

Package ‘explore’

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Type Package

Title Simplifies Exploratory Data Analysis

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Description Interactive data exploration with one line of code or use an easy to remember set of tidy functions for exploratory data analysis. Introduces three main verbs. `explore()` to graphically explore a variable or table, `describe()` to describe a variable or table and `report()` to create an automated report.

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Encoding UTF-8

URL <https://github.com/rolkra/explore/>

Imports assertthat, broom, dplyr, DT, forcats, ggplot2 (>= 3.0.0), gridExtra, magrittr, MASS, rlang, rmarkdown, rpart, rpart.plot, shiny, stats, stringr, tibble, tidyr

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VignetteBuilder knitr

NeedsCompilation no

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R topics documented:

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| | |
|----------------|---|
| add_random_var | <i>Add a random variable to dataset</i> |
|----------------|---|

Description

Add a random variable to dataset

Usage

```
add_random_var(  
    data,  
    name = "random",  
    min_val = 0,  
    max_val = 1,  
    overwrite = TRUE  
)
```

Arguments

| | |
|-----------|--|
| data | A dataset |
| name | Name of new variable (as string) |
| min_val | Minimum random integers |
| max_val | Maximum random integers |
| overwrite | Can new random variable overwrite an existing variable in dataset? |

Value

Dataset containing new random variable

Examples

```
add_random_var(iris)  
add_random_var(iris, name = "random_var")  
add_random_var(iris, min_val = 1, max_val = 10)  
add_random_var(iris, min_val = 1, max_val = 100, overwrite = FALSE)
```

| | |
|----------------|--------------------------------|
| balance_target | <i>Balance target variable</i> |
|----------------|--------------------------------|

Description

Balances the target variable in your dataset. Target must be 0/1, FALSE/TRUE ore no/yes

Usage

```
balance_target(data, target, min_prop = 0.1)
```

Arguments

| | |
|----------|--|
| data | A dataset |
| target | Target variable (0/1, TRUE/FALSE, yes/no) |
| min_prop | Minimum proportion of one of the target categories |

Value

Data

Examples

```
iris$is_versicolor <- ifelse(iris$Species == "versicolor", 1, 0)
balanced <- balance_target(iris, target = is_versicolor, min_prop = 0.5)
describe(balanced, is_versicolor)
```

| | |
|-----------|-----------------------|
| clean_var | <i>Clean variable</i> |
|-----------|-----------------------|

Description

Clean variable (replace NA values, set min_val and max_val)

Usage

```
clean_var(
  data,
  var,
  na = NA,
  min_val = NA,
  max_val = NA,
  max_cat = NA,
  rescale01 = FALSE,
  simplify_text = FALSE,
  name = NA
)
```

Arguments

| | |
|---------|---|
| data | A dataset |
| var | Name of variable |
| na | Value that replaces NA |
| min_val | All values < min_val are converted to min_val (var numeric or character) |
| max_val | All values > max_val are converted to max_val (var numeric or character) |
| max_cat | Maximum number of different factor levels for categorical variable (if more, .OTHER is added) |

| | |
|---------------|--|
| rescale01 | Rescale into value between 0 and 1 (var must be numeric) |
| simplify_text | if TRUE, a character variable is simplified (trim, upper, ...) |
| name | New name of variable (as string) |

Value

Dataset

Examples

```
clean_var(iris, Sepal.Width, max_val = 3.5, name = "sepal_width")
```

| | |
|-----------|--|
| count_pct | <i>Adds percentage to dplyr::count()</i> |
|-----------|--|

Description

Adds variables total and pct (percentage) to dplyr::count()

Usage

```
count_pct(data, ...)
```

Arguments

| | |
|------|------------------------------------|
| data | A dataset |
| ... | Other parameters passed to count() |

Value

Dataset

Examples

```
count_pct(iris, Species)
```

create_fake_data *Create fake data*

Description

Fake data that can be used for unit-testing or demonstration

Usage

```
create_fake_data(  
  obs = 1000,  
  target_name = "target_ind",  
  factorise_target = FALSE,  
  target1_prob = 0.5,  
  add_extreme = TRUE,  
  flip_gender = FALSE,  
  add_id = FALSE,  
  seed = 123  
)
```

Arguments

| | |
|------------------|--|
| obs | Number of observations |
| target_name | Variable name of target |
| factorise_target | Should target variable be factorised? (from 0/1 to facotr no/yes)? |
| target1_prob | Probability that buy = 1 |
| add_extreme | Add an observation with extreme values? |
| flip_gender | Should Male/Female be flipped in data? |
| add_id | Add an id-variable to data? |
| seed | Seed for randomization |

Details

Variables in dataset:

- id = Identifier
- period = Year & Month (YYYYMM)
- city_ind = Indicating if customer is residing in a city (1 = yes, 0 = no)
- female_ind = Gender of customer is female (1 = yes, 0 = no)
- fixedvoice_ind = Customer has a fixed voice product (1 = yes, 0 = no)
- fixeddata_ind = Customer has a fixed data product (1 = yes, 0 = no)
- fixedtv_ind = Customer has a fixed tv product (1 = yes, 0 = no)

- mobilevoice_ind = Customer has a mobile voice product (1 = yes, 0 = no)
- mobiledata_ind = Customer has a mobile data product (1 = yes, 0 = no)
- bbi_speed_ind = Customer has a Broadband Internet (BBI) with speed
- bbi_usg_gb = Broadband Internet (BBI) usage in Gigabyte (GB) last month
- hh_single = Expected to be a Single Household (1 = yes, 0 = no)

Target in dataset:

- target_ind (may be renamed) = Did customer buy a new product in next month? (1 = yes, 0 = no)

Value

A dataframe

| | |
|--------------------|---------------------------|
| create_random_data | <i>Create random data</i> |
|--------------------|---------------------------|

Description

Random data that can be used for unit-testing or demonstration

Usage

```
create_random_data(
  obs = 1000,
  vars = 10,
  target_name = "target_ind",
  factorise_target = FALSE,
  target1_prob = 0.5,
  add_id = TRUE,
  seed = 123
)
```

Arguments

| | |
|------------------|--|
| obs | Number of observations |
| vars | Number of variables |
| target_name | Variable name of target |
| factorise_target | Should target variable be factorised? (from 0/1 to facotr no/yes)? |
| target1_prob | Probability that buy = 1 |
| add_id | Add an id-variable to data? |
| seed | Seed for randomization |

Details

Variables in dataset:

- id = Identifier
- var_X = variable containing values between 0 and 100

Target in dataset:

- target_ind (may be renamed) = random values (1 = yes, 0 = no)

Value

A dataframe

| | |
|--------------|---|
| data_dict_md | <i>Create a data dictionary Markdown file</i> |
|--------------|---|

Description

Create a data dictionary Markdown file

Usage

```
data_dict_md(  
  data,  
  title = "",  
  description = NA,  
  output_file = "data_dict.md",  
  output_dir  
)
```

Arguments

| | |
|-------------|---|
| data | A dataframe (data dictionary for all variables) |
| title | Title of the data dictionary |
| description | Detailed description of variables in data (dataframe with columns 'variable' and 'description') |
| output_file | Output filename for Markdown file |
| output_dir | Directory where the Markdown file is saved |

Value

Create Markdown file

Examples

```
# Data dictionary of a dataframe
data_dict_md(iris,
             title = "iris flower data set",
             output_dir = tempdir())

# Data dictionary of a dataframe with additional description of variables
description <- data.frame(
  variable = c("Species"),
  description = c("Species of Iris flower"))
data_dict_md(iris,
             title = "iris flower data set",
             description = description,
             output_dir = tempdir())
```

decrypt

decrypt text

Description

decrypt text

Usage

```
decrypt(text, codeletters = c(toupper(letters), letters, 0:9), shift = 18)
```

Arguments

| | |
|-------------|--|
| text | A text (character) |
| codeletters | A string of letters that are used for decryption |
| shift | Number of elements shifted |

Value

Decrypted text

Examples

```
decrypt("zw336 E693v")
```

| | |
|----------|---------------------------------------|
| describe | <i>Describe a dataset or variable</i> |
|----------|---------------------------------------|

Description

Describe a dataset or variable (depending on input parameters)

Usage

```
describe(data, var, n, target, out = "text", ...)
```

Arguments

| | |
|--------|---|
| data | A dataset |
| var | A variable of the dataset |
| n | Weights variable for count-data |
| target | Target variable (0/1 or FALSE/TRUE) |
| out | Output format ("text" "list") of variable description |
| ... | Further arguments |

Value

Description as table, text or list

Examples

```
# Load package
library(magrittr)

# Describe a dataset
iris %>% describe()

# Describe a variable
iris %>% describe(Species)
iris %>% describe(Sepal.Length)
```

| | |
|--------------|--|
| describe_all | <i>Describe all variables of a dataset</i> |
|--------------|--|

Description

Describe all variables of a dataset

Usage

```
describe_all(data = NA, out = "large")
```

Arguments

| | |
|------|---------------------------------|
| data | A dataset |
| out | Output format ("small" "large") |

Value

Dataset (tibble)

