Package ‘clinUtils’

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Type Package

Title General Utility Functions for Analysis of Clinical Data

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Description Utility functions to facilitate the import, the reporting and analysis of clinical data. Example datasets in 'SDTM' and 'ADaM' format, containing a subset of patients/domains from the 'CDISC Pilot 01 study' are also available as R datasets to demonstrate the package functionalities.

Imports crosstalk, data.table, DT, haven, htmlwidgets, knitr, plyr, tools, utils, viridisLite

Suggests dplyr, tibble, ggplot2, plotly, htmltools, pander, rmarkdown, testthat, flextable

SystemRequirements pandoc

URL https://github.com/openanalytics/clinUtils

BugReports https://github.com/openanalytics/clinUtils/issues

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checkVarInData

Check if specified variables are in the data. If they are not, they are removed from specified variables and a message is printed.

Description

Check if specified variables are in the data. If they are not, they are removed from specified variables and a message is printed.

Usage

checkVarInData(var, data, label)

Arguments

var Character vector with variables.
data Data.frame with data.
label String with label used in message.

Value

var present in data, NULL if empty

clinColors

Colors of 'clinUtils'

Description

Default color palette is the color-blind viridis palette. See documentation of viridis.

Usage

clinColors(n, alpha = 1, begin = 0, end = 1, direction = 1, option = "D")

Arguments

n The number of colors (≥ 1) to be in the palette.
alpha The alpha transparency, a number in [0,1], see argument alpha in hsv.
begin The (corrected) hue in [0,1] at which the viridis colormap begins.
end The (corrected) hue in [0,1] at which the viridis colormap ends.
direction Sets the order of colors in the scale. If 1, the default, colors are ordered from darkest to lightest. If -1, the order of colors is reversed.
option A character string indicating the colormap option to use. Four options are available: "magma" (or "A"), "inferno" (or "B"), "plasma" (or "C"), "viridis" (or "D", the default option) and "cividis" (or "E").
**clinLinetypes**

*Linetypes of 'clinUtils'*

**Description**

A set of default linetypes are available as a vector.

**Usage**

clinLinetypes

**Format**

An object of class character of length 6.

**Value**

A character vector of linetypes.

---

**clinShapes**

*Shapes of 'clinUtils'*

**Description**

A set of default shapes are available as named vector.

**Usage**

clinShapes

**Format**

An object of class integer of length 24.

**Value**

A numeric vector of shapes.
**clinShapesText**

*Shapes of `clinUtils` as text*

---

**Description**

A set of default shapes are available as character vector.

**Usage**

clinShapesText

**Format**

An object of class character of length 24.

**Value**

A character vector of shapes.

---

**clinUtils-palette**

*Parameters for all palette functions for clinical visualizations.*

---

**Description**

Parameters for all palette functions for clinical visualizations.

**Arguments**

- **n**
  - Integer of length 1, number of elements in palette.

- **x**
  - Vector with elements used for palette. If factor, the levels are used, otherwise the unique elements of the vector. Missing values are automatically removed, excepted if `includeNA` is set to TRUE.

- **includeNA**
  - Logical (FALSE by default), should NA elements be retained in the palette in case x is specified?

**Value**

Specific palettes used in `clinUtils`. 
**colorPaletteNRIND**

Color palette for a standard CDISC Normal/Reference Range Indicator.

**Description**

Color palette for a standard CDISC Normal/Reference Range Indicator.

**Usage**

```r
colorPaletteNRIND
```

**Format**

A named character vector with color for typical Normal Reference Range Indicator variable:

- "LOW": orange
- "NORMAL": green
- "HIGH": orange
- "ABNORMAL": red
- "UNKNOWN" or 'NA': grey
- "NA": grey

**compareDiff**

Get differences between two data.frames

**Description**

Get differences between two data.frames

**Usage**

```r
compareDiff(
  newData,
  oldData,
  referenceVars = intersect(colnames(newData), colnames(oldData)),
  changeableVars = NULL
)
```
compareDiff

Arguments

newData data.frame object representing the new data
oldData data.frame object representing the old data
referenceVars character vector of the columns in the data that are used as reference for the comparison. If not specified, all columns present both in newData and oldData are considered.
changeableVars character vector of the columns in the data for which you want to assess the change, e.g., variables that might have changed from the old to the new data. If not specified, only 'Addition' and 'Removal' are detected.

Value

Object of class 'diff.data', i.e., a data.frame with columns:

- 'Comparison type': type of difference between the old and new data, either:
  - 'Change': records present both in new and old data, based on the reference variables, but with difference(s) in changeable vars
  - 'Addition': records with reference variables present in new but not in old data
  - 'Removal': records with reference variables present in old but not in new data
- 'Version': 'Previous' or 'Current' depending if record represents content from old or new data respectively
- referenceVars
- changeableVars

Identification of the differences between datasets

To identify the differences between datasets, the following steps are followed:

1. removal of records identical between the old and new dataset (will be considered as 'Identical' later on)
2. records with a reference value present in the old dataset but not in the new dataset are considered 'Removal'
3. records with a reference value present in the new dataset but not in the old dataset are considered 'Addition'
4. records with reference value present both in the new and old dataset, after filtering of identical records and with difference in the changeable variables are considered 'Change'

Author(s)

Laure Cougnaud
compareTables

Compare tables

Description

Compare tables

Usage

```r
compareTables(
  newData,
  oldData,
  referenceVars = intersect(colnames(newData), colnames(oldData)),
  changeableVars = NULL,
  outputType = c("table-comparison", "newData-diff", "oldData-diff",
                 "table-comparison-interactive", "newData-diff-interactive",
                 "oldData-diff-interactive"),
  ...)
```

Arguments

- **newData**: data.frame object representing the new data
- **oldData**: data.frame object representing the old data
- **referenceVars**: character vector of the columns in the data that are used as reference for the comparison. If not specified, all columns present both in `newData` and `oldData` are considered.
- **changeableVars**: character vector of the columns in the data for which you want to assess the change, e.g., variables that might have changed from the old to the new data. If not specified, only 'Addition' and 'Removal' are detected.
- **outputType**: String describing which output should be returned, (multiple are possible), either:
  - 'table-comparison': data.frame containing difference between two datasets, see 'output' of `compareDiff` function.
  - 'table-comparison-interactive': `datatable` object with differences between the two datasets, see 'output' of `exportDiffData`.
  - 'newData-diff' or 'oldData-diff': data.frame with new/old data respectively, containing the information if each record differs in the old/new datasets respectively. See output of `mergeDiffWithData`.
  - 'newData-diff-interactive' or 'oldData-diff-interactive': `datatable` with new/old data respectively, containing the information if each record differs in the old/new datasets respectively. See output of `exportDiffData`.
- **...**: Any parameters passed to the `exportDiffData` function. These are only used if 'table-comparison-interactive' is specified in `outputType`. 
Value

One of the output types specified in outputType. By default, all outputs are returned. If multiple output types are specified, a list of those are returned (named by output type).

