

Package ‘explore’

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Type Package

Title Simplifies Exploratory Data Analysis

Version 1.4.0

Description Interactive data exploration with one line of code, automated reporting or use an easy to remember set of tidy functions for low code exploratory data analysis.

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

BugReports <https://github.com/rolkra/explore/issues>

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abtest	<i>A/B testing</i>
--------	--------------------

Description

A/B testing

Usage

```
abtest(data, expr, n, target, sign_level = 0.05, color = "grey")
```

Arguments

data	A dataset. If no data is provided, a shiny app is launched
expr	Logical expression, that return in a FALSE/TRUE
n	A Variable for number of observations (count data)
target	Target variable
sign_level	Significance Level (typical 0.01/0.05/0.10)
color	Fill color of bar/violin-plot

Value

Plot that shows if difference is significant

Examples

```
## Using chi2-test or t-test depending on target type
data <- create_data_buy(obs = 100)
abtest(data, female_ind == 1, target = buy) # chi2 test
abtest(data, city_ind == 1, target = age)   # t test

## If small number of observations, Fisher's Exact test
## is used for a binary target (if <= 5 observations in a subgroup)
data <- create_data_buy(obs = 25, seed = 1)
abtest(data, female_ind == 1, target = buy) # Fisher's Exact test
```

abtest_shiny	<i>A/B testing interactive</i>
--------------	--------------------------------

Description

Launches a shiny app to A/B test

Usage

```
abtest_shiny(  
  size_a = 100,  
  size_b = 100,  
  success_a = 10,  
  success_b = 20,  
  success_unit = "percent",  
  sign_level = 0.05  
)
```

Arguments

size_a	Size of Group A
size_b	Size of Group B
success_a	Success of Group A
success_b	Success of Group B
success_unit	"count" "percent"
sign_level	Significance Level (typical 0.01/0.05/0.10)

Examples

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

abtest_targetnum	<i>A/B testing comparing two mean</i>
------------------	---------------------------------------

Description

A/B testing comparing two mean

Usage

```
abtest_targetnum(data, expr, target, sign_level = 0.05, color = "grey")
```

Arguments

data	A dataset
expr	Expression, that results in a FALSE/TRUE
target	Target variable (must be numeric)
sign_level	Significance Level (typical 0.01/0.05/0.10)
color	fill color

Value

Plot that shows if difference is significant

Examples

```
data <- create_data_buy(obs = 100)
abtest(data, city_ind == 1, target = age)
```

abtest_targetpct	<i>A/B testing comparing percent per group</i>
------------------	--

Description

A/B testing comparing percent per group

Usage

```
abtest_targetpct(
  data,
  expr,
  n,
  target,
  sign_level = 0.05,
  group_label,
  ab_label = FALSE,
  color = "grey"
)
```

Arguments

data	A dataset
expr	Expression, that results in a FALSE/TRUE
n	A Variable for number of observations (count data)
target	Target variable (must be 0/1 or FALSE/TRUE)
sign_level	Significance Level (typical 0.01/0.05/0.10)
group_label	Label of groups (default = expr)
ab_label	Label Groups as A and B (default = FALSE)
color	color of bar

Value

Plot that shows if difference is significant

Examples

```
data <- create_data_buy(obs = 100)
abtest(data, female_ind == 1, target = buy)
abtest(data, age >= 40, target = buy)
```

`add_var_id`*Add a variable id at first column in dataset*

Description

Add a variable id at first column in dataset

Usage

```
add_var_id(data, name = "id", overwrite = FALSE)
```

Arguments

<code>data</code>	A dataset
<code>name</code>	Name of new variable (as string)
<code>overwrite</code>	Can new id variable overwrite an existing variable in dataset?

