

# Package ‘HKRbook’

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**Type** Package

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**Version** 0.1.2

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---

checkPackages	<i>checkPackages</i>
---------------	----------------------

---

## Description

Checks if a package is installed without loading it. Returns a logical vector with TRUE or FALSE for each package checked.

## Usage

```
checkPackages(
  ...,
  add = c("highlight", "formatR", "shiny", "shinydashboard", "shinydashboardPlus", "DT")
)
```

## Arguments

...	character: name(s) of package
add	character: names of default packages to check (default: c("highlight", "formatR", "shiny", "shinydashboard", "shinydashboardPlus", "DT"))

## Value

TRUE if successful otherwise an error will be thrown

## Examples

```
checkPackages("graphics", add=NULL) # checks if 'graphics' is installed
if (interactive()) checkPackages("graphics") # checks if 'graphics', 'shiny', ... are installed
```

---

distributionParams      *distributionParams*

---

### Description

Computes approximate distribution parameters for the binomial, hypergeometric, Poisson, Exponential and normal distribution for a given mean (and standard deviation). With the sample and the population size the computation can be influenced.

### Usage

```
distributionParams(mean, sd, n = 30, N = 60)
```

### Arguments

mean	numeric: mean
sd	numeric: standard deviation (only used for the normal distribution)
n	integer: sample size (default: 30)
N	integer: population size (default: 60)

### Value

a list of parameters for each distribution

### Examples

```
# Compute approx. paramaters for a binomial distribution
distributionParams(mean=30*0.5, sd=sqrt(30*0.5*0.5))
```

---

gettext      *gettext*

---

### Description

Returns a translation from loaded PO-file. If the message is not found in the PO-file then original text will be returned.

### Usage

```
gettext(msg, utype = "vector")
```

### Arguments

msg	character: message(s) to translate
utype	character: how to return the translated message as vector or named list

**Value**

translated messages

**Examples**

```
msgs <- c("two.sided", "less", "greater")
gettext(msgs)
# for use in Shiny "choices"
gettext(msgs, "name")
gettext(msgs, "numeric")
```

---

hm\_cell

*hm\_cell*

---

**Description**

- `hm_cell` or `hm_index` modify a data cell format (`fmt="%s"`), value (unnamed parameter) or style (`text_align="left"`)
- `hm_col` or `hm_row` modify a row or column format (`fmt="%s"`), value (unnamed parameter) or style (`text_align="left"`)

**Usage**

```
hm_cell(x, row = NULL, col = NULL, ..., byrow = FALSE)
```

```
hm_index(x, ind, ...)
```

```
hm_title(x, ...)
```

```
hm_colmargintitle(x, ...)
```

```
hm_rowmargintitle(x, ...)
```

```
hm_total(x, ...)
```

```
hm_table(x, ...)
```

```
hm_row(x, ind, ...)
```

```
hm_col(x, ind, ...)
```

```
hm_colmargin(x, ind, ...)
```

```
hm_rowmargin(x, ind, ...)
```

```
hm_tr(x, ind, ...)
```

**Arguments**

x	html_matrix object
row	integer: row(s) to access
col	integer: column(s) to access
...	elements to change
byrow	logical: order indices by row or column (default: FALSE)
ind	integer vector or matrix: access various (row and columns) elements (first column: row, second column: column)

**Value**

modified html\_matrix object

**Examples**

```
l <- html_matrix(matrix(1:6, ncol=2))
# replace l[1,1] by NA
hm_cell(l, 1, 1, NA)
# replace l[1,1] by NA and set the text_align to center
hm_cell(l, 1, 1, NA, text_align="center")
# replace l[1,3] and l[2,1] by NA
rcind <- cbind(c(1,3), c(2, 1))
hm_index(l, rcind, NA)
# set a new title
hm_title(l, "new title")
# set a new row or column title
hm_row(l, 2, "row 2")
hm_col(l, 1, "col 1")
# set fmt by column or row
print(hm_cell(l, fmt=c("%.0f", "%.1f", "%.2f"), byrow=FALSE), which="fmt")
print(hm_cell(l, fmt=c("%.0f", "%.1f"), byrow=TRUE), which="fmt")
```

---

htmlTable

*htmlTable*


---

**Description**

Creates a HTML table from a two dimensional table object.

