

Package ‘**lares**’

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Type Package

Title Lean Analytics and Robust Exploration Sidekick

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Maintainer Bernardo Lares <laresbernardo@gmail.com>

Description Auxiliary package for better/faster analytics, visualization, data mining, and machine learning tasks. With a wide variety of family functions, like Machine Learning, Data Wrangling, Marketing Mix Modeling (Robyn), Exploratory, API, and Scrapper, it helps the analyst or data scientist to get quick and robust results, without the need of repetitive coding or advanced R programming skills.

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Author Bernardo Lares [aut, cre]

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autoline*New Line Feed for Long Strings (Wrapper)*

Description

Add a break or new line without breaking words. Automatically, the function can detect your plot's width and will dynamically set an auto width. You can adjust the relation (rel) parameter for different fonts and sizes until perfect harmony found. Quite similar to `stringr::str_wrap` but, if the text vector is a factor, the levels will be kept in order and transformed.

Usage

```
autoline(text, top = "auto", rel = 9)
```

Arguments

<code>text</code>	Character or factor vector.
<code>top</code>	Integer. How many characters aprox. should be on each line?
<code>rel</code>	Numeric. Relation of pixels and characters per line

Value

Character. String (vector) including some \n within.

See Also

Other Tools: `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
cat(autoline("This is a long text that may not fit into a single line", 8))

text <- factor(c("First value", "Second value", "First value"),
               levels = c("First value", "Second value"))
)
autoline(text, 1)

path <- file.path(R.home("doc"), "THANKS")
text <- paste(readLines(path), collapse = " ")
cat(autoline(text))
```

balance_data*Balance Binary Data by Resampling: Under-Over Sampling*

Description

This function lets the user balance a given data.frame by resampling with a given relation rate and a binary feature.

Usage

```
balance_data(df, var, rate = 1, target = "auto", seed = 0, quiet = FALSE)
```

Arguments

df	Vector or Dataframe. Contains different variables in each column, separated by a specific character
var	Variable. Which variable should we used to re-sample dataset?
rate	Numeric. How many X for every Y we need? Default: 1. If there are more than 2 unique values, rate will represent percentage for number of rows
target	Character. If binary, which value should be reduced? If kept in "auto", then the most frequent value will be reduced.
seed	Numeric. Seed to replicate and obtain same values
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

data.frame. Reduced sampled data.frame following the rate of appearance of a specific variable.

See Also

Other Data Wrangling: [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
data(dft) # Titanic dataset
df <- balance_data(dft, Survived, rate = 1)
df <- balance_data(dft, .data$Survived, rate = 0.5, target = "TRUE")
```

<code>bind_files</code>	<i>Bind Files into Dataframe</i>
-------------------------	----------------------------------

Description

This function imports and binds multiple files into a single data.frame. Files must be inserted with absolute roots files names.

Usage

```
bind_files(files)
```

Arguments

<code>files</code>	Character vector. Files names.
--------------------	--------------------------------

Value

data.frame with data joined from all files passed.

See Also

Other Tools: [autoline\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

<code>bring_api</code>	<i>Get API (JSON) and Transform into data.frame</i>
------------------------	---

Description

This function lets the user bring API data as JSON format and transform it into data.frame.

Usage

```
bring_api(url, status = TRUE)
```

Arguments

<code>url</code>	Character. API's URL to GET.
<code>status</code>	Boolean. Display status message?

Value

data.frame of url GET results or NULL if no results returned by API.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)
Other API: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

cache_write

Cache Save and Load (Write and Read)

Description

This function lets the user save and load a cache of any R object to improve timings and UX.