Value

Data set containing new id variable

Examples

```
library(magrittr)
iris %>% add_var_id() %>% head()
iris %>% add_var_id(name = "iris_nr") %>% head()
```

add_var_random_01	<i>Add a random 0/1 variable to dataset</i>
-------------------	---

Description

Add a random 0/1 variable to dataset

Usage

```
add_var_random_01(  
  data,  
  name = "random_01",  
  prob = c(0.5, 0.5),  
  overwrite = TRUE,  
  seed  
)
```

Arguments

data	A dataset
name	Name of new variable (as string)
prob	Vector of probabilities
overwrite	Can new random variable overwrite an existing variable in dataset?
seed	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)  
iris %>% add_var_random_01() %>% head()  
iris %>% add_var_random_01(name = "my_var") %>% head()
```

add_var_random_cat	<i>Add a random categorical variable to dataset</i>
--------------------	---

Description

Add a random categorical variable to dataset

Usage

```
add_var_random_cat(
  data,
  name = "random_cat",
  cat = LETTERS[1:6],
  prob,
  overwrite = TRUE,
  seed
)
```

Arguments

data	A dataset
name	Name of new variable (as string)
cat	Vector of categories
prob	Vector of probabilities
overwrite	Can new random variable overwrite an existing variable in dataset?
seed	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)
iris %>% add_var_random_cat() %>% head()
iris %>% add_var_random_cat(name = "my_cat") %>% head()
iris %>% add_var_random_cat(cat = c("Version A", "Version B")) %>% head()
iris %>% add_var_random_cat(cat = c(1,2,3,4,5)) %>% head()
```

add_var_random_dbl	<i>Add a random double variable to dataset</i>
--------------------	--

Description

Add a random double variable to dataset

Usage

```
add_var_random_dbl(
  data,
  name = "random_dbl",
  min_val = 0,
  max_val = 100,
  overwrite = TRUE,
  seed
)
```

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?
seed	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)
iris %>% add_var_random_dbl() %>% head()
iris %>% add_var_random_dbl(name = "random_var") %>% head()
iris %>% add_var_random_dbl(min_val = 1, max_val = 10) %>% head()
```

add_var_random_int	<i>Add a random integer variable to dataset</i>
--------------------	---

Description

Add a random integer variable to dataset

Usage

```
add_var_random_int(
  data,
  name = "random_int",
  min_val = 1,
  max_val = 10,
  overwrite = TRUE,
  seed
)
```

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?
seed	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)
iris %>% add_var_random_int() %>% head()
iris %>% add_var_random_int(name = "random_var") %>% head()
iris %>% add_var_random_int(min_val = 1, max_val = 10) %>% head()
```

add_var_random_moon	<i>Add a random moon variable to dataset</i>
---------------------	--

Description

Add a random moon variable to dataset

Usage

```
add_var_random_moon(data, name = "random_moon", overwrite = TRUE, seed)
```

Arguments

data	A dataset
name	Name of new variable (as string)
overwrite	Can new random variable overwrite an existing variable in dataset?
seed	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)
iris %>% add_var_random_moon() %>% head()
```

`add_var_random_starsign`*Add a random starsign variable to dataset*

Description

Add a random starsign variable to dataset

Usage

```
add_var_random_starsign(  
  data,  
  name = "random_starsign",  
  lang = "en",  
  overwrite = TRUE,  
  seed  
)
```

Arguments

<code>data</code>	A dataset
<code>name</code>	Name of new variable (as string)
<code>lang</code>	Language used for starsign (en = English, de = Deutsch, es = Espanol)
<code>overwrite</code>	Can new random variable overwrite an existing variable in dataset?
<code>seed</code>	Seed for random number generation (integer)

Value

Dataset containing new random variable

Examples

```
library(magrittr)  
iris %>% add_var_random_starsign() %>% head()  
iris %>% add_var_random_starsign(lang = "de") %>% head()
```

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

Description

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

Usage

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

Arguments

data	A dataset
target	Target variable (0/1, TRUE/FALSE, yes/no)
min_prop	Minimum proportion of one of the target categories
seed	Seed for random number generator

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)
```

check_vec_low_variance	<i>Check vector for low variance</i>
------------------------	--------------------------------------

Description

Check vector for low variance

Usage

```
check_vec_low_variance(values, max_prop = 0.99)
```

Arguments

values	Vector of values
max_prop	Maximum proportion of values without variance

Value

TRUE/FALSE (low variance)

Examples

```
## Not run:
values <- c(1, rep(0 ,1000))
check_vec_low_variance(values, max_prop = 0.9)

## End(Not run)
```