Identification of the differences between datasets

To identify the differences between datasets, the following steps are followed:

1. removal of records identical between the old and new dataset (will be considered as 'Identical' later on)
2. records with a reference value present in the old dataset but not in the new dataset are considered 'Removal'
3. records with a reference value present in the new dataset but not in the old dataset are considered 'Addition'
4. records with reference value present both in the new and old dataset, after filtering of identical records and with difference in the changeable variables are considered 'Change'

Author(s)

Laure Cougnaud, Michela Pasetto

Examples

```r
## Example 1
# In this case the referenceVar 'a' is the same
# the comparison highlights only as change in the variables 'c' and 'd'
newData <- data.frame("a" = c(1, 2, 3, 4),
                      "b" = c(5, 6, 7, 8),
                      "c" = rep(1, 4),
                      "d" = rep(2, 4))
oldData <- data.frame("a" = c(1, 2, 3, 4),
                      "b" = c(3, 4, 7, 8),
                      "c" = rep(2, 4),
                      "d" = rep(1, 4))
compareTables(
  newData = newData,
  oldData = oldData,
  referenceVars = "a",
  changeableVars = c("c", "d")
)

## Example 2
# In this case the referenceVar 'a' changes in the last two rows
# the comparison highlights as change the second and third rows in the variables 'c' and 'd'
# whereas the last rows are additions/removals with respect to the reference 'a'
newData <- data.frame(
```
"a" = c(7, 1, 2, 3, 4),
"b" = c(2, 1, 6, 7, 8),
"c" = rep(1, 5),
"d" = rep(2, 5)
}
oldData <- data.frame(
  "a" = c(7, 1, 2, 5, 6),
  "b" = c(2, 3, 4, 7, 8),
  "c" = c(1, rep(2, 4)),
  "d" = c(2, rep(1, 4))
)
compareTables(
  newData = newData,
  oldData = oldData,
  referenceVars = "a",
  changeableVars = c("c", "d")
)

## Example 3
# In this case the referenceVar 'a' is the same
# also the variable 'c' is the same and it's the only changeable var evaluated
newData <- data.frame(
  "a" = c(1, 2, 3, 4),
  "b" = c(5, 6, 7, 8),
  "c" = rep(1, 4),
  "d" = rep(2, 4)
)
oldData <- data.frame(
  "a" = c(1, 2, 3, 4),
  "b" = c(3, 4, 7, 8),
  "c" = rep(1, 4),
  "d" = rep(1, 4)
)
compareTables(
  newData = newData,
  oldData = oldData,
  referenceVars = "a",
  changeableVars = "c"
)

## In case only a specific output should be returned:

newData <- data.frame(
  "a" = c(7, 1, 2, 3, 4),
  "b" = c(2, 1, 6, 7, 8),
  "c" = rep(1, 5),
  "d" = rep(2, 5)
)
oldData <- data.frame(
  "a" = c(7, 1, 2, 5, 6),
  "b" = c(2, 3, 4, 7, 8),
  "c" = c(1, rep(2, 4)),
  "d" = c(2, rep(1, 4))
)
# get only the differences between datasets:

# as a data.frame
compareTables(newData = newData, oldData = oldData,
              referenceVars = "a", changeableVars = c("c", "d"),
              outputType = "table-comparison")

# as an interactive DataTable
compareTables(newData = newData, oldData = oldData,
              referenceVars = "a", changeableVars = c("c", "d"),
              outputType = "table-comparison-interactive"
)

# only the new data
compareTables(
    newData = newData, oldData = oldData,
    referenceVars = "a", changeableVars = c("c", "d"),
    outputType = "newData-diff"
)

# only the new data in interactive mode
compareTables(
    newData = newData, oldData = oldData,
    referenceVars = "a", changeableVars = c("c", "d"),
    outputType = "newData-diff-interactive"
)

# only the new data in static and interactive mode
compareTables(
    newData = newData, oldData = oldData,
    referenceVars = "a", changeableVars = c("c", "d"),
    outputType = c("newData-diff", "newData-diff-interactive")
)

# only the old data
compareTables(newData = newData, oldData = oldData,
              referenceVars = "a", changeableVars = c("c", "d"),
              outputType = "oldData-diff"
)

# only the old data in interactive mode
compareTables(
    newData = newData, oldData = oldData,
    referenceVars = "a", changeableVars = c("c", "d"),
    outputType = "oldData-diff-interactive"
)

# only the old data in static and interactive mode
compareTables(
    newData = newData, oldData = oldData,
    referenceVars = "a", changeableVars = c("c", "d"),
    outputType = c("oldData-diff", "oldData-diff-interactive")
)

## no changeable vars

newData <- data.frame(
    "a" = c(7, 1, 2, 3, 4),
    ...)
```r
b" = c(2, 1, 6, 7, 8),
"c" = rep(1, 5),
"d" = rep(2, 5)
}
oldData <- data.frame(
  "a" = c(7, 1, 2, 5, 6),
  "b" = c(2, 3, 4, 7, 8),
  "c" = c(1, rep(2, 4)),
  "d" = c(2, rep(1, 4))
)
compareTables(newData = newData, oldData = oldData,
  referenceVars = "a"
)

## duplicated records

## with labels in the interactive format, see ? getClinDT
```
General parameters used for the comparison table functionality