**Usage**

```
htmlTable(
  tab,
  vars = NULL,
  lines = NULL,
  cex = 1,
  title = "",
```

```

    rowsum = NULL,
    colsum = NULL,
    fmt = "%.0f",
    total = NULL,
    ...
)

```

### Arguments

tab	two dimensional table object
vars	character: names of row and column variable
lines	character: final line (default: NULL)
cex	numeric: font size (default: 1)
title	character: table title (default: '')
rowsum	character: add row sums at the right (default: NULL)
colsum	character: add column sums at the bottom (default: NULL)
fmt	character: format string for sprintf (default: "%.0f")
total	character: add the grand total at the bottom left (default: NULL)
...	further parameters given to html_matrix

### Value

html\_matrix object

### Examples

```

htab <- htmlTable(apply(Titanic,1:2,sum), c("Sex", "Class"), title="Titanic")
toHTML(htab, browser=interactive())

```

---

html\_matrix

*html\_matrix*

---

### Description

Creates from a vector, matrix, array, or table a HTML representation of it. The HTML representation has one column and row more than the data. The additional row and column are used to have a title (top left), the column names (top), and the row names (left).

You can set the style attributes (<td style="...">) via hm\_cell, hm\_title, hm\_col, and hm\_row. For example: hm\_cell(hm, 1, 1, text\_align="right") will lead to (<td style="text-align:right;">) for the cell (1,1) and any unnamed element will change the cell value. Note: since - is an operator in R, we use \_ instead. Of course, you could use "text-align="right", but I'm lazy.

**Usage**

```
html_matrix(x, ...)

## Default S3 method:
html_matrix(
  x,
  ...,
  byrow = FALSE,
  numeric = list(text_align = "right"),
  integer = list(text_align = "right"),
  char = list(text_align = "left"),
  logical = list(text_align = "right"),
  border = "#999999"
)
```

**Arguments**

x	vector, matrix, array, table or html_matrix: input
...	further parameters
byrow	logical: create a row or column matrix if x is one-dimensional (default: FALSE)
numeric	list: list of HTML style properties for a cell if class(x[i,j])=="numeric" (default: list(text_align="right"))
integer	list: list of HTML style properties for a cell if class(x[i,j])=="integer" (default: list(text_align="right"))
char	list: list of HTML style properties for a cell if class(x[i,j])=="character" (default: list(text_align="left"))
logical	list: list of HTML style properties for a cell if class(x[i,j])=="logical" (default: list(text_align="right"))
border	character: vector of background color for a border cell (default: "#999999")

**Value**

html\_matrix returns a html\_matrix, print returns invisible a character matrix

**Examples**

```
m <- matrix(1:6, ncol=2)
m
l <- html_matrix(m)
l
```



---

in_range	<i>in_range</i>
----------	-----------------

---

**Description**

Checks if x is between lower and upper,

**Usage**

```
in_range(x, lower, upper, rightmost.closed = TRUE, left.open = FALSE)
```

**Arguments**

x	numeric: values to check
lower	numeric: lower bound
upper	numeric: upper bound
rightmost.closed	logical: if true then $x \leq \text{upper}$ is checked otherwise $x < \text{upper}$ (default: TRUE)
left.open	logical: if true then $\text{upper} < x$ is checked otherwise $\text{lower} \leq x$ (default: FALSE)

**Value**

a logical vector whether x is in range or not

**Examples**

```
in_range(-1:2, 0, 1)
```

---

is.ASCII	<i>is.ASCII</i>
----------	-----------------

---

**Description**

Checks if txt contains only ASCII characters.

**Usage**

```
is.ASCII(txt)
```

**Arguments**

txt	character: text to check
-----	--------------------------

**Value**

logical

**Examples**

```
is.ASCII("Congratulations")
is.ASCII("Herzlichen Glückwunsch")
```

---

 men\_asso

*Association*


---

**Description**

Shiny app for the association coefficients between two categorical variables. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_asso(...)
```

**Arguments**

```
...           one or more data sets
```

**Value**

```
nothing
```

**Examples**

```
if (interactive()) men_asso()
if (interactive()) men_asso(HairEyeColor, Titanic)
```

---

 men\_bin

*men\_bin*


---

**Description**

Visualization of the probability mass and the cumulative distribution function of a binomial distribution.

**Usage**

```
men_bin(size = 10, prob = 0.5)
```

**Arguments**

```
size           integer: number of trials (zero or more)
prob           numeric: probability of success on each trial
```

**Value**

nothing

**Examples**

```
if (interactive()) men_bin()
if (interactive()) men_bin(20, 0.25)
```

---

men\_ci1

*men\_ci1*

---

**Description**

Shiny app for a confidence interval for the mean. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_ci1(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_ci1()
if (interactive()) men_ci1(stackloss)
```

---

men\_ci2

*men\_ci2*

---

**Description**

Shiny app for a confidence interval for the difference of two means. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_ci2(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_ci2()
if (interactive()) men_ci2(CO2)
```

---

men_cilen	<i>men_cilen</i>
-----------	------------------

---

**Description**

Shiny app for a length of a confidence interval for the mean.

**Usage**