Examples

```
describe_all(iris)
```

| | |
|--------------|--------------------------------------|
| describe_cat | <i>Describe categorical variable</i> |
|--------------|--------------------------------------|

Description

Describe categorical variable

Usage

```
describe_cat(data, var, n, max_cat = 10, out = "text", margin = 0)
```

Arguments

| | |
|---------|--|
| data | A dataset |
| var | Variable or variable name |
| n | Weights variable for count-data |
| max_cat | Maximum number of categories displayed |
| out | Output format ("text" "list") |
| margin | Left margin for text output (number of spaces) |

Value

Description as text or list

Examples

```
describe_cat(iris, Species)
```

| | |
|--------------|------------------------------------|
| describe_num | <i>Describe numerical variable</i> |
|--------------|------------------------------------|

Description

Describe numerical variable

Usage

```
describe_num(data, var, n, out = "text", margin = 0)
```

Arguments

| | |
|--------|--|
| data | A dataset |
| var | Variable or variable name |
| n | Weights variable for count-data |
| out | Output format ("text" "list") |
| margin | Left margin for text output (number of spaces) |

Value

Description as text or list

Examples

```
describe_num(iris, Sepal.Length)
```

| | |
|--------------|-----------------------|
| describe_tbl | <i>Describe table</i> |
|--------------|-----------------------|

Description

Describe table (e.g. number of rows and columns of dataset)

Usage

```
describe_tbl(data, n, target, out = "text")
```

Arguments

| | |
|--------|---------------------------------|
| data | A dataset |
| n | Weights variable for count-data |
| target | Target variable (binary) |
| out | Output format ("text" "list") |

Value

Description as text or list

Examples

```
describe_tbl(iris)

iris$is_virginica <- ifelse(iris$Species == "virginica", 1, 0)
describe_tbl(iris, is_virginica)
```

| | |
|-------------|-----------------------|
| dwh_connect | <i>connect to DWH</i> |
|-------------|-----------------------|

Description

connect to datawarehouse (DWH) using ODBC

Usage

```
dwh_connect(dsn, user = NA, pwd = NA, pwd_crypt = FALSE, ...)
```

Arguments

| | |
|-----------|--|
| dsn | DSN string |
| user | user name |
| pwd | password of user |
| pwd_crypt | is password encryption used? |
| ... | Further arguments to be passed to DBI::dbConnect() |

Value

connection

Examples

```
## Not run:  
con <- dwh_connect(dsn = "DWH1", user = "u12345")  
  
## End(Not run)
```

| | |
|----------------|----------------------------|
| dwh_disconnect | <i>disconnect from DWH</i> |
|----------------|----------------------------|

Description

disconnect from datawarehouse (DWH) using a ODBC connection

Usage

```
dwh_disconnect(connection, ...)
```

Arguments

| | |
|------------|---|
| connection | channel (ODBC connection) |
| ... | Further arguments to be passed to DBI::dbDisconnect() |

Examples

```
## Not run:  
dwh_disconnect(con)  
  
## End(Not run)
```

| | |
|--------------|----------------------------------|
| dwh_fastload | <i>write data to a DWH table</i> |
|--------------|----------------------------------|

Description

write data fast to a DWH table using a ODBC connection Function uses packages DBI/odbc to write data faster than RODBC Connects, writes data and disconnects

Usage

```
dwh_fastload(data, dsn, table, overwrite = FALSE, append = FALSE, ...)
```

Arguments

| | |
|-----------|--|
| data | dataframe |
| dsn | DSN string |
| table | table name (character string) |
| overwrite | Overwrite table if already exist |
| append | Append data to table |
| ... | Further arguments to be passed to DBI::dbConnect() |

Value

status

Examples

```
## Not run:
dwh_fastload(data, "DWH", "database.table_test")

## End(Not run)
```

| | |
|---------------|---------------------------|
| dwh_read_data | <i>read data from DWH</i> |
|---------------|---------------------------|

Description

read data from DWH using a ODBC connection

Usage

```
dwh_read_data(connection, sql, names_lower = TRUE, ...)
```

Arguments

| | |
|-------------|---|
| connection | DWH connection |
| sql | sql (character string) |
| names_lower | convert field names to lower (default = TRUE) |
| ... | Further arguments to be passed to DBI::dbGetQuery() |

Value

dataframe containing table data

Examples

```
## Not run:
dwh_read_data(con, "select * from database.table_test")

## End(Not run)
```

| | |
|----------------|------------------------------|
| dwh_read_table | <i>read a table from DWH</i> |
|----------------|------------------------------|

Description

read a table from DWH using a ODBC connection

Usage

```
dwh_read_table(connection, table, names_lower = TRUE, ...)
```

Arguments

| | |
|-------------|---|
| connection | DWH connection |
| table | table name (character string) |
| names_lower | convert field names to lower (default = TRUE) |
| ... | Further arguments to be passed to DBI::dbGetQuery() |