## Description

General parameters used for the comparison table functionality

## Arguments

- **newData**: data.frame object representing the new data
- **oldData**: data.frame object representing the old data
- **referenceVars**: character vector of the columns in the data that are used as reference for the comparison. If not specified, all columns present both in `newData` and `oldData` are considered.
- **changeableVars**: character vector of the columns in the data for which you want to assess the change, e.g., variables that might have changed from the old to the new data. If not specified, only 'Addition' and 'Removal' are detected.
- **diffData**: Object of class `diff.data` containing differences between datasets, as returned by the `compareDiff` function.
- **outputType**: String describing which output should be returned, (multiple are possible), either:
  - 'table-comparison': data.frame containing difference between two datasets, see 'output' of `compareDiff` function.
  - 'table-comparison-interactive': `datatable` object with differences between the two datasets, see 'output' of `exportDiffData`.
  - 'newData-diff' or 'oldData-diff': data.frame with new/old data respectively, containing the information if each record differs in the old/new datasets respectively. See output of `mergeDiffWithData`.
  - 'newData-diff-interactive' or 'oldData-diff-interactive': `datatable` with new/old data respectively, containing the information if each record differs in the old/new datasets respectively. See output of `exportDiffData`.  

```r
oldData = oldData,
referenceVars = "a",
changeableVars = c("c", "d"),
# parameters passed to datatable
colnames = c(
  "My reference variable" = "a",
  "Changeable variable c" = "c",
  "Changeable variable d" = "d"
)
)
**convertToDatatable**  
*Convert to data.table*

**Description**  
Convert a data frame into a data.table object.

**Usage**  
```r  
convertToDatatable(data)  
```

**Arguments**

- `data`  
  A data.frame

**Value**  
A data.table object.

---

**convertToDateTime**  
*Convert character vector to date/time object*

**Description**  
Convert character vector to date/time object.

**Usage**  
```r  
convertToDateTime(  
  x,  
  format = c("%Y-%m-%dT%H:%M", "%Y-%m-%d"),  
  colName = NULL,  
  verbose = TRUE  
)  
```
Arguments

- **x**: character vector to convert to date/time
- **format**: string with possible format(s) of the input date/time in the ADaM dataset. If multiple are specified, each format is tested successively, until at least one element in the input vector is converted with the specified format (non missing, following the approach described in the `format` parameter of the `strptime` function). See the 'Details' section of the help of the function, for more information about this format.
- **colName**: string with name of column, used in message (if any).
- **verbose**: logical, if TRUE (by default) progress messages are printed during execution.

Value

Vector of class `POSIXct`

Author(s)

Laure Cougnaud

---

dataADaMCDISCP01

Example of ADaM datasets from the CDISC original Pilot 01 study

Description

This contains a subset of the CDISC Pilot 01 study dataset for:

- a selected subset of subjects
- a selected subset of domains:
  - subject-level (`adsl`)
  - adverse event (`adae`)
  - laboratory chemistry data (`adlbc`)
  - vital signs (`advs`)
  - concomitant medications (`adcm`)
- efficacy:
  - ADAS-COG Data (`adqadas`), containing one of the primary endpoint: ADAS-Cog (11)
  - (Alzheimer’s Disease Assessment Scale - Cognitive Subscale)
  - CIBIC+ questionnaire data (`adqscibc`), containing one of the primary endpoint: CIBIC+
    - (Video-referenced Clinician’s Interview-based Impression of Change)
  - pharmacokinetic parameters (`adpp`)

Please note that this dataset contains different sets of subjects than the other example datasets.

This dataset was created following the ADaM Version 2.0 standard. This dataset contains the 'Modified \& augmented version of cdiscpilot01' dataset.
**Format**

List of data.frames containing the ADaM dataset for each selected domain.
Labels for the different variables across datasets is available via the `labelVars` attribute.

**Author(s)**

Laure Cougnaud

**Source**

Original (and entire) datasets are available in: 

**See Also**

`loadDataADaMSDTM`

---

**dataSDTMCDISCP01**  
**Example of SDTM datasets from the CDISC original Pilot 01 study**

**Description**

This contains a subset of the CDISC original Pilot 01 study dataset for:

- a selected subset of subjects
- a selected subset of domains:
  - demographics (`dm`)
  - treatment exposure (`ex`)
  - concomitant medications (`cm`)
  - medical history (`mh`)
  - adverse event (`ae`)
  - laboratory (`lb`)
  - vital signs (`vs`)
  - subject visit (`sv`)

This dataset was created following the SDTM Version 1.1 standard.

**Format**

List of data.frames containing the SDTM dataset for each selected domain.
Labels for the different variables across datasets is available via the `labelVars` attribute.

**Author(s)**

Laure Cougnaud
Source

Original (and entire) datasets are available in: https://github.com/phuse-org/phuse-scripts/tree/master/data/sdtm/cdiscpilot01 See in particular the define.xml file for further description of the datasets and variables name.

See Also

loadDataADaMSDTM

---

### exportDiffData

**Export the 'diff.data' object from compareDiff to a user-friendly format**

#### Description

Export the 'diff.data' object from compareDiff to a user-friendly format

#### Usage

```r
exportDiffData(
  diffData,  # Object of class 'diff.data' containing differences between datasets, as returned by the compareDiff function.
  newDataDiff,  # data.frame with new data with differences as returned by the mergeDiffWithData. The data set contains the new data with the information if each record differs in the new dataset.
  oldDataDiff,  # data.frame with old data with differences as returned by the mergeDiffWithData. The data set contains the old data with the information if each record differs in the old dataset.
  referenceVars = attr(diffData, "referenceVars"),  # character vector of the columns in the data that are the used as reference for the comparison. If not specified, all columns present both in newData and oldData are considered.
  changeableVars = attr(diffData, "changeableVars"),  # character vector of the columns in the data for which you want to assess the change, e.g. variables that might have changed from the old to the new data. If not specified, only 'Addition' and 'Removal' are detected.
  to = "DT",  #...
  ...
)
```

#### Arguments

- **diffData**: Object of class 'diff.data' containing differences between datasets, as returned by the compareDiff function.
- **newDataDiff**: data.frame with new data with differences as returned by the mergeDiffWithData. The data set contains the new data with the information if each record differs in the new dataset.
- **oldDataDiff**: data.frame with old data with differences as returned by the mergeDiffWithData. The data set contains the old data with the information if each record differs in the old dataset.
- **referenceVars**: character vector of the columns in the data that are the used as reference for the comparison. If not specified, all columns present both in newData and oldData are considered.
- **changeableVars**: character vector of the columns in the data for which you want to assess the change, e.g. variables that might have changed from the old to the new data. If not specified, only 'Addition' and 'Removal' are detected.
Value

Depending on the `to` parameter:

- `'DT'`: a `datatable` with the difference between datasets, with:
  - highlighting depending on the difference between datasets:
    * 'Addition’ in green
    * 'Removal’ in yellow
    * 'Change’ in lightblue
    * 'Identical’ are not highlighted
  - records only present in the old dataset are displayed in italic

```r
formatDetailsComparison
dt
```

Description

Format details comparison

Usage

```r
formatDetailsComparison(
diffData,
referenceVars = attr(diffData, "referenceVars"),
changeableVars = attr(diffData, "changeableVars")
)
```

Arguments

- `diffData` Object of class `diff.data` containing differences between datasets, as returned by the `compareDiff` function.
- `referenceVars` character vector of the columns in the data that are the used as reference for the comparison.
  If not specified, all columns present both in `newData` and `oldData` are considered.
- `changeableVars` character vector of the columns in the data for which you want to assess the change, e.g. variables that might have changed from the old to the new data.
  If not specified, only 'Addition' and 'Removal' are detected.
Value

diffData with extra columns: `[]`.diff` for the `referenceVars` and `changeableVars` columns, and attributes: `colsDiff` as a named vector with mapping with input variables (names) and corresponding diff variables.

Author(s)

Laure Cougnaud

formatDTBarVar

Format a variable in a **datatable** as a barplot.

Description

Format a variable in a **datatable** as a barplot.

Usage