```
men_cilen()
```

**Value**

nothing

**Examples**

```
if (interactive()) men_cilen()
```

---

men_cipi	<i>men_cipi</i>
----------	-----------------

---

**Description**

Shiny app for a confidence interval for the proportion. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_cipi(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_cipi()
if (interactive()) men_cipi(Titanic)
```

---

men\_cisig

*men\_cisig*

---

**Description**

Shiny app for a confidence interval for the variance. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_cisig(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_cisig()
if (interactive()) men_cisig(stackloss)
```

---

men\_corr

*Correlation*

---

**Description**

Shiny app for the correlation coefficients between two numeric variables. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_corr(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_corr()
if (interactive()) men_corr(iris)
```

---

men_die	<i>men_die</i>
---------	----------------

---

**Description**

Shiny app for detecting if a die is fair or unfair.

**Usage**

```
men_die()
```

**Value**

nothing

**Examples**

```
if (interactive()) men_die()
```

---

men_dot	<i>men_dot</i>
---------	----------------

---

**Description**

Shiny app for visualizing a univariate numeric variable as dotplot including univariate parameters. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_dot(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_dot()
if (interactive()) men_dot(iris)
```

---

men\_exp

*men\_exp*

---

**Description**

Visualization of the density and the cumulative distribution function of a exponential distribution.

**Usage**

```
men_exp(rate = 1)
```

**Arguments**

rate                    numeric: rate

**Value**

nothing

**Examples**

```
if (interactive()) men_exp()
if (interactive()) men_exp(3)
```

---

men\_hall

*men\_hall*

---

**Description**

Shiny app for the **Monty Hall problem**:

**Usage**

```
men_hall(pointdoor = 1, afteropen = 1)
```

**Arguments**

pointdoor            integer: to which door to point (default: 1)  
afteropen            integer: play strategy 1=keep door, 2=change door (default: 1)

**Details**

Suppose you're on a game show, and you're given the choice of three doors: Behind one door is a car; behind the others, goats. You pick a door, say No. 1, and the host, who knows what's behind the doors, opens another door, say No. 3, which has a goat. He then says to you, "Do you want to pick door No. 2?" Is it to your advantage to switch your choice?

**Value**

nothing

**Examples**

```
if (interactive()) men_hall()
if (interactive()) men_hall(4, 2)
```

---

men\_hist

*men\_hist*

---

**Description**

Shiny app for visualizing a univariate numeric variable as histogram. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory. #'

**Usage**

```
men_hist(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_hist()
if (interactive()) men_hist(iris)
```



---

men_hyp	<i>men_hyp</i>
---------	----------------

---

**Description**

Visualization of the probability mass and the cumulative distribution function of a hypergeometric distribution.

**Usage**

```
men_hyp(N = 60, M = 30, n = 20)
```

**Arguments**

N	integer: the number of black and white balls in the urn
M	integer: the number of white balls in the urn
n	integer: the number of balls drawn from the urn

**Value**

nothing

**Examples**

```
if (interactive()) men_hyp()
if (interactive()) men_hyp(50, 25, 10)
```

---

men_norm	<i>men_norm</i>
----------	-----------------

---

**Description**

Visualization of the density and the cumulative distribution function of a normal distribution.

**Usage**

```
men_norm(mean = 0, sd2 = 1)
```

**Arguments**

mean	numeric: mean
sd2	numeric: variance

**Value**

nothing

**Examples**

```
if (interactive()) men_norm()
if (interactive()) men_norm(1, 0.5)
```

---

men_parn	<i>men_parn</i>
----------	-----------------

---

**Description**

Shiny app for the distribution of sample parameters. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_parn(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_parn()
if (interactive()) men_parn(iris)
```

---

men_poi	<i>men_poi Visualization of the probability mass and the cumulative distribution function of a Poisson distribution.</i>
---------	--

---

**Description**

men\_poi Visualization of the probability mass and the cumulative distribution function of a Poisson distribution.

