL)
```

Arguments

`variables` vector of character strings, indicating names of variables
`orderwithinorigin`
 a list, see example

Value

a list.

Examples

```
TK<-Tsampladata(TRUE)
Sparameters.variables.reorder.default(names(TK$TtableA))
#Second example: no transposing variables
TK<-Tsampladata(FALSE)
orderwithinorigin=c("AA.factor1", "AA.factor2")
Sparameters.variables.reorder.default(names(TK$TtableA),orderwithinorigin)
```

Tsampledata	<i>Transposed sample data.</i>
-------------	--------------------------------

Description

Transposed sample data.

Usage

Tsampledata(transposingvariables = TRUE)

Arguments

transposingvariables
a boolean. If TRUE, transposing id variables are created.

Details

Tsampledata(x) is Generaltransposefunction(Tsampledata(x))

Value

a data frame with id variables, numeric, factor and character variables.

TTsampledata	<i>Transposed sample data.</i>
--------------	--------------------------------

Description

Transposed sample data.

Usage

TTsampledata(transposingvariables = TRUE)

Arguments

transposingvariables
a boolean. If TRUE, transposing id variables are created.

Details

Tsampledata(x) is Generaltransposefunction(Tsampledata(x))

Value

a data frame with id variables, numeric, factor and character variables.

var.summary	Summary for each variable in table.
-------------	-------------------------------------

Description

Summary for each variable in table.

Usage

```
var.summary(X, datadic = NULL)
```

Arguments

X	a data frame
datadic	a data dictionary

Value

a list

Examples

```
data(cars)
var.summary(cars)
```

var.summaryConnect	Summary for each variable in table.
--------------------	-------------------------------------

Description

Summary for each variable in table.

Usage

```
var.summaryConnect(X, datadic = NULL)
```

Arguments

X	a data frame
datadic	a data dictionary

Value

a list

Examples

```
data(cars)  
var.summaryConnect(cars)
```


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