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	IF TRUE, value is rescaled 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

```
library(magrittr)
iris %>% clean_var(Sepal.Width, max_val = 3.5, name = "sepal_width") %>% head()
iris %>% clean_var(Sepal.Width, rescale01 = TRUE) %>% head()
```

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_data_abtest	<i>Create data of A/B testing</i>
--------------------	-----------------------------------

Description

Data that can be used for unit-testing or teaching

Usage

```
create_data_abtest(
  n_a = 100,
  n_b = 100,
  success_a = 10,
  success_b = 5,
  success_unit = "count",
  count = TRUE
)
```

Arguments

n_a	Total size of group A
n_b	Total size of group B
success_a	Success in group A
success_b	Success in group B
success_unit	Unit ("count" "percent")
count	Create as count-data (FALSE/TRUE)

Value

A dataset as tibble

Examples

```
library(dplyr)
create_data_abtest() %>% abtest()
create_data_abtest(
  n_a = 100,
  n_b = 100,
  success_a = 20,
  success_b = 30,
  success_unit = "count"
) %>% abtest()
```

create_data_app

Create data app

Description

Artificial data that can be used for unit-testing or teaching

Usage

```
create_data_app(obs = 1000, add_id = FALSE, seed = 123)
```

Arguments

obs	Number of observations
add_id	Add an id-variable to data?
seed	Seed for randomization (integer)

Value

A dataset as tibble

Examples

```
create_data_app()
```

create_data_buy	<i>Create data buy</i>
-----------------	------------------------

Description

Artificial data that can be used for unit-testing or teaching

Usage

```
create_data_buy(  
  obs = 1000,  
  target_name = "buy",  
  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 factor no/yes)?
target1_prob	Probability that target = 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_prd = Customer has a mobile data product (NO/MOBILE STICK/BUSINESS)
- bbi_speed_ind = Customer has a Broadband Internet (BBI) with extra 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:

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

Value

A dataset as tibble

Examples

```
create_data_buy()
```

create_data_churn	<i>Create data churn</i>
-------------------	--------------------------

Description

Artificial data that can be used for unit-testing or teaching

Usage

```
create_data_churn(
  obs = 1000,
  target_name = "churn",
  factorise_target = FALSE,
  target1_prob = 0.4,
  add_id = FALSE,
  seed = 123
)
```

Arguments

obs	Number of observations
target_name	Variable name of target
factorise_target	Should target variable be factorised?
target1_prob	Probability that target = 1
add_id	Add an id-variable to data?
seed	Seed for randomization (integer)

Value

A dataset as tibble

Examples

```
create_data_churn()
```

create_data_empty	<i>Create an empty dataset</i>
-------------------	--------------------------------

Description

Create an empty dataset

Usage

```
create_data_empty(obs = 1000, add_id = FALSE)
```

Arguments

obs	Number of observations
add_id	Add an id

Value

Dataset as tibble

Examples

```
create_data_empty(obs = 100)
create_data_empty(obs = 100, add_id = TRUE)
```

`create_data_esoteric` *Create data esoteric*

Description

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

Usage

```
create_data_esoteric(obs = 1000, add_id = FALSE, seed = 123)
```

Arguments

<code>obs</code>	Number of observations
<code>add_id</code>	Add an id-variable to data?
<code>seed</code>	Seed for randomization

Details

Variables in dataset:

- `id` = Identifier
- `starsign` = random starsign
- `chinese` = random chinese zodiac
- `moon` = random moon phase
- `blood` = random blood type
- `fingers_crossed` = random fingers crossed (1 = yes, 0 = no)
- `success` = random success (1 = yes, 0 = no)

Value

A dataset as tibble

Examples

```
create_data_esoteric(obs = 100)
```

`create_data_newsletter`*Create data newsletter*

Description

Artificial data that can be used for unit-testing or teaching (fairness & AI bias)

Usage

```
create_data_newsletter(obs = 1000, add_id = FALSE, seed = 123)
```

Arguments

<code>obs</code>	Number of observations
<code>add_id</code>	Add an id-variable to data?
<code>seed</code>	Seed for randomization (integer)

Value

A dataset as tibble

Examples