**Usage**

```
men_poi(lambda = 5)
```

**Arguments**

lambda numeric: (non-negative) mean

**Value**

nothing

**Examples**

```
if (interactive()) men_poi()
if (interactive()) men_poi(3)
```

---

men\_rank

*men\_rank*

---

**Description**

Shiny app for the rank correlation coefficients between two ordered variables. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_rank(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_rank()
if (interactive()) {
  data("plantTraits", package="cluster")
  men_rank(plantTraits)
}
```

---

men_regr	<i>men_regr</i>
----------	-----------------

---

**Description**

Shiny app for a simple linear regression. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_regr(...)
```

**Arguments**

```
...           one or more data sets
```

**Value**

nothing

**Examples**

```
if (interactive()) men_regr()  
if (interactive()) men_regr(stackloss)
```

---

men_tab	<i>Frequency tables</i>
---------	-------------------------

---

**Description**

Shiny app for frequency tables for two categorical variables. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_tab(...)
```

**Arguments**

```
...           one or more data sets
```

**Value**

nothing

**Examples**

```
if (interactive()) men_tab()
if (interactive()) men_tab(HairEyeColor, Titanic)
```

---

men_terr	<i>men_terr</i>
----------	-----------------

---

**Description**

Shiny app for a test for the true mean. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_terr(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_terr()
if (interactive()) men_terr(iris)
```

---

men_time	<i>men_time</i>
----------	-----------------

---

**Description**

Shiny app for classical time series analysis. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_time(...)
```

**Arguments**

... one or more time series

**Value**

nothing

**Examples**

```
if (interactive()) men_time()
if (interactive()) men_time(co2)
```

---

men_tmu1	<i>men_tmu1</i>
----------	-----------------

---

**Description**

Shiny app for a test for the true mean. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_tmu1(...)
```

**Arguments**

...                    one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_tmu1()
if (interactive()) men_tmu1(iris)
```

---

men_tmu2	<i>men_tmu2</i>
----------	-----------------

---

**Description**

Shiny app for a test on difference of two true means. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_tmu2(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_tmu2()
if (interactive()) men_tmu2(CO2)
```

---

*men\_tprop*

*men\_tprop*

---

**Description**

Shiny app for test on the proportion. The data used is considered as a population from which random samples can be drawn. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_tprop(...)
```

**Arguments**

... one or more data sets

**Value**

nothing

**Examples**

```
if (interactive()) men_tprop()
if (interactive()) men_tprop(Titanic)
```

---

men_vis	<i>men_vis</i>
---------	----------------

---

**Description**

Shiny app for visualizing the univariate numeric variable, e.g. boxplot, stripchart, histogram, and cumulative distribution function. If no data are given then the default data from the book will be used. Otherwise the data will be stored as RDS file in a temporary directory.

**Usage**

```
men_vis(...)
```

**Arguments**

...	one or more data sets
-----	-----------------------

**Value**

nothing

**Examples**

```
if (interactive()) men_vis()
if (interactive()) men_vis(iris)
```

---

mmstat.attrVar	<i>mmstat.attrVar</i>
----------------	-----------------------

---

**Description**

Returns the parameters for a variable. If type="numeric" then descriptive measures will be returned. Otherwise absolute and relative frequencies will be returned. For using a subset of observation set index.

**Usage**

```
mmstat.attrVar(var, type, index = NULL)
```

**Arguments**

var	vector: values of a mmstat variable
type	character: type of values, allowed are numvars, binvars, ordvars or facvars
index	integer: observation numbers to use for computation, default is to use all observations



**Value**

descriptive measures

**Examples**

```
# make sure that no other data sets are loaded
mmstat.set(datasets=NULL)
mmstat.getDataNames(mmstat.rds("CARS"))
# summary of first numeric variable in first data set in mmstat
var <- mmstat.getVar(1, 1, 'numeric')
mmstat.attrVar(var, "numeric")
# summary of first factor variable in first data set in mmstat
var <- mmstat.getVar(1, 1, 'factor')
mmstat.attrVar(var, 'factor')
```

---

mmstat.axis

*mmstat.axis*


---

**Description**

Based on range the position of the labels are determined and the axis is plotted.

**Usage**

```
mmstat.axis(side, range, at, labels, ...)
```

**Arguments**

side	an integer specifying which side of the plot the axis is to be drawn on. The axis is placed as follows: 1=below, 2=left, 3=above and 4=right.
range	range: a data range
at	the points at which tick-marks are to be drawn. Non-finite (infinite, NaN or NA) values are omitted. By default (when NULL) tickmark locations are computed, see ‘Details’ below.
labels	this can either be a logical value specifying whether (numerical) annotations are to be made at the tickmarks, or a character or expression vector of labels to be placed at the tickpoints. (Other objects are coerced by <a href="#">as.graphicsAnnot.</a> ) If this is not logical, at should also be supplied and of the same length. If labels is of length zero after coercion, it has the same effect as supplying TRUE.
...	further parameters to <a href="#">graphics::axis</a>

**Value**

adds a axis to a plot

**Examples**

```
oldpar <- par(mfrow=c(1,2))
plot(iris[,1])
plot(iris[,1], axes=FALSE)
mmstat.axis(2, iris[,1])
par(oldpar)
```

---

<code>mmstat.baraxis</code>	<i>mmstat.baraxis</i>
-----------------------------	-----------------------

---

**Description**

Based on range the position of the labels are determined and the axis is plotted.

**Usage**