```r
formatDTBarVar(
  tableDT,
  data,
  barVar = NULL,
  barColorThr = NULL,
  barRange = NULL,
  getCol = function(x) x
)
```

Arguments

tableDT **datatable** object
data Data.frame with content of `tableDT`.
barVar Character vector with numeric variable of `data` which should be represented as bar in the table.
barColorThr Numeric vector with threshold to consider to color the bar, either:
  - a numeric vector of length 1, same threshold for all bars
  - named vector with threshold for each bar, named with the variable in `barVar`
barRange (optional) range for the bars, either:
  - a numeric vector of length 2, same range for all bars
  - list with range for each bar, named with the variable in `barVar`
getCol Function, which for an index of a column in `data` returns the index of the column to be passed to `formatStyle`
Value

Updated tableDT

Author(s)

Laure Cougnaud

formatLabel

Concatenate and format text strings to a label

Description

This function concatenates and formats text strings to a label e.g. to use for chunk and table/figures

Usage

formatLabel(...)

Arguments

... string(s) to be concatenated to form label or data.frame with only one row. If an unique data.frame is specified, the different columns are collapsed to form one label.

Value

String with chunk label

Author(s)

Laure Cougnaud

formatLabelChunk

Concatenate and format text strings to a chunk label

Description

Concatenate and format text strings to a chunk label

Usage

formatLabelChunk(...)

Arguments

... string to be concatenated to form chunk label
**formatLongLabel**

**Value**
String with chunk label

**Author(s)**
Laure Cougnaud

---

**formatLongLabel**  
*Format a variable with long labels*

---

**Description**
This function formats a variable with long labels by wrapping its elements into multiple lines.

**Usage**

```r
formatLongLabel(x, width = 20)
```

**Arguments**

- `x` character vector with labels to format
- `width` target maximum size. Note: a word longer than this width won’t be split (see `strwrap`).

**Value**
Vector with formatted labels

**Author(s)**
Laure Cougnaud

```r
longLabel <- "This is a very long description of the variable in the dataset"
cat(longLabel) cat(formatLongLabel(longLabel))
```

---

**formatTableLabel**  
*Concatenate and format text strings to a label of a table*

---

**Description**
This function concatenates and formats text strings to a label of a table for bookdown package

**Usage**

```r
formatTableLabel(...)```
formatVarForPlotLabel

Arguments

... string to be concatenated to form label

Value

String with chunk label

Author(s)

Laure Cougnaud

formatVarForPlotLabel  Format parameter variable to be displayed in the labels of a plot

Description

The following workflow is used:

1. format the variable as a factor
2. wrap it across multiple lines if needed
3. sort (its levels) according to a grouping variable

Usage

formatVarForPlotLabel(
  data,
  paramVar = NULL,
  paramGroupVar = NULL,
  revert = FALSE,
  width = 20
)

Arguments

data  data.frame with data
paramVar  string, variable of data with parameter
paramGroupVar  (optional) character vector with variable(s) of data with grouping. If specified, the parameters will be grouped by this(these) variable(s) in the y-axis.
revert  logical, if TRUE revert the order of the levels of the variable
width  max number of characters in the codeparamVar parameter.

Value

Vector with re-formatted paramVar, NULL if empty
Author(s)

Laure Cougnaud

library(ggplot2)
data(dataADaMCDISCP01)
dataAE <- dataADaMCDISCP01$ADAE

# by default, groups are sorted alphabetically in ggplot2 (from bottom to top for an histogram)
ggplot(data = dataAE, aes(y = AEDECOD, fill = AEBODSYS)) + geom_histogram(stat="count")

# by default: labels are set to a new line if more than 20 characters: dataAE$AEDECOD <- formatVarForPlotLabel(data = dataAE, paramVar = "AEDECOD")
levels(dataAE$AEDECOD) ggplot(data = dataAE, aes(y = AEDECOD, fill = AEBODSYS)) + geom_histogram(stat="count")

# revert order of the variable dataAE$AEDECOD <- formatVarForPlotLabel(data = dataAE, paramVar = "AEDECOD", revert = TRUE) levels(dataAE$AEDECOD) ggplot(data = dataAE, aes(y = AEDECOD, fill = AEBODSYS)) + geom_histogram(stat="count")

# group based on body system dataAE$AEDECOD <- formatVarForPlotLabel(data = dataAE, paramVar = "AEDECOD", paramGroupVar = "AEBODSYS") ggplot(data = dataAE, aes(y = AEDECOD, fill = AEBODSYS)) + geom_histogram(stat="count")

---

**getClinDT**  
*Convert a data.frame into interactive table*

**Description**

This function converts a data.frame from R into **datatable** object with sensitive defaults.