Value

dataframe containing table data

Examples

```
## Not run:
dwh_read_table(con, "database.table_test")

## End(Not run)
```

| | |
|---------|---------------------|
| encrypt | <i>encrypt text</i> |
|---------|---------------------|

Description

encrypt text

Usage

```
encrypt(text, codeletters = c(toupper(letters), letters, 0:9), shift = 18)
```

Arguments

| | |
|-------------|--|
| text | A text (character) |
| codeletters | A string of letters that are used for encryption |
| shift | Number of elements shifted |

Value

Encrypted text

Examples

```
encrypt("hello world")
```

| | |
|----------------|--|
| explain_logreg | <i>Explain a binary target using a logistic regression (glm). Model chosen by AIC in a Stepwise Algorithm (MASS::stepAIC).</i> |
|----------------|--|

Description

Explain a binary target using a logistic regression (glm). Model chosen by AIC in a Stepwise Algorithm (MASS::stepAIC).

Usage

```
explain_logreg(data, target, out = "tibble", ...)
```

Arguments

| | |
|--------|--|
| data | A dataset |
| target | Target variable (binary) |
| out | Output of the function: "tibble" "model" |
| ... | Further arguments |

Value

Dataset with results (term, estimate, std.error, z.value, p.value) or the model (if out = "model")

Examples

```
data <- iris
data$is_versicolor <- ifelse(iris$Species == "versicolor", 1, 0)
data$Species <- NULL
explain_logreg(data, target = is_versicolor)
```

| | |
|--------------|---|
| explain_tree | <i>Explain a target using a simple decision tree (classification or regression)</i> |
|--------------|---|

Description

Explain a target using a simple decision tree (classification or regression)

Usage

```
explain_tree(
  data,
  target,
  n,
  max_cat = 10,
  max_target_cat = 5,
  maxdepth = 3,
  minsplit = NA,
  cp = 0,
  weights = NA,
  size = 0.7,
  out = "plot",
  ...
)
```

Arguments

| | |
|----------------|---|
| data | A dataset |
| target | Target variable |
| n | weights variable (for count data) |
| max_cat | Drop categorical variables with higher number of levels |
| max_target_cat | Maximum number of categories to be plotted for target (except NA) |
| maxdepth | Maximal depth of the tree (rpart-parameter) |
| minsplit | The minimum number of observations that must exist in a node to split. |
| cp | Complexity parameter (rpart-parameter) |
| weights | Vector containing weight of each observation (rpart-parameter). Can not be used in combination with parameter n (variable containing weight for count-data) |
| size | Textsize of plot |
| out | Output of function: "plot" "model" |
| ... | Further arguments |

Value

Plot or additional the model (if out = "model")

Examples

```
data <- iris
data$is_versicolor <- ifelse(iris$Species == "versicolor", 1, 0)
data$Species <- NULL
explain_tree(data, target = is_versicolor)
```

 explore

Explore a dataset or variable

Description

Explore a dataset or variable

Usage

```
explore(
  data,
  var,
  var2,
  n,
  target,
  split,
  min_val = NA,
  max_val = NA,
  auto_scale = TRUE,
  na = NA,
  ...
)
```

Arguments

| | |
|------------|--|
| data | A dataset |
| var | A variable |
| var2 | A variable for checking correlation |
| n | A Variable for number of observations (count data) |
| target | Target variable (0/1 or FALSE/TRUE) |
| split | Split by target variable (FALSE/TRUE) |
| min_val | All values < min_val are converted to min_val |
| max_val | All values > max_val are converted to max_val |
| auto_scale | Use 0.2 and 0.98 quantile for min_val and max_val (if min_val and max_val are not defined) |
| na | Value to replace NA |
| ... | Further arguments (like flip = TRUE/FALSE) |

Value

Plot object

Examples

```
## Launch Shiny app (in interactive R sessions)
if (interactive()) {
  explore(iris)
}

## Explore grafically

# Load library
library(magrittr)

# Explore a variable
iris %>% explore(Species)
iris %>% explore(Sepal.Length)
iris %>% explore(Sepal.Length, min_val = 4, max_val = 7)

# Explore a variable with a target
iris$is_virginica <- ifelse(iris$Species == "virginica", 1, 0)
iris %>% explore(Species, target = is_virginica)
iris %>% explore(Sepal.Length, target = is_virginica)

# Explore correlation between two variables
iris %>% explore(Species, Petal.Length)
iris %>% explore(Sepal.Length, Petal.Length)

# Explore correlation between two variables and split by target
iris %>% explore(Sepal.Length, Petal.Length, target = is_virginica)
```

explore_all

Explore all variables

Description

Explore all variables of a dataset (create plots)