```
mmstat.baraxis(side, range, at, labels, ...)
```

**Arguments**

side	an integer specifying which side of the plot the axis is to be drawn on. The axis is placed as follows: 1=below, 2=left, 3=above and 4=right.
range	range: a data range
at	the points at which tick-marks are to be drawn. Non-finite (infinite, NaN or NA) values are omitted. By default (when NULL) tickmark locations are computed, see ‘Details’ below.
labels	this can either be a logical value specifying whether (numerical) annotations are to be made at the tickmarks, or a character or expression vector of labels to be placed at the tickpoints. (Other objects are coerced by <a href="#">as.graphicsAnnot.</a> ) If this is not logical, at should also be supplied and of the same length. If labels is of length zero after coercion, it has the same effect as supplying TRUE.
...	further parameters to <a href="#">graphics::axis</a>

**Value**

adds a axis to a plot

**Examples**

```
oldpar <- par(mfrow=c(1,2))
x <- 0:15
px <- dbinom(x, 10, 0.5)
plot(x, px, type="h")
plot(x, px, type="h", axes=FALSE)
mmstat.baraxis(1, range(x), at=x, labels=as.character(x))
par(oldpar)
```

---

mmstat.dec	<i>mmstat.dec</i>
------------	-------------------

---

**Description**

Computes the number of the significant digits based on the smallest non-zero difference of the sorted data.

**Usage**

```
mmstat.dec(x, ord = NULL)
```

**Arguments**

x	numeric: data vector
ord	index: subset of the ordered data (default: NULL)

**Value**

The number of significant digits and (the subset of) the order of the data.

**Examples**

```
x <- rnorm(20)
d <- mmstat.dec(x)
# create strings so that they are unique (if they were)
sprintf("%.*f", d$dec, x)
```

---

mmstat.getDataNames	<i>mmstat.getDataNames</i>
---------------------	----------------------------

---

**Description**

Returns the names of data sets and stores them in the internal environment. The name of the data set is base name without extension.

**Usage**

```
mmstat.getDataNames(...)
```

**Arguments**

...	character: names of the data sets.
-----	------------------------------------

**Value**

the names of the data sets

**Examples**

```
files <- mmstat.rds("HAIR.EYE.COLOR", "TITANIC")
mmstat.getDataNames(files)
```

---

mmstat.getDataSets      *mmstat.getDataSets*

---

**Description**

Reads data set(s) into the mmstat object.

**Usage**

```
mmstat.getDataSets(...)
```

**Arguments**

...                      character: file name(s) of RDS data file(s)

**Value**

the names of the data set(s)

**Examples**

```
# not used, deprecated??
```

---

mmstat.getLog              *mmstat.getLog*

---

**Description**

Returns the internal log message as HTML. In a Shiny app the log message are updated every 100 milliseconds

**Usage**

```
mmstat.getLog(session)
```

**Arguments**

session                      session object

**Value**

HTML code

## Examples

```
# will work only in A Shiny app
if (interactive()) {
  require("shiny")
  ui <- fluidPage(
    titlePanel("getLog example"),
    sidebarLayout(sidebarPanel(
      actionButton("quit", "Quit")),
      mainPanel(textOutput("log")))
  )
}
#
server <- function(input, output, session) {
  observeEvent(input$quit, { stopApp() })
  output$log <- renderText({ mmstat.getLog(session) })
}
#
shinyApp(ui, server)
}
```

---

mmstat.getValues

*mmstat.getValues*

---

## Description

`mmstat.getValues` returns a list with named elements. If the parameter is `NULL` then a default value stored local will be used. `mmstat.getValue` returns a value. If the parameter is `NULL` or `NA` then `def` will be returned

## Usage

```
mmstat.getValues(local, ...)
```

```
mmstat.getValue(val, def)
```

## Arguments

`local` list: default values for the named parameter

`...` list of named parameters

`val` value for a parameter

`def` default value for a parameter

## Value

a list of requested parameters

**Examples**

```
def <- list(a=3)
mmstat.getValues(def, b=3, a=NULL)
mmstat.getValue(NA, 5)
mmstat.getValue(NULL, 5)
mmstat.getValue(3, 5)
```

---

mmstat.getVar	<i>mmstat.getVar</i>
---------------	----------------------

---

**Description**

Returns the from a data set a variable from the given type.

**Usage**

```
mmstat.getVar(
  dataname = NULL,
  varname = NULL,
  vartype = NULL,
  na.action = stats::na.omit
)
```

**Arguments**

dataname	integer: number of data set
varname	integer: number of variable
vartype	character: variable type, one of numeric, binary, ordered, or factor
na.action	function: indicate what should happen when the data contain NAs (default: <a href="#">stats::na.omit</a> )

**Value**

a variable of the given type

**Examples**