**Usage**

```r
getClinDT(
  data,
  nonVisibleVar = NULL,
  nonVisible = NULL,
  percVar = NULL,
  barVar = NULL,
  barColorThr = NULL,
  barRange = NULL,
  filter = "top",
  searchBox = FALSE,
  pageLength,
  fixedColumns = NULL,
  columnsWidth = NULL,
  options = list(),
  expandVar = NULL,
  expandIdx = NULL,
  escape = TRUE,
  rowGroup = NULL,
  rowGroupVar = NULL,
  vAlign = "top",
  callback = NULL,
  buttons = c("copy", "csv", "excel", "pdf", "print"),
)```

scrollX = TRUE, 
file = NULL, 
verbose = TRUE, 
... 
)

Arguments

data Data.frame, matrix or SharedData object with input data for the table.
nonVisibleVar Character vector with column(s) in data to hide in the output table (column is hidden).
nonVisible This parameter is deprecated, use the new interface with the nonVisibleVar parameter. Numeric vector with column(s) in data to not display in the output table (column is hidden), in Javascript unit: first column is 0, second column is 1, ...
percVar Character vector with percentage columns. These columns should contain the percentage from 0 to 1. The content of these columns will be rounded to 2 digits.
barVar Character vector with numeric variable of data which should be represented as bar in the table.
barColorThr Numeric vector with threshold to consider to color the bar, either:
  • a numeric vector of length 1, same threshold for all bars
  • named vector with threshold for each bar, named with the variable in barVar
barRange (optional) range for the bars, either:
  • a numeric vector of length 2, same range for all bars
  • list with range for each bar, named with the variable in barVar
If not specified, the range of each barVar variable in data is used.
filter String with position of the filter boxes (filter parameter of the datatable function), 'top' by default. Set to 'none' to not included any filtering boxes.
searchBox Logical, if TRUE (FALSE by default) a general search box is included.
pagelength Numeric with number of records to include in one page, by default set to 10. Set to Inf to include all records.
fixedColumns List with fixed columns, see corresponding parameter in the options parameter of the datatable function.
columnsWidth Character vector with column width, of length 1 (used for all columns) or of length: ncol(data)
options List with additional datatable options. This parameter overwrites the default options set internally in the function (an indicative message mentions it if that is the case).
expandVar Character vector with expandable variables of data. These columns won’t be included in the table, but displayed for each row when the ‘+’ icon in the first column of the table will be clicked on.
expandIdx Matrix named with: ‘row’/’column’ containing row/column indices to expand.
Column(s) to escape in the table (e.g. containing raw HTML code), either character, numeric or logical of length 1. See corresponding parameter in the `datatable` function.

This parameter is deprecated, please use `rowGroupVar` instead.

Character vector with colname(s) of data containing variables to group rows by. This creates row header containing this column. Please note that the original row order in data is respected, so you might want to order rows based on the grouping variables upfront.

String with vertical alignment for the cells, 'top' by default.

String with custom Javascript callback function.

DataTable buttons (passed to the 'buttons' element of the options parameter of `datatable`). To remove all buttons, set this parameter to NULL.

Logical, if TRUE (by default) a horizontal scrolling bar is included. Note: this differs from the `datatable` default (FALSE), because required for bookdown::gitbook output if table is too wide.

(optional) String with name of html file to which the created DT should be exported.

Logical, if TRUE (by default) informative messages are displayed, e.g. if specified options overwrite the internal default.

Additional parameters for the `datatable` function, e.g. table width.

The defaults set in this function allow to create specific interactive tables where columns/cells of interest can be collapsed/expanded via the `expandVar/expandIdx` parameters.

A `datatable` object.

Laure Cougnaud

library(dplyr)

data(dataADaMCDISCP01)
labelVars <- attr(dataADaMCDISCP01, "labelVars")

# example of simple adverse event table
dataAE <- dataADaMCDISCP01$ADAE

subjectsSafety <- subset(dataADaMCDISCP01$ADSL, SAFFL == "Y")$USUBJID

# compute counts of subjects presenting each AE
tableAE <- dataAE %>%

group_by(AESOC, AEDECOD) %>%
summarise(
  N = n_distinct(USUBJID),
  perc = round(N/length(subjectsSafety)*100, 3)
) %>%
arrange(desc(perc))

# extract new variables labels
tableAELabels <- getLabelVar(
  var = colnames(tableAE),
  labelVars = labelVars,
  label = c(N = '# subjects', perc = '% subjects')
)
# 'colnames' should be specified as c('new name' = 'old name', ...)
tableAELabelsDT <- setNames(names(tableAELabels), tableAELabels)
tableAEBase <- tableAE[, setdiff(colnames(tableAE), "USUBJID")]

## create table with bar

### default:
getClinDT(
  data = tableAEBase,
  barVar = "perc",
  colnames = tableAELabelsDT
)

### specify range for the bar
getClinDT(
  data = tableAEBase,
  filter = "none",
  barVar = "perc",
  barRange = c(0, 100),
  colnames = tableAELabelsDT
)

### change color according to threshold
getClinDT(
  data = tableAEBase,
  filter = "none",
  barVar = "perc",
  barColorThr = seq(from = 0, to = 100, by = 25),
  colnames = tableAELabelsDT
)

### group per system organ class:
tableAEGroup <- tableAEBase %>%
group_by(AESOC) %>%
mutate(NAESOC = sum(N)) %>%
arrange(desc(NAESOC), AESOC, desc(perc)) %>%
select(-NAESOC)
getClinDT(
  data = tableAEGroup,
  filter = "none",
  barVar = "perc",
  barRange = c(0, 100),
  colnames = tableAELabelsDT,
  rowGroupVar = "AESOC",
  pageLength = Inf
)

# expand the subject ID column, will
# be accessible when clicking on the '+’ button
# Format URL correctly with: ‘escape’,
# please note that indexing starts at 0!
getClinDT(
  data = tableAE,
  barVar = "perc",
  colnames = tableAELabelsDT,
  expandVar = "USUBJID",
  escape = grep("USUBJID", colnames(tableAE))-1
)

# with fixed columns:
getClinDT(
  data = tableAEGroup,
  colnames = tableAELabelsDT,
  fixedColumns = list(leftColumns = 1),
  columnsWidth = c(0.1, 0.7, 0.1, 0.1),
  width = "350px" # change dimension table
)

# change default buttons
getClinDT(
  data = tableAEGroup,
  colnames = tableAELabelsDT,
  # remove general filter
  filter = "none",
  # custom set of buttons
  buttons = c("csv", "excel", "pdf"),
  # change number of records displayed at each page
  pageLength = 20
)

# hide the first column:
getClinDT(
  data = tableAEGroup,
  nonVisibleVar = "AESOC"
)

# with specific caption
library(htmltools)
caption <- tags$caption("Number of subjects with adverse events grouped by system organ class."),

getColorPalette

Get a color palette for clinical visualizations.

Description

Get a color palette of specified length, either from a vector of names for the palette, or from a
specified length.

Usage

g getColorPalette(n = NULL, x = NULL, includeNA = FALSE, palette = clinColors)

Arguments

n  Integer of length 1, number of elements in palette.

x  Vector with elements used for palette. If factor, the levels are used, otherwise
the unique elements of the vector. Missing values are automatically removed,
excepted if includeNA is set to TRUE.

includeNA Logical (FALSE by default), should NA elements be retained in the palette in
 case x is specified?

palette  A vector of custom colors, or a function returning this vector from a specific
number of colors.
 Default is the the colorblind viridis color palette.

Value

Vector of colors, named with the elements in x if x is specified.

Author(s)

Laure Cougnaud and Michela Pasetto
getLabelParamcd

Examples

# extract longest palette available
getColorPalette(n = 11)
# extract palette for a vector
getColorPalette(x = paste('treatment', 1:4))
# possibility to include missing values:
getColorPalette(x = c(NA_character_, "group1"), includeNA = FALSE)
getColorPalette(x = c(NA_character_, "group1"), includeNA = TRUE)
# change default settings
getColorPalette(n = 3, palette = c("red", "green", "grey"))

getLabelParamcd

Get label for a parameter code

Description

This function gets the label for a parameter code extracted from the 'PARAM' column.

Usage

getLabelParamcd(paramcd, data, paramcdVar = "PARAMCD", paramVar = "PARAM")

Arguments

paramcd Character vector with parameter code(s).
data Data.frame with data.
paramcdVar String with column containing the paramcd parameter, 'PARAMCD' by default
(for ADaM format).