```
create_data_newsletter()
```

`create_data_person`*Create data person*

Description

Artificial data that can be used for unit-testing or teaching

Usage

```
create_data_person(obs = 1000, add_id = FALSE, seed = 123)
```

Arguments

<code>obs</code>	Number of observations
<code>add_id</code>	Add an id
<code>seed</code>	Seed for randomization (integer)

Value

A dataset as tibble

Examples

```
create_data_person()
```

```
create_data_random
```

Create data random

Description

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

Usage

```
create_data_random(
  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 target = 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 dataset as tibble

Examples

```
create_data_random(obs = 100, vars = 5)
```

create_data_unfair	Create data unfair
--------------------	--------------------

Description

Artificial data that can be used for unit-testing or teaching (fairness & AI bias)

Usage

```
create_data_unfair(  
  obs = 1000,  
  target_name = "target_ind",  
  factorise_target = FALSE,  
  target1_prob = 0.25,  
  add_id = FALSE,  
  seed = 123  
)
```

Arguments

obs	Number of observations
target_name	Variable name of target
factorise_target	Should target variable be factorised?
target1_prob	Probability that target = 1
add_id	Add an id-variable to data?
seed	Seed for randomization (integer)

Value

A dataset as tibble

Examples

```
create_data_unfair()
```

`create_notebook_explore`*Generate a notebook*

Description

Generate an RMarkdown Notebook template for a report. You must provide a output-directory (parameter `output_dir`). The default file-name is "notebook-explore.Rmd" (may overwrite existing file with same name)

Usage

```
create_notebook_explore(output_file = "notebook-explore.Rmd", output_dir)
```

Arguments

<code>output_file</code>	Filename of the html report
<code>output_dir</code>	Directory where to save the html report

Examples

```
create_notebook_explore(output_file = "explore.Rmd", output_dir = tempdir())
```

`cut_vec_num_avg`*Cut a variable*

Description

Cut a variable

Usage

```
cut_vec_num_avg(values, bins = 8)
```

Arguments

<code>values</code>	Variable
<code>bins</code>	Number of bins

Value

Data frame

data_dict_md	Create a data dictionary Markdown file
--------------	--

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	<i>decrypt text</i>
---------	---------------------

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, 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" "tibble" "df")
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[,1] <- NA  
describe_tbl(iris)
```

drop_obs_if	<i>Drop all observations where expression is true</i>
-------------	---

Description

Drop all observations where expression is true

Usage

```
drop_obs_if(data, expr)
```

Arguments

data	Data frame
expr	Expression

Value

Data frame

Examples

```
drop_obs_if(iris, Species == "setosa")
drop_obs_if(iris, Sepal.Length < 5 | Sepal.Length > 7)
```

drop_obs_with_na	<i>Drop all observations with NA-values</i>
------------------	---

Description

Drop all observations with NA-values

Usage

```
drop_obs_with_na(data)
```

Arguments

data	Data frame
------	------------

Value

Data frame

Examples

```
data <- data.frame(a = 1:10, b = rep("A",10))
data[1,1] <- NA
drop_obs_with_na(data)
```

drop_var_by_names	<i>Drop variables by name</i>
-------------------	-------------------------------

Description

Drop variables by name

Usage

```
drop_var_by_names(data, var_names)
```

Arguments

data	Data frame
var_names	Vector of variable names (as string)

Value

Data frame

Examples

```
drop_var_by_names(iris, "Species")  
drop_var_by_names(iris, c("Sepal.Length", "Sepal.Width"))
```

drop_var_low_variance	<i>Drop all variables with low variance</i>
-----------------------	---

Description

Drop all variables with low variance

Usage

```
drop_var_low_variance(data, max_prop = 0.99)
```

Arguments

data	Data frame
max_prop	Maximum proportion of values without variance

Value

Data frame

Examples

```
data <- data.frame(a = 1:100, b = c(0, rep(1, 99)))
drop_var_low_variance(data, max_prop = 0.9)
```

`drop_var_not_numeric` *Drop all not numeric variables*

Description

Drop all not numeric variables

Usage

```
drop_var_not_numeric(data)
```

Arguments

`data` Data frame

Value

Data frame

Examples