Usage

```
cache_write(  
  data,  
  base = "temp",  
  cache_dir = getOption("LARES_CACHE_DIR"),  
  ask = FALSE,  
  overwrite = TRUE,  
  quiet = FALSE,  
  ...  
)  
  
cache_read(  
  base,  
  cache_dir = getOption("LARES_CACHE_DIR"),  
  ask = FALSE,  
  overwrite = TRUE,  
  quiet = FALSE,  
  ...  
)  
  
cache_exists(base = NULL, cache_dir = getOption("LARES_CACHE_DIR"), ...)  
cache_clear(cache_dir = getOption("LARES_CACHE_DIR"), quiet = FALSE, ...)  
cache_pipe(data, base = "cache_pipe", read = TRUE, write = TRUE, ...)
```

Arguments

data	Object
base	Character vector. Unique name for your cache file. You can pass a character vector with multiple elements that will be concatenated. All cache files with start with lares_cache_* automatically to quickly detect these cache files.
cache_dir	Character. Where do you want to save you cache files? By default they'll be stored on tempdir() but you can change it using this parameter or setting a global option called "LARES_CACHE_DIR".
ask	Boolean. If cache exists, when reading: (interactive) ask the user if the cache should be used to proceed or ignored; when writing, (interactive) ask the user if the cache should be overwritten. Note that you can only ask for one cache file at a time because vectors are concatenated.
overwrite	Boolean. Set to overwrite existing cache file. When reading, this parameter answers to ask prompt instead.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.
read, write	Boolean. Do you wish to read and or write cache?

Value

cache_write. No return value, called for side effects.
 cache_read. R object. Data from cache file or NULL if no cache found.
 cache_exists. Boolean. Result of base existence.
 cache_clear. Invisible vector containing cache file names removed.
 cache_pipe. Same as x or cached result.

Examples

```

x <- list(a = 1, b = 2:4)
base <- c(as.character(Sys.Date()), "A", "B")
cache_write(x, base)
cache_read(base, ask = FALSE)
cache_exists(base)
cache_clear()
  
```

Description

This function lets the user reduce categorical values in a vector. It is tidyverse friendly for use on pipelines

Usage

```
categ_reducer(
  df,
  var,
  nmin = 0,
  pmin = 0,
  pcummax = 100,
  top = NA,
  pvalue_max = 1,
  cor_var = "tag",
  limit = 20,
  other_label = "other",
  ...
)
```

Arguments

df	Categorical Vector
var	Variable. Which variable do you wish to reduce?
nmin	Integer. Number of minimum times a value is repeated
pmin	Numerical. Percentage of minimum times a value is repeated
pcummax	Numerical. Top cumulative percentage of most repeated values
top	Integer. Keep the n most frequently repeated values
pvalue_max	Numeric (0-1]. Max pvalue categories
cor_var	Character. If pvalue_max < 1, you must define which column name will be compared with (numerical or binary).
limit	Integer. Limit one hot encoding to the n most frequent values of each column. Set to NA to ignore argument.
other_label	Character. With which text do you wish to replace the filtered values with?
...	Additional parameters.

Value

data.frame df on which var has been transformed

See Also

Other Data Wrangling: [balance_data\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zeroivar\(\)](#)

Examples

```
data(dft) # Titanic dataset
categ_reducer(dft, Embarked, top = 2) %>% freqs(Embarked)
categ_reducer(dft, Ticket, nmin = 7, other_label = "Other Ticket") %>% freqs(Ticket)
categ_reducer(dft, Ticket, pvalue_max = 0.05, cor_var = "Survived") %>% freqs(Ticket)
```

check_opts

Validate inputs (attributions, options, ...)

Description

This function validates if inputs match all/any of your options and return error/message with possible options to use. Similar to `match.arg()` but more flexible.

This function checks if an object has a specific attribute and stops if not.

Check whether a value or vector is or is not following a set of rules. For example: is an URL, is an ID vector, are non-variant or constant values, are binary values... Notice that `is_` will return the result for each observation and `are_` for the whole vector.

Usage

```
check_opts(
  inputs,
  opts,
  input_name = "input",
  type = "all",
  not = "stop",
  quiet = TRUE
)

check_attr(object, attr = "type", check = NULL, stop = TRUE)

is_url(x, ...)

is_ip(x, ...)

are_id(x)

are_constant(x)

are_binary(x)

is_even(x)

is_odd(x)
```

Arguments

inputs	Vector character. Check options.
opts	Vector character. Valid options.
input_name	Character. Custom your message and change "input" for any other string. For example: "column names".
type	Character. Options: "all", "any."
not	Character. Options: "stop", "message", "print", "return".
quiet	Boolean. Keep quiet? If not, returns logical value.
object	Object of any kind
attr	Character. Attribute to check
check	Character. Attribute value
stop	Boolean. Stop if doesn't check?
x	Vector
...	Additional parameters passed to grep1()

Value

Boolean. Result of inputs in opts (options). Depending on type and/or stop arguments, errors or messages will be shown.