```
# make sure that no other data sets are loaded
mmstat.set(datasets=NULL)
mmstat.getDataNames(mmstat.rds("CARS"))
# summary of first numeric variable in first data set in mmstat
str(mmstat.getVar(1, 1, 'numeric'))
# summary of first factor variable in first data set in mmstat
str(mmstat.getVar(1, 1, 'factor'))
```

---

```
mmstat.getVariableNames  
mmstat.getVariableNames
```

---

**Description**

Returns all variable names of data set stored in the internal environment.

**Usage**

```
mmstat.getVariableNames(name)
```

**Arguments**

name                    character or numeric: name or index of data set

**Value**

vector of names

**Examples**

```
# Delete all stored data sets  
mmstat.set(datasets=NULL)  
# Load CAR data set into mmstat  
mmstat.getDataNames(mmstat.rds("CARS"))  
# Extract names of all variables  
mmstat.getVariableNames(1)
```

---

```
mmstat.getVarNames     mmstat.getVarNames
```

---

**Description**

Returns the variable names of a specific type from a mmstat data set.

**Usage**

```
mmstat.getVarNames(dataname, vartype, which = NULL)
```

**Arguments**

dataname                character: name of data set  
vartype                 character: type of variable, either numeric, ordered, factor, or binary  
which                    integer: index number

**Value**

a vector or element of variable names which have the type var type

**Examples**

```
# Load CAR data set into mmstat
mmstat.getDataNames(mmstat.rds("CARS"))
# Extract names of numeric variables
mmstat.getVarNames(1, "numeric")
```

---

mmstat.lang	<i>mmstat.lang</i>
-------------	--------------------

---

**Description**

Loads a **PO file** for a translation into the internal environment.

**Usage**

```
mmstat.lang(pof = NULL)
```

**Arguments**

pof                    character: file name

**Value**

nothing

**Examples**

```
mmstat.lang()
```

---

mmstat.log	<i>mmstat.log</i>
------------	-------------------

---

**Description**

Writes a message into the internal log.

**Usage**

```
mmstat.log(txt)
```

**Arguments**

txt                    character: message to write



**Value**

nothing

**Examples**

```
mmstat.log("Test")
```

---

mmstat.math

*mmstat.math*


---

**Description**

Returns a math expression based on HTML special characters notation.

**Usage**

```
mmstat.math(txt)
```

**Arguments**

txt                    character: input text

**Value**

expression

**Examples**

```
mmstat.math(" &bar(X);~&N(mu[0], sigma^2/n); ")
mmstat.math("&H[0];: &mu==mu[0]; vs. &H[1];: &mu!=mu[0]; ")
```

---

mmstat.merge

*mmstat.merge*


---

**Description**

Computes a new range from by a union of the two ranges.

**Usage**

```
mmstat.merge(range1, range2)
```

**Arguments**

range1                range: first range  
range2                range: second range

**Value**

new range

**Examples**

```
mmstat.merge(c(0,1), c(0.5, 2)) # returns c(0, 2)
```

---

```
mmstat.plotTestRegions  
      mmstat.plotTestRegions
```

---

**Description**

Plots the test regions in a plot

**Usage**

```
mmstat.plotTestRegions(  
  crit,  
  xlim,  
  ylim,  
  cex,  
  close = FALSE,  
  col = "black",  
  label = NULL,  
  pos = 1  
)
```

**Arguments**

crit	numeric(2): critical value(s)
xlim	numeric(2): the x limits of the plot
ylim	numeric(2): the y limits of the plot
cex	numeric: amount by which plotting text should be magnified relative to the default
close	logical: should the region box be closed by vertical lines (default: FALSE)
col	color: specification for the default plotting color (default: "black")
label	unused
pos	unused

**Value**

adds test regions to a plot

**Examples**

```
x <- (-30:30)/10
px <- dnorm(x)
plot(x, px, type="l", ylim=c(-0.25, max(px)), xlim=range(x))
mmstat.plotTestRegions(crit=c(-1.96, +1.96), xlim=range(x), ylim=c(-0.2, -0.1), cex=1)
```

---

`mmstat.pos`*mmstat.pos*

---

**Description**

Returns a linear interpolation based on minmax.

**Usage**

```
mmstat.pos(minmax, pos)
```

**Arguments**

<code>minmax</code>	numeric(2): range to interpolate between
<code>pos</code>	numeric: proportion(s) to interpolate, usually between zero and one

**Value**

interpolated values

**Examples**