Usage

```
explore_all(data, n, target, ncol = 2, split = TRUE)
```

Arguments

| | |
|--------|--|
| data | A dataset |
| n | Weights variable (only for count data) |
| target | Target variable (0/1 or FALSE/TRUE) |

| | |
|-------|-------------------------------------|
| ncol | Layout of plots (number of columns) |
| split | Split by target (TRUE FALSE) |

Value

Plot

Examples

```
explore_all(iris)

iris$is_virginica <- ifelse(iris$Species == "virginica", 1, 0)
explore_all(iris, target = is_virginica)
```

| | |
|-------------|--|
| explore_bar | <i>Explore categorical variable using bar charts</i> |
|-------------|--|

Description

Create a barplot to explore a categorical variable. If a target is selected, the barplot is created for all levels of the target.

Usage

```
explore_bar(
  data,
  var,
  target,
  flip = NA,
  title = "",
  numeric = NA,
  max_cat = 30,
  max_target_cat = 5,
  legend_position = "right",
  label,
  label_size = 2.7,
  ...
)
```

Arguments

| | |
|--------|---|
| data | A dataset |
| var | variable |
| target | target (can have more than 2 levels) |
| flip | Should plot be flipped? (change of x and y) |
| title | Title of the plot (if empty var name) |

| | |
|-----------------|---|
| numeric | Display variable as numeric (not category) |
| max_cat | Maximum number of categories to be plotted |
| max_target_cat | Maximum number of categories to be plotted for target (except NA) |
| legend_position | Position of the legend ("bottom" "top" "none") |
| label | Show labels? (if empty, automatic) |
| label_size | Size of labels |
| ... | Further arguments |

Value

Plot object (bar chart)

| | |
|-------------|--|
| explore_cor | <i>Explore the correlation between two variables</i> |
|-------------|--|

Description

Explore the correlation between two variables

Usage

```
explore_cor(
  data,
  x,
  y,
  target,
  bins = 8,
  min_val = NA,
  max_val = NA,
  auto_scale = TRUE,
  title = NA,
  color = "grey",
  ...
)
```

Arguments

| | |
|---------|---|
| data | A dataset |
| x | Variable on x axis |
| y | Variable on y axis |
| target | Target variable (categorical) |
| bins | Number of bins |
| min_val | All values < min_val are converted to min_val |

| | |
|------------|--|
| max_val | All values > max_val are converted to max_val |
| auto_scale | Use 0.2 and 0.98 quantile for min_val and max_val (if min_val and max_val are not defined) |
| title | Title of the plot |
| color | Color of the plot |
| ... | Further arguments |

Value

Plot

Examples

```
explore_cor(iris, x = Sepal.Length, y = Sepal.Width)
```

| | |
|---------------|--|
| explore_count | <i>Explore count data (categories + frequency)</i> |
|---------------|--|

Description

Create a plot to explore count data (categories + frequency) Variable named 'n' is auto detected as Frequency

Usage

```
explore_count(
  data,
  cat,
  n,
  target,
  pct = FALSE,
  split = TRUE,
  title = NA,
  numeric = FALSE,
  max_cat = 30,
  max_target_cat = 5,
  flip = NA
)
```

Arguments

| | |
|--------|------------------------------------|
| data | A dataset (categories + frequency) |
| cat | Numerical variable |
| n | Number of observations (frequency) |
| target | Target variable |

| | |
|----------------|---|
| pct | Show as percent? |
| split | Split by target? |
| title | Title of the plot |
| numeric | Display variable as numeric (not category) |
| max_cat | Maximum number of categories to be plotted |
| max_target_cat | Maximum number of categories to be plotted for target (except NA) |
| flip | Flip plot? (for categorical variables) |

Value

Plot object

Examples

```
library(dplyr)
iris %>%
  count(Species) %>%
  explore_count(Species)
```

| | |
|-----------------|------------------------------------|
| explore_density | <i>Explore density of variable</i> |
|-----------------|------------------------------------|

Description

Create a density plot to explore numerical variable

Usage

```
explore_density(
  data,
  var,
  target,
  title = "",
  min_val = NA,
  max_val = NA,
  color = "grey",
  auto_scale = TRUE,
  max_target_cat = 5,
  ...
)
```