No return value, called for side effects.

is_url. Boolean. Result of checking if x is a valid URL string.

is_ip. Boolean. Result of checking if x is a valid IP string.

are_id. Boolean. Result of checking if x is a potential ID vector

are_constant. Boolean. Result of checking if x is a constant vector

are_binary. Boolean. Result of checking if x is a binary vector

Examples

```
opts <- c("A", "B", "C")
# Let's check the "all" logic
check_opts(inputs = c("A", "B"), opts, quiet = FALSE)
check_opts(inputs = c("X"), opts, not = "message", quiet = FALSE)
check_opts(inputs = c("A", "X"), opts, input_name = "value", not = "warning")
# Now let's check the "any" logic
check_opts(inputs = c("A", "X"), opts, type = "any")
check_opts(inputs = c("X"), opts, type = "any", not = "message")
check_opts(inputs = c("A", NA), opts, type = "any")
# Final trick: just ignore results
check_opts(inputs = "X", opts, not = "invisible")
test <- data.frame()
attributes(test)
check_attr(test, "class", "data.frame")
# check_attr(test, "class", "not.data.frame")
is_url(c("google.com", "http://google.com"))
```

```

is_ip(c("163.114.132.0", "7.114.132", "0.0.0.0", "1.1.1.1"))

are_id(1:10)
are_id(LETTERS[1:10])

are_constant(rep(1, 10))
are_constant(1:10)

are_binary(c("A", "B", "A"))

is_even(1:5)
is_odd(1:5)
is_odd(c(0, 1.5, 2.5, NA, Inf, NULL))

```

chr2num*Check character values for date/numeric/logical and change datatype*

Description

Automatically check a vector, data.frame or list for numeric, logical, date content and change their datatype. Note that factors are skipped in case the user requires character numeric values to be kept as they are.

Usage

```

chr2num(data)

chr2logical(data)

chr2date(data)

```

Arguments

data	Vector, data.frame or list
------	----------------------------

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
str(chr2num(c("1", "2", "3")))
df <- data.frame(A = c("1", "3"), B = c("A", "B"), c = c(pi, pi * 2))
str(chr2num(df))
lst <- list(A = c("1", "2", "3"), B = c("A", "B", "3"), C = pi, D = 3L)
str(chr2num(lst))
lst2 <- list(layer1 = ":D", layer2 = lst)
str(chr2num(lst2))
str(chr2logical(c(NA, "true", FALSE)))
```

ci_lower

Lower/Upper Confidence Intervals

Description

Calculate lower and upper confidence intervals given a mean, standard deviation, sample size, and confidence level. You may want to use `ci_var()` to calculate all values quickly.

Usage

```
ci_lower(mean, ssd, n, conf = 0.95)
ci_upper(mean, ssd, n, conf = 0.95)
```

Arguments

<code>mean</code>	Numeric. Mean: <code>mean(var, na.rm = TRUE)</code>
<code>ssd</code>	Numeric. Standard deviation: <code>sd(var, na.rm = TRUE)</code>
<code>n</code>	Integer. Amount of observations: <code>n()</code>
<code>conf</code>	Numeric (0-1). Confidence level.

Value

Vector with confidence limit value.

See Also

Other Confidence: `ci_var()`

Examples

```
ci_lower(100, 5, 10)
ci_upper(100, 5, 10)
```

ci_var*Confidence Intervals on Dataframe***Description**

Calculate confidence intervals for a continuous numerical column on a dataframe, given a confidence level. You may also group results using another variable. Tidyverse friendly.

Usage

```
ci_var(df, var, group_var = NULL, conf = 0.95)
```

Arguments

<code>df</code>	Dataframe
<code>var</code>	Variable name. Must be a numerical column.
<code>group_var</code>	Variable name. Group results by another variable.
<code>conf</code>	Numeric. Confidence level (0-1).

Value

data.frame mean, standard deviation, counter, upper and lower CIs.

See Also

Other Confidence: [ci_lower\(\)](#)

Examples