```
data <- data.frame(a = 1:10, b = rep("A",10))
drop_var_not_numeric(data)
```

`drop_var_no_variance` *Drop all variables with no variance*

Description

Drop all variables with no variance

Usage

```
drop_var_no_variance(data)
```

Arguments

`data` Data frame

Value

Data frame

Examples

```
data <- data.frame(a = 1:10, b = rep(1,10))
drop_var_no_variance(data)
```

drop_var_with_na	<i>Drop all variables with NA-values</i>
------------------	--

Description

Drop all variables with NA-values

Usage

```
drop_var_with_na(data)
```

Arguments

data	Data frame
------	------------

Value

Data frame

Examples

```
data <- data.frame(a = 1:10, b = rep(NA,10))
drop_var_with_na(data)
```

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_forest	<i>Explain a target using Random Forest.</i>
----------------	--

Description

Explain a target using Random Forest.

Usage

```
explain_forest(data, target, ntree = 50, out = "plot", ...)
```

Arguments

data	A dataset
target	Target variable (binary)
ntree	Number of trees used for Random Forest
out	Output of the function: "plot" "model" "importance" all"
...	Further arguments

Value

Plot of importance (if out = "plot")

Examples

```
data <- create_data_buy()
explain_forest(data, target = buy)
```

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)

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 = 20,
  cp = 0,
  weights = NA,
  size = 0.7,
  out = "plot",
  ...
)

```

Arguments

<code>data</code>	A dataset
<code>target</code>	Target variable
<code>n</code>	weights variable (for count data)
<code>max_cat</code>	Drop categorical variables with higher number of levels
<code>max_target_cat</code>	Maximum number of categories to be plotted for target (except NA)
<code>maxdepth</code>	Set the maximum depth of any node of the final tree, with the root node counted as depth 0. Maximum value 30 (as node numbers have to be stored in 32-bit signed integer types).
<code>minsplit</code>	the minimum number of observations that must exist in a node in order for a split to be attempted.
<code>cp</code>	complexity parameter. Any split that does not decrease the overall lack of fit by a factor of <code>cp</code> is not attempted. For instance, with anova splitting, this means that the overall R-squared must increase by <code>cp</code> at each step. The main role of this parameter is to save computing time by pruning off splits that are obviously not worthwhile. Essentially, the user informs the program that any split which does not improve the fit by <code>cp</code> will likely be pruned off by cross-validation, and that hence the program need not pursue it.
<code>weights</code>	optional case weights.
<code>size</code>	Text size of plot
<code>out</code>	Output of function: "plot" "model"
<code>...</code>	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)
```

explain_xgboost

Explain a binary target using xgboost

Description

Based on the hyperparameters defined in the setup parameter, XGBoost hyperparameter-tuning is carried out using cross-validation. The best model is chosen and returned. As default, the function returns the feature-importance plot. To get the all outputs, use parameter out = "all"

Usage

```
explain_xgboost(
  data,
  target,
  log = TRUE,
  nthread = 1,
  setup = list(cv_nfold = 2, max_nrounds = 1000, early_stopping_rounds = 50, grid_xgboost =
    list(eta = c(0.3, 0.1, 0.01), max_depth = c(3, 5), gamma = 0, colsample_bytree =
      0.8, subsample = 0.8, min_child_weight = 1, scale_pos_weight = 1)),
  out = "plot"
)
```

Arguments

data	Data frame, must contain variable defined in target, but should not contain any customer-IDs or date/period columns
target	Target variable (must be binary 0/1, FALSE/TRUE, no/yes)
log	Log?
nthread	Number of threads used for training
setup	Setup of model
out	Output of the function: "plot" "model" "importance" all"

Value

Plot of importance (if out = "plot")

Examples

```
data <- use_data_iris()
data$is_versicolor <- ifelse(data$Species == "versicolor", 1, 0)
data$Species <- NULL
explain_xgboost(data, target = is_versicolor, log = FALSE)
```

explore

*Explore a dataset or variable***Description**

Explore a dataset or variable

Usage

```
explore(
  data,
  var,
  var2,
  n,
  target,
  targetpct,
  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)
targetpct	Plot variable as target% (FALSE/TRUE)
split	Alternative to targetpct (split = !targetpct)
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,
  targetpct,
  color = c("#ADD8E6", "#7BB8DA"),
  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)
targetpct	Plot variable as target% (FALSE/TRUE)
color	Force a default color (if possible)
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

Explore categorical variable using bar charts

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,
  color = c("#ADD8E6", "#7BB8DA"),
  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)
color	Color for bar
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_col	<i>Explore data without aggregation (label + value)</i>
-------------	---

Description

Label and Value are in the data. Create a bar plot where the heights of the bars represent the values for each label.