Arguments

| | |
|----------------|---|
| data | A dataset |
| var | Variable |
| target | Target variable (0/1 or FALSE/TRUE) |
| title | Title of the plot (if empty var name) |
| min_val | All values < min_val are converted to min_val |
| max_val | All values > max_val are converted to max_val |
| color | Color of plot |
| auto_scale | Use 0.02 and 0.98 percent quantile for min_val and max_val (if min_val and max_val are not defined) |
| max_target_cat | Maximum number of levels of target shown in the plot (except NA). |
| ... | Further arguments |

Value

Plot object (density plot)

Examples

```
explore_density(iris, "Sepal.Length")
iris$is_virginica <- ifelse(iris$Species == "virginica", 1, 0)
explore_density(iris, Sepal.Length, target = is_virginica)
```

explore_shiny

Explore dataset interactive

Description

Launches a shiny app to explore a dataset

Usage

```
explore_shiny(data, target)
```

Arguments

| | |
|--------|-------------------------------------|
| data | A dataset |
| target | Target variable (0/1 or FALSE/TRUE) |

Examples

```
# Only run examples in interactive R sessions
if (interactive()) {
  explore_shiny(iris)
}
```

explore_targetpct *Explore variable + binary target (values 0/1)*

Description

Create a plot to explore relation between a variable and a binary target as target percent. The target variable is chosen automatically if possible (name starts with 'target')

Usage

```
explore_targetpct(
  data,
  var,
  target = NULL,
  title = NULL,
  min_val = NA,
  max_val = NA,
  auto_scale = TRUE,
  na = NA,
  flip = NA,
  ...
)
```

Arguments

| | |
|------------|--|
| data | A dataset |
| var | Numerical variable |
| target | Target variable (0/1 or FALSE/TRUE) |
| title | Title of the plot |
| min_val | All values < min_val are converted to min_val |
| max_val | All values > max_val are converted to max_val |
| auto_scale | Use 0.2 and 0.98 quantile for min_val and max_val (if min_val and max_val are not defined) |
| na | Value to replace NA |
| flip | Flip plot? (for categorical variables) |
| ... | Further arguments |

Value

Plot object

Examples

```
iris$target01 <- ifelse(iris$Species == "versicolor",1,0)
explore_targetpct(iris)
```

| | |
|-------------|----------------------|
| explore_tbl | <i>Explore table</i> |
|-------------|----------------------|

Description

Explore a table. Plots variable types, variables with no variance and variables with NA

Usage

```
explore_tbl(data, n)
```

Arguments

| | |
|------|--------------------------------|
| data | A dataset |
| n | Weight variable for count data |

Examples

```
explore_tbl(iris)
```

| | |
|-----------------|---|
| format_num_auto | <i>Format number as character string (auto)</i> |
|-----------------|---|

Description

Formats a number depending on the value as number with space, scientific or big number as k (1 000), M (1 000 000) or B (1 000 000 000)

Usage

```
format_num_auto(number = 0, digits = 1)
```

Arguments

| | |
|--------|----------------------------|
| number | A number (integer or real) |
| digits | Number of digits |

Value

Formatted number as text

Examples

```
format_num_kMB(5500, digits = 2)
```

| | |
|----------------|--|
| format_num_kMB | <i>Format number as character string (kMB)</i> |
|----------------|--|

Description

Formats a big number as k (1 000), M (1 000 000) or B (1 000 000 000)

Usage

```
format_num_kMB(number = 0, digits = 1)
```

Arguments

| | |
|--------|----------------------------|
| number | A number (integer or real) |
| digits | Number of digits |

Value

Formatted number as text

Examples

```
format_num_kMB(5500, digits = 2)
```

| | |
|------------------|--|
| format_num_space | <i>Format number as character string (space as big.mark)</i> |
|------------------|--|

Description

Formats a big number using space as big.mark (1000 = 1 000)

Usage

```
format_num_space(number = 0, digits = 1)
```

Arguments

| | |
|--------|----------------------------|
| number | A number (integer or real) |
| digits | Number of digits |

Value

Formatted number as text

Examples

```
format_num_space(5500, digits = 2)
```

| | |
|---------------|----------------------|
| format_target | <i>Format target</i> |
|---------------|----------------------|

Description

Formats a target as a 0/1 variable. If target is numeric, 1 = above average.