```
data(dft) # Titanic dataset
ci_var(dft, Fare)
ci_var(dft, Fare, Pclass)
ci_var(dft, Fare, Pclass, conf = 0.99)
```

cleanText*Clean text strings automatically***Description**

`cleanText`: Clean character strings automatically. Options to keep ASCII characters only, keep certain characters, lower caps, title format, are available.

`cleanNames`: Resulting names are unique and consist only of the _ character, numbers, and ASCII letters. Capitalization preferences can be specified using the `lower` parameter.

Usage

```
cleanText(
  text,
  spaces = TRUE,
  keep = "",
  lower = TRUE,
  ascii = TRUE,
  title = FALSE
)

cleanNames(df, num = "x", keep = "_", ...)
```

Arguments

text	Character Vector
spaces	Boolean. Keep spaces? If character input, spaces will be transformed into passed argument.
keep	Character. String (concatenated or as vector) with all characters that are accepted and should be kept, in addition to alphanumeric.
lower	Boolean. Transform all to lower case?
ascii	Boolean. Only ASCII characters?
title	Boolean. Transform to title format (upper case on first letters).
df	data.frame/tibble.
num	Add character before only-numeric names.
...	Additional parameters passed to <code>cleanText()</code> .

Details

Inspired by `janitor::clean_names`.

Value

Character vector with transformed strings.
data.frame/tibble with transformed column names.

See Also

Other Data Wrangling: `balance_data()`, `categ_reducer()`, `date_cuts()`, `date_feats()`, `file_name()`, `formatHTML()`, `holidays()`, `impute()`, `left()`, `normalize()`, `num_abbr()`, `ohe_commas()`, `ohse()`, `quants()`, `removenacols()`, `replaceall()`, `replacefactor()`, `textFeats()`, `textTokenizer()`, `vector2text()`, `year_month()`, `zerovar()`

Other Text Mining: `ngrams()`, `remove_stopwords()`, `replaceall()`, `sentimentBreakdown()`, `textCloud()`, `textFeats()`, `textTokenizer()`, `topics_rake()`

Examples

```
cleanText("Bernardo Lares 123")
cleanText("Bérnárdo LáreS 123", lower = FALSE)
cleanText("Bernardo Lare$", spaces = ".", ascii = FALSE)
cleanText("\@\@í÷å %ñS ..-X", spaces = FALSE)
cleanText(c("maría", "€", "núñez_a."), title = TRUE)
cleanText("29_Feb-92()#", keep = c("#", "_"), spaces = FALSE)

# For a data.frame directly:
df <- dft[1:5, 1:6] # Dummy data
colnames(df) <- c("ID.", "34", "x_2", "Num 123", "Nòn-äscì", "white Spaces ")
print(df)
cleanNames(df)
cleanNames(df, lower = FALSE)
```

Description

This function lets the user cluster a whole data.frame automatically. As you might know, the goal of kmeans is to group data points into distinct non-overlapping subgroups. If needed, one hot encoding will be applied to categorical values automatically with this function. For consideration: Scale/standardize the data when applying kmeans. Also, kmeans assumes spherical shapes of clusters and does not work well when clusters are in different shapes such as elliptical clusters.

Usage