Usage

```
explore_col(  
  data,  
  var_label,  
  var_value,  
  title = NA,  
  subtitle = "",  
  numeric = FALSE,  
  max_cat = 30,  
  na = 0,  
  flip = NA,  
  color = "#ADD8E6"  
)
```

Arguments

<code>data</code>	A dataset (categories + frequency)
<code>var_label</code>	Variable containing the label
<code>var_value</code>	Variable containing the value
<code>title</code>	Title of the plot
<code>subtitle</code>	Subtitle of the plot
<code>numeric</code>	Display variable as numeric (not category)
<code>max_cat</code>	Maximum number of categories to be plotted
<code>na</code>	Value to use for NA
<code>flip</code>	Flip plot? (for categorical variables)
<code>color</code>	Color for bar

Value

Plot object

Examples

```
library(magrittr)
data <- data.frame(label = LETTERS[1:5], value = c(1.5,2,1.2,3,2.6))
data %>% explore_col(label, value)
```

<code>explore_cor</code>	<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 = c("#ADD8E6", "#7BB8DA"),
  ...
)
```

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,  
  color = c("#ADD8E6", "#7BB8DA"),  
  flip = NA  
)
```

Arguments

<code>data</code>	A dataset (categories + frequency)
<code>cat</code>	Numerical variable
<code>n</code>	Number of observations (frequency)
<code>target</code>	Target variable
<code>pct</code>	Show as percent?
<code>split</code>	Split by target (FALSE/TRUE)
<code>title</code>	Title of the plot
<code>numeric</code>	Display variable as numeric (not category)
<code>max_cat</code>	Maximum number of categories to be plotted
<code>max_target_cat</code>	Maximum number of categories to be plotted for target (except NA)
<code>color</code>	Color for bar
<code>flip</code>	Flip plot? (for categorical variables)

Value

Plot object

Examples

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

<code>explore_density</code>	<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 = c("#ADD8E6", "#7BB8DA"),
  auto_scale = TRUE,
  max_target_cat = 5,
  ...
)
```

Arguments

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

Description

Launches a shiny app to explore a dataset

Usage

```
explore_shiny(data, target, color = c("#ADD8E6", "#7BB8DA"))
```

Arguments

<code>data</code>	A dataset
<code>target</code>	Target variable (0/1 or FALSE/TRUE)
<code>color</code>	Color for plots (vector)

Examples

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

explore_targetpct	<i>Explore variable + binary target (values 0/1)</i>
-------------------	--

Description

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

Usage

```
explore_targetpct(
  data,
  var,
  target = NULL,
  title = NA,
  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 (intdblglchrldat)

Usage

```
format_type(type)
```

Arguments

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

Value

Formatted type description (intdblglchrldat)

Examples

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

get_color	<i>Get predefined colors</i>
-----------	------------------------------

Description

Get predefined colors

Usage

```
get_color(name, fill = FALSE, fill_color = "#DDDDDD", fill_n = 10)
```

Arguments

name	Name of color/color-vector
fill	Fill color vector?
fill_color	Color to use to fill color vector
fill_n	Number of color codes to return

Value

Vector of color-codes

Examples

```
get_color("mario")

get_color("mario")
show_color(get_color("mario"))
show_color(get_color("mario", fill = TRUE, fill_n = 10))

col <- get_color("mario")
explore(iris, Sepal.Length, target = Species,
  color = col)
explore(iris, Sepal.Length, target = Species,
  color = c(col["peach"], col["bowser"], col["donkeykong"]))
```