Usage

```
format_target(target)
```

Arguments

| | |
|--------|--------------------|
| target | Variable as vector |
|--------|--------------------|

Value

Formatted target

Examples

```
iris$is_virginica <- ifelse(iris$Species == "virginica", "yes", "no")
iris$target <- format_target(iris$is_virginica)
table(iris$target)
```

| | |
|-------------|--------------------------------|
| format_type | <i>Format type description</i> |
|-------------|--------------------------------|

Description

Format type description of variable to 3 letters (intdblglchrdat)

Usage

```
format_type(type)
```

Arguments

| | |
|------|--|
| type | Type description ("integer", "double", "logical", "character", "date") |
|------|--|

Value

Formatted type description (intdblglchrdat)

Examples

```
format_type(typeof(iris$Species))
```

| | |
|----------|--|
| get_nrow | <i>Get number of rows for a grid plot (deprecated, use total_fig_height() instead)</i> |
|----------|--|

Description

Get number of rows for a grid plot (deprecated, use total_fig_height() instead)

Usage

```
get_nrow(varnames, exclude = 0, ncol = 2)
```

Arguments

| | |
|----------|---|
| varnames | List of variables to be plotted |
| exclude | Number of variables that will be excluded from plot |
| ncol | Number of columns (default = 2) |

Value

Number of rows

Examples

```
get_nrow(names(iris), ncol = 2)
```

| | |
|----------|--------------------------------|
| get_type | <i>Return type of variable</i> |
|----------|--------------------------------|

Description

Return value of typeof, except if variable contains <hide>, then return "other"

Usage

```
get_type(var)
```

Arguments

| | |
|-----|-----------------------------|
| var | A vector (dataframe column) |
|-----|-----------------------------|

Value

Value of typeof or "other"

Examples

```
get_type(iris$Species)
```

| | |
|-----------------|---|
| get_var_buckets | <i>Put variables into "buckets" to create a set of plots instead one large plot</i> |
|-----------------|---|

Description

Put variables into "buckets" to create a set of plots instead one large plot

Usage

```
get_var_buckets(data, bucket_size = 100, var_name_target = NA, var_name_n = NA)
```

Arguments

| | |
|-----------------|--|
| data | A dataset |
| bucket_size | Maximum number of variables in one bucket |
| var_name_target | Name of the target variable (if defined) |
| var_name_n | Name of the weight (n) variable (if defined) |

Value

Buckets as a list

Examples

```
get_var_buckets(iris)
get_var_buckets(iris, bucket_size = 2)
get_var_buckets(iris, bucket_size = 2, var_name_target = "Species")
```

| | |
|---------------|--|
| guess_cat_num | <i>Return if variable is categorial or nomerical</i> |
|---------------|--|

Description

Guess if variable is categorial or numerical based on name, type and values of variable

Usage

```
guess_cat_num(var, descr)
```

Arguments

| | |
|-------|--|
| var | A vector (dataframe column) |
| descr | A description of the variable (optional) |

Value

"cat" (categorical), "num" (numerical) or "oth" (other)

Examples

```
guess_cat_num(iris$Species)
```

plot_legend_targetpct *Plots a legend that can be used for explore_all with a binary target*

Description

Plots a legend that can be used for explore_all with a binary target

Usage

```
plot_legend_targetpct(border = TRUE)
```

Arguments

border Draw a border?

Value

Base plot ' @importFrom graphics legend par plot

Examples

```
plot_legend_targetpct(border = TRUE)
```

plot_text *Plot a text*

Description

Plots a text (base plot) and let you choose text-size and color

Usage

```
plot_text(text = "hello world", size = 1.2, color = "black")
```

Arguments

text Text as string
size Text-size
color Text-color

Value

Plot

Examples

```
plot_text("hello", size = 2, color = "red")
```

| | |
|---------------|-----------------------------|
| plot_var_info | <i>Plot a variable info</i> |
|---------------|-----------------------------|

Description

Creates a ggplot with the variable-name as title and a text

Usage

```
plot_var_info(data, var, info = "")
```

Arguments

| | |
|------|--------------|
| data | A dataset |
| var | Variable |
| info | Text to plot |

Value

Plot (ggplot)

| | |
|-----------------|-------------------|
| replace_na_with | <i>Replace NA</i> |
|-----------------|-------------------|

Description

Replace NA values of a variable in a dataframe

Usage

```
replace_na_with(data, var_name, with)
```

Arguments

| | |
|----------|---|
| data | A dataframe |
| var_name | Name of variable where NAs are replaced |
| with | Value instead of NA |

Value

Updated dataframe

Examples

```
data <- data.frame(nr = c(1,2,3,NA,NA))
replace_na_with(data, "nr", 0)
```

| | |
|--------|---|
| report | <i>Generate a report of all variables</i> |
|--------|---|

Description

Generate a report of all variables If target is defined, the relation to the target is reported

Usage

```
report(data, n, target, split = TRUE, output_file, output_dir)
```

Arguments

| | |
|-------------|---|
| data | A dataset |
| n | Weights variable for count data |
| target | Target variable (0/1 or FALSE/TRUE) |
| split | Split by target? (TRUE/FALSE) |
| output_file | Filename of the html report |
| output_dir | Directory where to save the html report |