```
clusterKmeans(
  df,
  k = NULL,
  wss_var = 0,
  limit = 15,
  drop_na = TRUE,
  ignore = NULL,
  ohse = TRUE,
  norm = TRUE,
  algorithm = c("Hartigan-Wong", "Lloyd", "Forgy", "MacQueen"),
  dim_red = "PCA",
  comb = c(1, 2),
  seed = 123,
  quiet = FALSE,
  ...
)
```

Arguments

df	Dataframe
k	Integer. Number of clusters
wss_var	Numeric. Used to pick automatic k value, when k is NULL based on WSS variance while considering limit clusters. Values between (0, 1). Default value could be 0.05 to consider convergence.
limit	Integer. How many clusters should be considered?
drop_na	Boolean. Should NA rows be removed?
ignore	Character vector. Names of columns to ignore.
ohse	Boolean. Do you wish to automatically run one hot encoding to non-numerical columns?
norm	Boolean. Should the data be normalized?
algorithm	character: may be abbreviated. Note that "Lloyd" and "Forgy" are alternative names for one algorithm.
dim_red	Character. Select dimensionality reduction technique. Pass any of: c("PCA", "tSNE", "all", "none").
comb	Vector. Which columns do you wish to plot? Select which two variables by name or column position.
seed	Numeric. Seed for reproducibility
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters to pass sub-functions.

Value

List. If no k is provided, contains nclusters and nclusters_plot to determine optimal k given their WSS (Within Groups Sum of Squares). If k is provided, additionally we get:

- df data.frame with original df plus cluster column
- clusters integer which is the same as k
- fit kmeans object used to fit clusters
- means data.frame with means and counts for each cluster
- correlations plot with correlations grouped by clusters
- PCA list with PCA results (when dim_red="PCA")
- tSNE list with t-SNE results (when dim_red="tSNE")

See Also

Other Clusters: [clusterOptimalK\(\)](#), [clusterVisualK\(\)](#), [reduce_pca\(\)](#), [reduce_tsne\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data("iris")
df <- subset(iris, select = c(-Species))

# If dataset has +5 columns, feel free to reduce dimensionalities
# with reduce_pca() or reduce_tsne() first

# Find optimal k
check_k <- clusterKmeans(df, limit = 10)
check_k$nclusters_plot
# Or pick k automatically based on WSS variance
check_k <- clusterKmeans(df, wss_var = 0.05, limit = 10)
# You can also use our other functions:
# clusterOptimalK(df) and clusterVisualK(df)

# Run with selected k
clusters <- clusterKmeans(df, k = 3)
names(clusters)

# Cross-Correlations for each cluster
plot(clusters$correlations)

# PCA Results (when dim_red = "PCA")
plot(clusters$PCA$plot_explained)
plot(clusters$PCA$plot)
```

clusterOptimalK

Visualize K-Means Clusters for Several K Methods

Description

Visualize cluster data for assorted values of k and methods such as WSS, Silhouette and Gap Statistic. See `factoextra::fviz_nbclust` for more.

Usage

```
clusterOptimalK(
  df,
  method = c("wss", "silhouette", "gap_stat"),
  drop_na = TRUE,
  ohse = TRUE,
  norm = TRUE,
  quiet = TRUE,
  ...
)
```

Arguments

df	Dataframe
method	Character vector.
drop_na	Boolean. Should NA rows be removed?
ohse	Boolean. Do you wish to automatically run one hot encoding to non-numerical columns?
norm	Boolean. Should the data be normalized?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters passed to factoextra::fviz_nbclust

Value

Plot. Optimal number of clusters of df data.frame given a selected method.

See Also

Other Clusters: [clusterKmeans\(\)](#), [clusterVisualK\(\)](#), [reduce_pca\(\)](#), [reduce_tsne\(\)](#)

Examples

```
# You must have "factoextra" library to use this auxiliary function:
## Not run:
data("iris")
df <- subset(iris, select = c(-Species))
# Calculate and plot optimal k clusters
clusterOptimalK(df)

## End(Not run)
```

Description

Visualize cluster data for assorted values of k.

Usage

```
clusterVisualK(df, ks = 2:6, ...)
```

Arguments

df	Dataframe
ks	Integer vector. Which k should be tested?
...	Additional parameters passed to clusterKmeans

Value

List. Plot and data.frame results of clustering df data.frame into ks integer clusters.

See Also

Other Clusters: [clusterKmeans\(\)](#), [clusterOptimalK\(\)](#), [reduce_pca\(\)](#), [reduce_tsne\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data("iris")
df <- subset(iris, select = c(-Species))
df <- df[sample(nrow(df)), ]

# Calculate and plot
result <- clusterVisualK(df, ks = 2:4)
plot(result$plot)

# You can use the data generated as well
lapply(result$data, function(x) head(x$cluster, 10))
```

conf_mat

Confussion Matrix

Description

This function calculates a Confussion Matrix using crosstab for 2 or more categories. You can either set the score and threshold or the labels you wish to cross with.

Usage

```
conf_mat(tag, score, thresh = 0.5, sense = ">=", diagonal = TRUE, plot = FALSE)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result
thresh	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
sense	Character. Inequation sense for threshold: <, <=, >=, >
diagonal	Boolean. FALSE to convert diagonal numbers to zeroes. Ideal to detect must confusing categories.
plot	Boolean. Plot result? Uses <code>mplot_conf()</code>

Details

You may use `mplot_conf()` or set `plot=TRUE`.