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 categorical or numerical</i>
---------------	---

Description

Guess if variable is categorical 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)
```

interact	<i>Make a explore-plot interactive</i>
----------	--

Description

Make a explore-plot interactive

Usage

```
interact(obj, lower_title = TRUE, hide_geom_text = TRUE)
```

Arguments

obj	A object (e.g. ggplot2-object)
lower_title	Lowering the title in ggplot2-object(FALSE/TRUE)
hide_geom_text	Hiding geom_text in ggplot2-object (FALSE/TRUE)

Value

Plot object

Examples

```
library(dplyr)
if (interactive()) {
  iris %>% explore(Sepal.Length, target = Species) %>% interact()
}
```

log_info_if	<i>Log conditional</i>
-------------	------------------------

Description

Log conditional

Usage

```
log_info_if(log = TRUE, text = "log")
```

Arguments

log	log (TRUE FALSE)
text	text string to be logged

Value

prints log on screen (if log == TRUE).

mix_color	<i>Mix colors</i>
-----------	-------------------

Description

Mix colors

Usage

```
mix_color(color1, color2 = NA, n = 5)
```

Arguments

color1	Color 1
color2	Color 2
n	Number of different colors that should be generated

Value

Vector of color-codes

Examples

```
mix_color("blue", n = 10)
mix_color("gold", "red", n = 4)
```

plot_legend_targetpct	<i>Plots a legend that can be used for explore_all with a binary target</i>
-----------------------	---

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

Examples

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

plot_text	<i>Plot a text</i>
-----------	--------------------

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", ggplot = FALSE)
```

Arguments

text	Text as string
size	Text-size
color	Text-color
ggplot	return a ggplot-object? (or base plot)

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)

predict_target	<i>Predict target using a trained model.</i>
----------------	--

Description

Predict target using a trained model.

Usage

```
predict_target(data, model, name = "prediction")
```

Arguments

data	A dataset (data.frame or tbl)
model	A model created with explain_*() function
name	Prefix of variable-name for prediction

Value

data containing predicted probabilities for target values

Examples

```
data_train <- create_data_buy(seed = 1)
data_test <- create_data_buy(seed = 2)
model <- explain_tree(data_train, target = buy, out = "model")
data <- predict_target(data = data_test, model = model)
describe(data)
```

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, targetpct, split, color, output_file, output_dir)
```

Arguments

data	A dataset
n	Weights variable for count data
target	Target variable (0/1 or FALSE/TRUE)
targetpct	Plot variable as target% (FALSE/TRUE)
split	Alternative to targetpct (split = !targetpct)
color	User defined colors for plots (vector)
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)
```

show_color	Show color vector as ggplot
------------	-----------------------------

Description

Show color vector as ggplot

Usage

```
show_color(color)
```

Arguments

color Vector of colors

Value

ggplot

Examples

```
show_color("gold")
show_color(c("blue", "red", "green"))
```

simplify_text	Simplifies a text string
---------------	--------------------------

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

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 = 25,
  color = c("#ECEFF1", "#CFD8DC", "#B0BEC5", "#90A4AE"),
  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
color	Color vector (4 colors)
legend_position	Position of legend ("right" "bottom" "non")
Value	
	Plot object
target_explore_num	<i>Explore Nuberical 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,  
  bins = 10,  
  flip = TRUE,  
  title = NA,  
  auto_scale = TRUE,  
  na = NA,  
  color = c("#ECEFF1", "#CFD8DC", "#B0BEC5", "#90A4AE"),  
  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
bins	Nuber of bins
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
color	Color vector (4 colors)
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)
```

use_data_beer	<i>Use the beer data set</i>
---------------	------------------------------

Description

This data set is an incomplete collection of popular beers in Austria, Germany and Switzerland. Data are collected from various websites in 2023. Some of the collected data may be incorrect.

Usage

```
use_data_beer()
```

Value

Dataset as tibble

Examples

```
use_data_beer()
```

use_data_diamonds	<i>Use the diamonds data set</i>
-------------------	----------------------------------

Description

This data set comes with the ggplot2 package. It contains the prices and other attributes of almost 54,000 diamonds.

Usage