```
mmstat.pos(c(0,360), 0.5)
```

---

`mmstat.range`*mmstat.range*

---

**Description**

Computes a range from several R objects by union.

**Usage**

```
mmstat.range(...)
```

**Arguments**

<code>...</code>	R objects
------------------	-----------

**Value**

range

**Examples**

```
mmstat.range(-5:5, 0:10) # returns c(-5, 10)
```

---

 mmstat.rds

*mmstat.rds*


---

**Description**

Returns the full file names of all or specific data set that come with the package.

**Usage**

```
mmstat.rds(...)
```

**Arguments**

```
...          names of data sets
```

**Value**

full file names

**Examples**

```
mmstat.rds() # return all RDS file that come with the package
mmstat.rds("HAIR.EYE.COLOR", "TITANIC") # location of specific data sets
```

---

 mmstat.round.down

*mmstat.round.down*


---

**Description**

Rounds down.

**Usage**

```
mmstat.round.down(x, digits = 0)
```

**Arguments**

```
x          numeric: values for rounding
digits     numeric: digits for rounding (default: 0)
```

**Value**

down rounded values

**Examples**

```
x <- runif(5)
cbind(x, mmstat.round.down(x, 1))
```

---

mmstat.round.up	<i>mmstat.round.up</i>
-----------------	------------------------

---

**Description**

Rounds up.

**Usage**

```
mmstat.round.up(x, digits = 0)
```

**Arguments**

x	numeric: values for rounding
digits	numeric: digits for rounding (default: 0)

**Value**

uprounded values

**Examples**

```
x <- runif(5)
cbind(x, mmstat.round.up(x, 1))
```

---

mmstat.set	<i>mmstat.set</i>
------------	-------------------

---

**Description**

mmstat.set sets one (or more) parameter to the internal environment. mmstat.get return one or more parameters from the internal environment.

**Usage**

```
mmstat.set(...)
```

```
mmstat.get(...)
```

**Arguments**

...                    named parameters with values or names

**Value**

nothing

**Examples**

```
mmstat.set(debug=0)
mmstat.get("debug")
mmstat.get("debug", "shiny") # returns a list
```

---

mmstat.sliderInput     *mmstat.sliderInput*

---

**Description**

A modified sliderInput for mmstat which supports user defined tick marks.

**Usage**

```
mmstat.sliderInput(...)
```

**Arguments**

...                    parameters for [shiny::sliderInput](#)

**Value**

the HTML output

**Examples**

```
ticks <- c(80, 85, 90, 95, 98, 99, 99.5, 99.9)
mmstat.sliderInput("id", "label", min=1, max=length(ticks), value=3, step=1, ticks=ticks)
```

---

mmstat.ticks	<i>mmstat.ticks</i>
--------------	---------------------

---

**Description**

Returns tick marks for a log based scale between nmin and nin.

**Usage**

```
mmstat.ticks(nin, nmin = 3, tin = 11)
```

**Arguments**

nin	integer: maximum of scale
nmin	integer: minimum of scale
tin	integer: number of desired tick marks

**Value**

vector of tick marks

**Examples**

```
mmstat.ticks(506)
```

---

mmstat.ui.call	<i>mmstat.ui.call</i>
----------------	-----------------------

---

**Description**

Calls the underlying Shiny UI element (selectInput, ...).

**Usage**

```
mmstat.ui.call(inputId, ...)
```

**Arguments**

inputId	character: the input slot called
...	further parameters given to the call

**Value**

whatever the call to the underlying Shiny UI element returns

## Examples

```
mmstat.ui.elem(inputId="alpha", type="significance")  
mmstat.ui.call("alpha")
```

---

mmstat.ui.elem

*mmstat.ui.elem*

---

## Description

Adds a new UI element to the app interface. The following types from Shiny are allowed:

- [actionButton](#),
- [checkboxInput](#),
- [checkboxGroupInput](#),
- [dateInput](#),
- [dateRangeInput](#),
- [fileInput](#),
- [helpText](#),
- [numericInput](#),
- [radioButtons](#),
- [selectInput](#),
- [sliderInput](#),
- [submitButton](#), and
- [textInput](#).

Additionally some standard statistical UI elements are supported (links go to the Shiny element used):

- [sampleSize](#),
- [drawSample](#),
- [speedSlider](#),
- [confidenceLevel](#),
- [significance](#),
- [testHypotheses](#),
- [dataSet](#),
- [variable1](#),
- [variableN](#), and
- [fontSize](#).

Partially these elements have default settings which can be overwritten.



**Usage**

```
mmstat.ui.elem(inputId, type, ...)
```

**Arguments**

inputId	character: input slot that will be used to access the value
type	character: element type
...	further named parameter to Shiny UI elements

**Value**

nothing

**Examples**