Value

`data.frame`. Result of counting tag and score's tag given a threshold, similar to `base::table()`.

See Also

Other Machine Learning: [ROC\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

Other Model metrics: [ROC\(\)](#), [errors\(\)](#), [gain_lift\(\)](#), [loglossBinary\(\)](#), [model_metrics\(\)](#)

Examples

```
data(dfr) # Results for AutoML Predictions
lapply(dfr[c(1, 2)], head)

# Results for Binomial Model
conf_mat(dfr$class2$tag, dfr$class2$scores)
conf_mat(dfr$class2$tag, dfr$class2$scores, thresh = 0.3)
conf_mat(dfr$class2$tag, dfr$class2$scores, sense = "<=")

# Results for Multi-Categorical Model
conf_mat(dfr$class3$tag, dfr$class3$score)
```

corr

Correlation table

Description

This function correlates a whole dataframe, running one hot smart encoding (ohse) to transform non-numerical features. Note that it will automatically suppress columns with less than 3 non missing values and warn the user.

Usage

```
corr(
  df,
  method = "pearson",
  use = "pairwise.complete.obs",
  pvalue = FALSE,
  padjust = NULL,
  half = FALSE,
  dec = 6,
  ignore = NULL,
  dummy = TRUE,
```

```

redundant = NULL,
logs = FALSE,
limit = 10,
top = NA,
...
)

```

Arguments

df	Dataframe. It doesn't matter if it's got non-numerical columns: they will be filtered.
method	Character. Any of: c("pearson", "kendall", "spearman").
use	Character. Method for computing covariances in the presence of missing values. Check <code>stats::cor</code> for options.
pvalue	Boolean. Returns a list, with correlations and statistical significance (p-value) for each value.
padjust	Character. NULL to skip or any of <code>p.adjust.methods</code> to calculate adjust p-values for multiple comparisons using <code>p.adjust()</code> .
half	Boolean. Return only half of the matrix? The redundant symmetrical correlations will be NA.
dec	Integer. Number of decimals to round correlations and p-values.
ignore	Vector or character. Which column should be ignored?
dummy	Boolean. Should One Hot (Smart) Encoding (<code>ohse()</code>) be applied to categorical columns?
redundant	Boolean. Should we keep redundant columns? i.e. If the column only has two different values, should we keep both new columns? Is set to NULL, only binary variables will dump redundant columns.
logs	Boolean. Calculate $\log(x)+1$ for numerical columns?
limit	Integer. Limit one hot encoding to the n most frequent values of each column. Set to NA to ignore argument.
top	Integer. Select top N most relevant variables? Filtered and sorted by mean of each variable's correlations.
...	Additional parameters passed to <code>ohse</code> , <code>corr</code> , and/or <code>cor.test</code> .

Value

`data.frame`. Squared dimensions (N x N) to match every correlation between every `df` `data.frame` column/variable. Notice that when using `ohse()` you may get more dimensions.

See Also

Other Calculus: [dist2d\(\)](#), [model_metrics\(\)](#), [quants\(\)](#)

Other Correlations: [corr_cross\(\)](#), [corr_var\(\)](#)

Examples

```
data(dft) # Titanic dataset
df <- dft[, 2:5]

# Correlation matrix (without redundancy)
corr(df, half = TRUE)

# Ignore specific column
corr(df, ignore = "Pclass")

# Calculate p-values as well
corr(df, pvalue = TRUE, limit = 1)

# Test when no more than 2 non-missing values
df$trash <- c(1, rep(NA, nrow(df) - 1))
# and another method...
corr(df, method = "spearman")
```

corr_cross

Ranked cross-correlation across all variables

Description

This function creates a correlation full study and returns a rank of the highest correlation variables obtained in a cross-table.