```
use_data_diamonds()
```

Value

Dataset

See Also

[ggplot2::diamonds](#)

Examples

```
use_data_diamonds()
```

use_data_iris	<i>Use the iris flower data set</i>
---------------	-------------------------------------

Description

This data set comes with base R. The data set gives the measurements in centimeters of the variables sepal length and width and petal length and width, respectively, for 50 flowers from each of 3 species of iris. The species are *Iris setosa*, *versicolor*, and *virginica*.

Usage

```
use_data_iris()
```

Value

Dataset as tibble

Examples

```
use_data_iris()
```

use_data_mpg	<i>Use the mpg data set</i>
--------------	-----------------------------

Description

This data set comes with the ggplot2 package. It contains a subset of the fuel economy data that the EPA makes available on <https://fuelconomy.gov/>. It contains only models which had a new release every year between 1999 and 2008 - this was used as a proxy for the popularity of the car.

Usage

```
use_data_mpg()
```

Value

Dataset

See Also

[ggplot2::mpg](#)

Examples

```
use_data_mpg()
```

use_data_mtcars	<i>Use the mtcars data set</i>
-----------------	--------------------------------

Description

This data set comes with base R. The data was extracted from the 1974 Motor Trend US magazine, and comprises fuel consumption and 10 aspects of automobile design and performance for 32 automobiles (1973–74 models).

Usage

```
use_data_mtcars()
```

Value

Dataset

Examples

```
use_data_mtcars()
```

use_data_penguins	<i>Use the penguins data set</i>
-------------------	----------------------------------

Description

This data set comes with the palmerpenguins package. It contains measurements for penguin species, island in Palmer Archipelago, size (flipper length, body mass, bill dimensions), and sex.

Usage

```
use_data_penguins(short_names = FALSE)
```

Arguments

short_names	Use short variable names
-------------	--------------------------

Value

Dataset

See Also

[palmerpenguins::penguins](#)

Examples

```
use_data_penguins()  
use_data_penguins(short_names = TRUE)
```

use_data_starwars	<i>Use the starwars data set</i>
-------------------	----------------------------------

Description

This data set comes with the dplyr package. It contains data of 87 star war characters

Usage

```
use_data_starwars()
```

Value

Dataset

See Also

[dplyr::starwars](#)

Examples

```
use_data_starwars()
```

use_data_titanic	<i>Use the titanic data set</i>
------------------	---------------------------------

Description

This data set comes with base R. Survival of passengers on the Titanic.

Usage

```
use_data_titanic(count = FALSE)
```

Arguments

count	use count data
-------	----------------

Value

Dataset

Examples

```
use_data_titanic(count = TRUE)  
use_data_titanic(count = FALSE)
```

use_data_wordle	<i>Use the wordle data set</i>
-----------------	--------------------------------

Description

This data set contains the result of a real wordle challange (in german language) between tow players. Wordle is a game where a player guesses a five-letter word in six tries. The variable "try" reflects the success of player A and B. Other variables like "noun", "aeiou", "unique", "common" and "rare" reflect the properties of the word.

Usage

```
use_data_wordle()
```

Value

Dataset

Examples

```
use_data_wordle()
```

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

Description

Create weights for the target variable in your dataset so that are equal weights 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)
versicolor <- iris$is_versicolor
table(versicolor, weights)
```

`yyyyymm_calc`*Calculate with periods (format yyyymm)*

Description

Calculate with periods (format yyyymm)

Usage

```
yyyyymm_calc(yyymm, add_month = 0, add_year = 0, diff_to = NA)
```

Arguments

<code>yyymm</code>	Input vector of periods (format yyyymm)
<code>add_month</code>	How many months to add (can be negative too)
<code>add_year</code>	How many years to add (can be negative too)
<code>diff_to</code>	Difference between date and yyyymm (format yyyymm)

Value

Vector of periods (format yyyymm) or number of months

Examples

```
yyyyymm_calc(202412, add_month = 1)
yyyyymm_calc(c(202411,202412,202501), add_month = -1, add_year = 1)
yyyyymm_calc(202410, diff_to = 202501)
yyyyymm_calc(c(202411,202412,202501,202502), diff_to = 202501)
```

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