```
mmstat.ui.elem(inputId="alpha", type="significance")
```

---

mmstat.ui.update	<i>mmstat.ui.update</i>
------------------	-------------------------

---

**Description**

Call for a update of an underlying Shiny UI element (selectInput, ...).

**Usage**

```
mmstat.ui.update(inputId, ...)
```

**Arguments**

inputId	character: the input slot called
...	further parameters given to the call

**Value**

whatever the update to the underlying Shiny UI element returns

**Examples**

```
mmstat.ui.elem(inputId="alpha", type="significance")  
mmstat.ui.call("alpha")
```

`mmstat.warn`*mmstat.warn*

---

**Description**

Writes a warning text into the log object in the internal mmstat object.

**Usage**

```
mmstat.warn(cond, txt)
```

**Arguments**

<code>cond</code>	logical: condition to test
<code>txt</code>	character: text to write if cond is true

**Value**

nothing

**Examples**

```
mmstat.warn(TRUE, "just a true seen")
```

---

`resetpar`*resetpar*

---

**Description**

Resets the par if necessary.

**Usage**

```
resetpar(oldpar)
```

**Arguments**

<code>oldpar</code>	graphical parameters
---------------------	----------------------

**Value**

nothing

**Examples**

```
par("mar")
oldpar <- par(no.readonly = TRUE)
par(mar = c(0,0,0,0))
par("mar")
resetpar(oldpar)
par("mar")
```

---

stopif	<i>stopif</i>
--------	---------------

---

**Description**

A equivalent to stopifnot: if cond is TRUE then a error is thrown.

**Usage**

```
stopif(cond, txt)
```

**Arguments**

cond	logical: condition to test
txt	character: error message

**Value**

nothing

**Examples**

```
if (interactive()) stopif(1+1==2, "1+1 can not be 2, this is fake science!")
```

---

table2dataframe	<i>table2dataframe</i>
-----------------	------------------------

---

**Description**

Converts a table to a full data frame.

**Usage**

```
table2dataframe(tab, ...)
```

**Arguments**

tab	table: contingency table
...	further parameters given to <a href="#">base::as.data.frame.table</a>

**Value**

a data frame with `sum(tab)` rows and `length(dim(tab))` cols

**Examples**

```
table2dataframe(Titanic)
```

---

```
toHTML.html_matrix    toHTML
```

---

**Description**

Returns a HTML representation of a matrix and optionally shows the result in the browser. If you decide to view the result in a browser then the HTML will be written to a temporary file and [utils::browseURL\(\)](#) called

**Usage**

```
## S3 method for class 'html_matrix'  
toHTML(x, browser = FALSE, ...)  
  
## S3 method for class 'table'  
toHTML(x, browser = FALSE, ...)  
  
## S3 method for class 'matrix'  
toHTML(x, browser = FALSE, ...)
```

**Arguments**

```
x                html_matrix object  
browser          logical: show HTML in a browser (default: FALSE)  
...              further parameters to utils::browseURL\(\)
```

**Value**

html\_matrix object

**Examples**

```
library("tools")  
m  <- matrix(1:12, ncol=4)  
hm <- html_matrix(m)  
html <- toHTML(hm, browser=interactive())
```

---

toRDS	<i>toRDS</i>
-------	--------------

---

**Description**

Saves one or more data sets in RDS format to a temporary directory (`tmpdir()`). Data sets must have the class `ts` or something that can be converted to a data frame, e.g. `matrix`, `table`, etc.

**Usage**

```
toRDS(...)
```

**Arguments**

```
...          data sets to save
```

**Value**

returns the name of the created files

**Examples**

```
toRDS(Titanic) # saves to tmpdir/Titanic.rds
```

---

ucfirst	<i>ucfirst</i>
---------	----------------

---

**Description**

Uppercases the first character in `txt`.

**Usage**

```
ucfirst(txt)
```

**Arguments**

```
txt          character:
```

**Value**

character

**Examples**

```
ucfirst("hello world")
```

---

zebra	<i>zebra</i>
-------	--------------

---

**Description**

zebra

**Usage**

```
zebra(x, col = c("#FFFFFF", "#CCCCCC"), byrow = TRUE)
```

**Arguments**

x	html_matrix object
col	a vector of colors to zebra with (default:c("#FFFFFF", "#CCCCCC"))
byrow	logical: zebra by row or by column (default: TRUE)

**Value**

html\_matrix object

**Examples**

```
library("magrittr")
library("tools")
m <- matrix(1:12, ncol=4)
hm <- html_matrix(m) %>% zebra()
html <- toHTML(hm, browser=interactive())
```

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