paramVar String with column containing the param parameter, 'PARAM' by default(for
ADaM format).

Value

Named character vector with label for parameter code or paramcd if label is missing.

Author(s)

Laure Cougnaud

Examples

# for ADaM
data(dataADaMCDISCP01)
getLabelParamcd(paramcd = "CHOL", data = dataADaMCDISCP01$ADLBC)
# for SDTM
data(dataSDTMCDISCP01)
getLabelParamcd(paramcd = "ALB", ...)
getData = dataSDTMCDISCP01$LB,
paramcdVar = "LBTESTCD",
paramVar = "LBTEST"
)

---

data getLabelVar

Get label for a variable of the dataset

Description

The label is extracted either (in this order):

1. if label is specified: from this label based on names, or directly from this label if label and var are of length 1 (if available)
2. if labelVars is specified: from the specified vector of labels, based on names (if available)
3. if data is specified: from the 'label' attribute of the corresponding column in data (if available)

If the label is not available, the input variable is returned.

Usage

getLabelVar(var, data = NULL, labelVars = NULL, label = NULL)

Arguments

var Character vector with variables of interest.
data Data frame with data.
labelVars Named character vector with variable labels (names are the variable code), usually extracted from data.
label (Named) Character vector with user-specified label for var. Label is extracted based on names if variable is available. If var is of length 1, label can also be specified as an unnamed character.

Value

Named character vector with label, var is no label is available

Author(s)

Laure Cougnaud
Examples

data(dataADaMCDISCP01)
labelVars <- attr(dataADaMCDISCP01, "labelVars")

# (upon reading the data with haven: attributes should directly available in each column)
getLabelVar(data = dataADaMCDISCP01, var = "AEREL")

# but if the data as data.frame is subsetted, label is lost
# so better to use 'labelVars':
getLabelVar(var = "AEREL", labelVars = labelVars)

data(dataADaMCDISCP01)
labelVars <- attr(dataADaMCDISCP01, "labelVars")

# extract label for all variables from specified datasets:
getLabelVars(data = dataADaMCDISCP01[c("ADLBC", "ADVS")], labelVars = labelVars)

# extracted from specified labelVars, e.g. to specify custom label for specific variable(s)
labelVarsCustom <- getLabelVars(
  data = dataADaMCDISCP01,
  labelVars = c(USUBJID = "Subject identifier for my study")
)
labelVarsCustom["USUBJID"]

Description
Get label of the variables in SAS dataset(s)

Usage
getLabelVars(data, labelVars = NULL)

Arguments
  data  Data.frame with SAS dataset(s) or list of those.
  labelVars  (optional) Named character vector with additional labels.

Value
Named vector with variable labels.

Author(s)
Laure Cougnaud

Examples

getLinetypePalette

Get a linetype palette for clinical visualizations.

Description

Get a linetype palette of specified length, either from a vector of names for the palette, or from a specified length.

Usage

getLinetypePalette(
  n = NULL,
  x = NULL,
  includeNA = FALSE,
  palette = clinLinetypes
)

Arguments

n       Integer of length 1, number of elements in palette.

x       Vector with elements used for palette. If factor, the levels are used, otherwise the unique elements of the vector. Missing values are automatically removed, excepted if includeNA is set to TRUE.

includeNA Logical (FALSE by default), should NA elements be retained in the palette in case x is specified?

palette A vector of custom linetypes, or a function returning this vector from a specific number of linetypes. Default is the clinLinetypes linetype palette.

Details

Note that 7 unique symbols are available at maximum (replicated if necessary).

Value

Vector with linetypes, named with the elements in x if x is specified.

Author(s)

Laure Cougnaud and Michela Pasetto
getPaletteCDISC

Examples

# extract longest linetype palette available
getLinetypePalette(n = 6)
# extract palette for a vector
getLinetypePalette(x = paste('treatment', 1:4))
# include missing
getLinetypePalette(x = c(NA_character_, "group1"), includeNA = TRUE)
getLinetypePalette(x = c(NA_character_, "group1"), includeNA = FALSE)
# set custom linetypes
lty <- getColorPalette(n = 3, palette = c("twodash", "dashed"))

getPaletteCDISC  Get standard palette for typical CDISC variables.

Description

The extraction of the palette elements is case-insensitive.

Usage

getPaletteCDISC(x, var, type, palette = NULL)

Arguments

x      Character vector of factor with variable to consider. The palette is built based on the unique elements of this vector, or levels if x is a factor.
var    String with type of variable, among:
        • 'NRIND': Normal Reference Range Indicator
        • 'shape': shape/symbol palette
type   String with type of palette:
        • 'shape': shape/symbol palette
palette (optional) Named vector with extra palette, e.g. to specify elements for non-standard categories. This palette is combined with the standard palette.

Details

The order of the palette depends on the type of the input variable (x):

• if a factor is specified, the palette is ordered based on its levels
• if a character vector is specified, the elements from the internal standard palette are used first, the remaining elements are then sorted alphabetically.

Value

Named vector with palette.
getSetDiff

Description
Get only additions and removals from two data sets (data.table objects). The additions/removals are extracted as x vs y. This function assumes that the objects x and y don’t share identical rows.

Usage
getSetDiff(x, y, referenceVars)

Arguments
x A data.table object
y A data.table object
referenceVars character vector of the columns in the data that are the used as reference for the comparison. If not specified, all columns present both in newData and oldData are considered.

Value
A data.table object with the additions/removals with respect of the comparison between x vs y.
getShapePalette

Get a shape palette for clinical visualizations.

Description

Get a shape palette of specified length, either from a vector of names for the palette, or from a specified length.

Usage

getShapePalette(
  n = NULL,
  x = NULL,
  includeNA = FALSE,
  asText = FALSE,
  palette = if (asText) { clinShapesText } else { clinShapes }
)

Arguments

n
  Integer of length 1, number of elements in palette.

x
  Vector with elements used for palette. If factor, the levels are used, otherwise the unique elements of the vector. Missing values are automatically removed, excepted if includeNA is set to TRUE.

includeNA
  Logical (FALSE by default), should NA elements be retained in the palette in case x is specified?

asText
  Logical (FALSE by default), should the palette be expressed as integer (base R plot and ggplot2 compatible) or in text format (e.g. required if combined with unicode symbols in ggplot2)?

palette
  A vector of custom shapes, or a function returning this vector from a specific number of shapes. The vector should be a character if asText is set to TRUE. Default is the clinShapes shape palette, or clinShapesText if asText is set to TRUE.

Details

Note that 19 unique symbols are available at maximum (replicated if necessary).

Value

Vector of shapes, named with the elements in x if x is specified.

Author(s)

Laure Cougnaud and Michela Pasetto
knitPrintListObjects

Include a list of objects in a knitr document

Description

This functions includes a list of objects in a knitr document allowing different chunk options (e.g. flextable object).

Usage

knitPrintListObjects(
  xList,
  generalLabel = "objectsList",
  labels = paste0(generalLabel, seq_along(xList)),
  titles = NULL,
  titleLevel = 2,
  printObject = FALSE,
  ...
)

Arguments

  xList List of objects to print.
  generalLabel String with general label for the chunks, used to build the labels. The labels are constructed as 'generalLabel[i]', with i the list index. Only use if labels is not specified.
  labels Character vector with labels, one for each chunk.
  titles Character vector with section titles, one for each chunk.
  titleLevel Integer with level for section header, 1 for top-level section header.
  printObject Logical, if TRUE (FALSE by default), each object within xList is explicitely printed with the print function.
  ... any chunk parameters, will be replicated if necessary, at the exception of 'results', set to 'asis' and 'echo' set to FALSE internally. See knitr[opts_chunk] for further details on available options.
Details

Note: the chunk in which this function is called should be set with the following option: results = 'asis'.

Value

No returned value, a text is printed with chunk content

Author(s)

Laure Cougnaud

Examples

## Not run:

```r
# Note: the following code should be included within a chunk of a knitr (e.g. RMarkdown) document
# to include/export a list of figures in the Rmarkdown output
data(iris)

# include a list of static plots
library(ggplot2)
plotsListStatic <- list(
  point = ggplot(data = cars, aes(x = speed, y = dist)) + geom_point(),
  line = ggplot(data = cars, aes(x = speed, y = dist)) + geom_line()
)
# with general label (used to name exported figure)
knitPrintListPlots(
  plotsList = plotsListStatic,
  generalLabel = "scatter-cars"
)
# with label for each plot (used to name exported figure)
knitPrintListPlots(
  plotsList = plotsListStatic,
  labels = names(plotsListStatic)
)
# with section header (header of level 1 in Markdown)
knitPrintListPlots(
  plotsList = plotsListStatic,
  titles = names(plotsListStatic),
  titleLevel = 3
)
# specify caption and dimension for each figure (figure height is replicated)
knitPrintListPlots(
  plotsList = plotsListStatic,
  fig.cap = names(plotsListStatic),
  fig.width = 3*seq_along(plotsListStatic),
  fig.height = 6
)
```

# include a list of interactive plots
library(plotly)
knitPrintListPlots <- list(
  point = plot_ly(data = cars, x = ~speed, y = ~dist, type = "scatter", mode = "marker"),
  line = plot_ly(data = cars, x = ~speed, y = ~dist, type = "scatter", mode = "line")
)
knitPrintListPlots(
  plotsList = plotsListInteractive,
  type = "plotly",
  titles = names(plotsListInteractive),
  titleLevel = 3
)

## End(Not run)
## Not run:
# Note: the following code should be included within a chunk of a knitr (e.g. RMarkdown) document
# to include/export a list of figures in the Rmarkdown output
# list of flextable objects
library(flextable)
listTables <- list(flextable(iris), flextable(cars))
knitPrintListObjects(xList = listTables)

## End(Not run)

---

**knitPrintListPlots**  
*Include a list of plots in a knitr document*

---

**Description**

This function includes a list of plots with different figure height, size, label in a knitr document. The reason is that the chunk options `fig.height` and `fig.width` are not vectorized.

**Usage**

```r
knitPrintListPlots(
  plotsList,
  generalLabel = "plotsList",
  type = c("ggplot2", "plotly"),
  ...
)
```

**Arguments**

- `plotsList`  
  list of plots, e.g. ggplot objects from the ggplot2 package or from the plotly packages.

- `generalLabel`  
  general label for the chunks, used to build the labels. The labels are constructed as `generalLabel[i]`, with `i` the plot number (from sequence spanning the length of `plotsList`). Only use if `labels` is not specified.
### knitPrintListPlots

**type** string with plot type: ‘ggplot2’ or ‘plotly’

Additional parameters for the `knitPrintListObjects` function.

### Details

Note: the chunk in which this function is called should be set with the following option: `results = 'asis'`.

Note that a (one-level) list of plotly plots can also be included directly via `htmltools::tagList(listPlots)`, but without the possibility to specify title and/or change chunk option(s). Each plot will be included in a separated chunk.

### Value

No returned value, a text is printed with chunk content

### Author(s)

Laure Cougnaud

### Examples

```r
## Not run:
# Note: the following code should be included within a chunk of a knitr (e.g. RMarkdown) document
# to include/export a list of figures in the Rmarkdown output
data(iris)

# include a list of static plots
library(ggplot2)
plotsListStatic <- list(
  point = ggplot(data = cars, aes(x = speed, y = dist)) + geom_point(),
  line = ggplot(data = cars, aes(x = speed, y = dist)) + geom_line()
)
# with general label (used to name exported figure)
knitPrintListPlots(
  plotsList = plotsListStatic,
  generallabel = "scatter-cars"
)
# with label for each plot (used to name exported figure)
knitPrintListPlots(
  plotsList = plotsListStatic,
  labels = names(plotsListStatic)
)
# with section header (header of level 1 in Markdown)
knitPrintListPlots(
  plotsList = plotsListStatic,
  titles = names(plotsListStatic),
  titleLevel = 3
)
# specify caption and dimension for each figure (figure height is replicated)
knitPrintListPlots(
```

```r
plotsList = plotsListStatic,
fig.cap = names(plotsListStatic),
fig.width = 3*seq_along(plotsListStatic),
fig.height = 6
)

# include a list of interactive plots
library(plotly)
plotsListInteractive <- list(
    point = plot_ly(data = cars, x = ~speed, y = ~dist, type = "scatter", mode = "marker"),
    line = plot_ly(data = cars, x = ~speed, y = ~dist, type = "scatter", mode = "line"
)
)
knitPrintListPlots(
    plotsList = plotsListInteractive,
    type = "plotly",
    titles = names(plotsListInteractive),
    titleLevel = 3
)

## End(Not run)

---

**loadDataADaMSDTM**  Load data from ADaM/SDTM file(s).

**Description**

Load data set from SAS format into R data.frames.