Examples

```
if (rmarkdown::pandoc_available("1.12.3")) {
  report(iris, output_dir = tempdir())
}
```

| | |
|-----------|--|
| rescale01 | <i>Rescales a numeric variable into values between 0 and 1</i> |
|-----------|--|

Description

Rescales a numeric variable into values between 0 and 1

Usage

```
rescale01(x)
```

Arguments

x numeric vector (to be rescaled)

Value

vector with values between 0 and 1

Examples

```
rescale01(0:10)
```

| | |
|---------------|---------------------------------|
| simplify_text | <i>Simplifies a text string</i> |
|---------------|---------------------------------|

Description

A text string is converted into a simplified version by trimming, converting to upper case, replacing german Umlaute, dropping special characters like comma and semicolon and replacing multiple spaces with one space.

Usage

```
simplify_text(text)
```

Arguments

text text string

Value

text string

Examples

```
simplify_text(" Hello World !, ")
```

target_explore_cat *Explore categorical variable + target*

Description

Create a plot to explore relation between categorical variable and a binary target

Usage

```
target_explore_cat(  
  data,  
  var,  
  target = "target_ind",  
  min_val = NA,  
  max_val = NA,  
  flip = TRUE,  
  num2char = TRUE,  
  title = NA,  
  auto_scale = TRUE,  
  na = NA,  
  max_cat = 30,  
  legend_position = "bottom"  
)
```

Arguments

| | |
|-----------------|--|
| data | A dataset |
| var | Categorical variable |
| target | Target variable (0/1 or FALSE/TRUE) |
| min_val | All values < min_val are converted to min_val |
| max_val | All values > max_val are converted to max_val |
| flip | Should plot be flipped? (change of x and y) |
| num2char | If TRUE, numeric values in variable are converted into character |
| title | Title of plot |
| auto_scale | Not used, just for compatibility |
| na | Value to replace NA |
| max_cat | Maximum numbers of categories to be plotted |
| legend_position | Position of legend ("right" "bottom" "non") |

Value

Plot object

| | |
|--------------------|--|
| target_explore_num | <i>Explore categorical variable + target</i> |
|--------------------|--|

Description

Create a plot to explore relation between numerical variable and a binary target

Usage

```
target_explore_num(  
  data,  
  var,  
  target = "target_ind",  
  min_val = NA,  
  max_val = NA,  
  flip = TRUE,  
  title = NA,  
  auto_scale = TRUE,  
  na = NA,  
  legend_position = "bottom"  
)
```

Arguments

| | |
|-----------------|---|
| data | A dataset |
| var | Numerical variable |
| target | Target variable (0/1 or FALSE/TRUE) |
| min_val | All values < min_val are converted to min_val |
| max_val | All values > max_val are converted to max_val |
| flip | Should plot be flipped? (change of x and y) |
| title | Title of plot |
| auto_scale | Use 0.02 and 0.98 quantile for min_val and max_val (if min_val and max_val are not defined) |
| na | Value to replace NA |
| legend_position | Position of legend ("right" "bottom" "non") |

Value

Plot object

| | |
|------------------|--|
| total_fig_height | <i>Get fig.height for RMarkdown-junk using explore_all()</i> |
|------------------|--|

Description

Get fig.height for RMarkdown-junk using explore_all()

Usage

```
total_fig_height(  
  data,  
  var_name_n,  
  var_name_target,  
  nvar = NA,  
  ncol = 2,  
  size = 3  
)
```

Arguments

| | |
|-----------------|---|
| data | A dataset |
| var_name_n | Weights variable for count data? (TRUE / MISSING) |
| var_name_target | Target variable (TRUE / MISSING) |
| nvar | Number of variables to plot |
| ncol | Number of columns (default = 2) |
| size | fig.height of 1 plot (default = 3) |

Value

Number of rows

Examples

```
total_fig_height(iris)  
total_fig_height(iris, var_name_target = "Species")  
total_fig_height(nvar = 5)
```

| | |
|---------------|-------------------------------|
| weight_target | <i>Weight target variable</i> |
|---------------|-------------------------------|

Description

Create weights for the target variable in your dataset so that are equal weiths for target = 0 and target = 1. Target must be 0/1, FALSE/TRUE ore no/yes

Usage

```
weight_target(data, target)
```

Arguments

| | |
|--------|---|
| data | A dataset |
| target | Target variable (0/1, TRUE/FALSE, yes/no) |

Value

Weights for each observation (as a vector)

Examples

```
iris$is_versicolor <- ifelse(iris$Species == "versicolor", 1, 0)
weights <- weight_target(iris, target = is_versicolor)
summary(weights)
```

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