**Usage**

```r
loadDataADaMSDTM(
    files,
    convertToDate = FALSE,
    dateVars = "DTC$",
    verbose = TRUE,
    encoding = "UTF-8",
    ...
)
```

**Arguments**

- `files`  Character vector with path to ADaM or SDTM file(s). Currently only import of files with extension: 'sas7bdat' or 'xpt' are supported.
- `convertToDate`  logical, if TRUE columns with date/time are converted to POSIXct format, which stores calendar date/time in R. Please note that most of the time this is not necessary, as date variables are automatically imported via the haven package if encoded correctly in the dataset.
**mergeDiffWithData**

Merge the 'diff.data' object from `compareDiff` with the original `newData` or `oldData`.

**Description**

The `newData`/`oldData` are merged with `diffData` based on the columns of `diffData` excepted 'Comparison type' and 'Version'.

**Usage**

```r
mergeDiffWithData(diffData, newData, oldData)
```
Arguments

diffData Object of class 'diff.data' containing differences between datasets, as returned by the `compareDiff` function.

newData data.frame object representing the new data

oldData data.frame object representing the old data

Value

The `newData` or `oldData` (as a data frame object) with the extra column 'Comparison type' specifying the type of change, either:

- 'Change': record present in both dataset based on the reference variables, but with changes in the changeable variables
- 'Addition': records present in new but not in old data
- 'Removal': records present in old but not in new data
- 'Identical': records identical in the old and new datasets (on both the reference and changeable variables)

Description

Custom merge (left join) of difference data with some input data

Usage

```
mergeInputDiff(diffData, inputData, typeData, colsBy)
```

Arguments

diffData A data.table object as output from `compareDiff`.

inputData A data.table object. For instance, the `newData` or the `oldData` argument from `compareTables`.

typeData String with type of data, as "new" for `newData` or "old" for `oldData`.

colsBy Character vector of columns for doing the merge by.

Value

A data.table object. The `inputData` is joined with the columns 'Comparison type' and 'Version' from the `diffData` argument.
**reorderColumns**

*Function for reordering columns*

**Description**

Function for reordering columns

**Usage**

```r
reorderColumns(data, vars)
```

**Arguments**

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>data</td>
<td>A data.frame</td>
</tr>
<tr>
<td>vars</td>
<td>Named vector indicating the position in the data frame of the specified variable</td>
</tr>
</tbody>
</table>

**Value**

The same data.frame specified in data, with ordered columns.

**Examples**

```r
someData <- data.frame(
  "Col1" = c(1, 2),
  "Col2" = c(2, 3),
  "Col3" = c(3, 4)
)
reorderColumns(
  data = someData,
  vars = c("Col3" = 1)
)
```

---

**roundHalfUp**

*Round a number with 'rounding up' strategy for rounding off a 5*

**Description**

This function rounds a number for a specified number of digits. It rounds off to the highest number for a 5. The default R `round` function rounds to the 'even digit' in case of rounding off a 5 (see 'Details' section in ? round). This function instead rounds up to the nearest number for a 5. It mimics a similar rounding strategy used in SAS. See examples for the difference between `round` and `roundHalfUp` below.

**Usage**

```r
roundHalfUp(x, digits = 0)
```
Arguments

- `x` Numeric vector to round.
- `digits` Integer with number of digits to consider, 0 by default.

Value

Rounded numeric vector.

Author(s)

stackoverflow question 6461209

Examples

```r
# numbers are rounded to the closest even number in case of .5
# with the round 'base' function
round(0.45, 1)
# 'roundHalfUp' always round to the next highest number in case of .5
roundHalfUp(0.45, 1)
# rounding is the same for uneven number:
round(0.55, 1)
roundHalfUp(0.55)
# other examples
round(1.456e-2, digits = 3)
round(1.456e-2, digits = 2)
round(1.456e-2, digits = 1)
```

Description

This function rounds numbers with a 'round-up' strategy for rounding off a 5. The function rounds for a specified number of digits and format number to a: 'xxx.xxx' text.

Usage

```r
roundHalfUpTextFormat(x, digits = 0)
```

Arguments

- `x` Numeric vector to round.
- `digits` Integer with number of digits to consider, 0 by default.
Details

The following workflow is used:

1. numbers are rounded with the `roundHalfUp` function, see the `? roundHalfUp` for more details on the rounding strategy
2. round numbers are formatted to character in the format: 'xxx.xxx' with pads leading zeros

Value

A character vector with the rounded number. NA values are returned as 'NA' as string.

Author(s)

Laure Cougnaud and Michela Pasetto

See Also

`roundHalfUp` for the rounding customization.

Examples

```r
# number of digits higher than number of decimal
droundHalfUpTextFormat(x = c(0.345, 0.567, -0.98), digits = 2)
# number of digits lower than number of decimal
droundHalfUpTextFormat(x = c(0.345, 0.567, -0.98), digits = 0)
# by default, 'digits' is 0!
droundHalfUpTextFormat(x = c(0.345, 0.567, -0.98))
# padding zeros
droundHalfUpTextFormat(1.23, 10)
```

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**shapePaletteNRIND**

*Shape palette for a standard CDISC Normal/Reference Range Indicator.*

Description

These symbols should be supported in Windows and Linux.

Usage

`shapePaletteNRIND`
Format

A named character vector with shape symbol for typical Normal Reference Range Indicator variable:

- "LOW": filled down-pointing arrow (25)
- "NORMAL": filled circle (21)
- "HIGH": filled up-pointing arrow (24)
- "ABNORMAL": diamond (18)
- "UNKNOWN" or 'NA': cross (3)
- "NA": cross (3)

simpleCap

Capitalize the first letter of a word/sentence.

Description

This implementation is inspired from the help of the toupper function.

Usage

simpleCap(x, onlyFirst = TRUE, rev = FALSE)

Arguments

x Character vector to capitalize

onlyFirst Logical, if TRUE (by default) capitalize the first letter of the first word only. Otherwise, capitalize the first letters of all words of the sentence. See also link[tools]{toTitleCase} for a more syntax-friendly implementation.

rev Logical, if TRUE (FALSE by default), set first letter to lower case (otherwise upper case)

Value

Character vector with first letter capitalized

Author(s)

author of the 'toupper' function?

See Also

link[tools]{toTitleCase}
Examples

# capitalize only the first word of the sentence
simpleCap(x = "this is the caption of my figure.")

# capitalize all words
simpleCap(x = "this is the caption of my figure.", onlyFirst = FALSE)

# opposite: set the first letter of the first word to lower case
simpleCap(x = "This is the caption of my figure.", rev = TRUE)
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