Usage

```
corr_cross(
  df,
  plot = TRUE,
  pvalue = TRUE,
  max_pvalue = 1,
  type = 1,
  max = 1,
  top = 20,
  local = 1,
  ignore = NULL,
  contains = NA,
  grid = TRUE,
  rm.na = FALSE,
  quiet = FALSE,
  ...
)
```

Arguments

<code>df</code>	Dataframe. It doesn't matter if it's got non-numerical columns: they will be filtered.
<code>plot</code>	Boolean. Show and return a plot?
<code>pvalue</code>	Boolean. Returns a list, with correlations and statistical significance (p-value) for each value.
<code>max_pvalue</code>	Numeric. Filter non-significant variables. Range (0, 1]
<code>type</code>	Integer. Plot type. 1 is for overall rank. 2 is for local rank.
<code>max</code>	Numeric. Maximum correlation permitted (from 0 to 1)
<code>top</code>	Integer. Return top n results only. Only valid when type = 1. Set value to NA to use all cross-correlations
<code>local</code>	Integer. Label top n local correlations. Only valid when type = 2
<code>ignore</code>	Vector or character. Which column should be ignored?
<code>contains</code>	Character vector. Filter cross-correlations with variables that contains certain strings (using any value if vector used).
<code>grid</code>	Boolean. Separate into grids?
<code>rm.na</code>	Boolean. Remove NAs?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters passed to <code>corr</code>

Details

DataScience+ Post: [Find Insights with Ranked Cross-Correlations](#)

Value

Depending on input `plot`, we get correlation and p-value results for every combination of features, arranged by descending absolute correlation value, with a data.frame `plot = FALSE` or plot `plot = TRUE`.

See Also

Other Correlations: [corr\(\)](#), [corr_var\(\)](#)

Other Exploratory: [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

# Only data with no plot
corr_cross(dft, plot = FALSE, top = 10)

# Show only most relevant results filtered by pvalue
```

```
corr_cross(dft, rm.na = TRUE, max_pvalue = 0.05, top = 15)

# Cross-Correlation for certain variables
corr_cross(dft, contains = c("Survived", "Fare"))

# Cross-Correlation max values per category
corr_cross(dft, type = 2, top = NA)
```

corr_var*Correlation between variable and dataframe***Description**

This function correlates a whole dataframe with a single feature. It automatically runs ohse (one-hot-smart-encoding) so no need to input only numerical values.

Usage

```
corr_var(
  df,
  var,
  ignore = NULL,
  trim = 0,
  clean = FALSE,
  plot = TRUE,
  top = NA,
  ceiling = 1,
  max_pvalue = 1,
  limit = 10,
  ranks = FALSE,
  zeroes = FALSE,
  save = FALSE,
  quiet = FALSE,
  ...
)

## S3 method for class 'corr_var'
plot(x, var, max_pvalue = 1, top = NA, limit = NULL, ...)
```

Arguments

- | | |
|------------|---|
| df | Dataframe. It doesn't matter if it's got non-numerical columns: they will be filtered. |
| var | Variable. Name of the variable to correlate. Note that if the variable var is not numerical, 1. you may define which category to select from using 'var_category'; 2. You may have to add redundant = TRUE to enable all categories (instead of n-1). |

<code>ignore</code>	Character vector. Which columns do you wish to exclude?
<code>trim</code>	Integer. Trim words until the nth character for categorical values (applies for both, target and values)
<code>clean</code>	Boolean. Use <code>lares::cleanText</code> for categorical values (applies for both, target and values)
<code>plot</code>	Boolean. Do you wish to plot the result? If set to TRUE, the function will return only the plot and not the result's data
<code>top</code>	Integer. If you want to plot the top correlations, define how many
<code>ceiling</code>	Numeric. Remove all correlations above... Range: (0-1]
<code>max_pvalue</code>	Numeric. Filter non-significant variables. Range (0, 1]
<code>limit</code>	Integer. Limit one hot encoding to the n most frequent values of each column. Set to NA to ignore argument.
<code>ranks</code>	Boolean. Add ranking numbers?
<code>zeroes</code>	Do you wish to keep zeroes in correlations too?
<code>save</code>	Boolean. Save output plot into working directory
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters passed to <code>corr</code> and <code>cor.test</code>
<code>x</code>	<code>corr_var</code> object

Value

`data.frame`. With variables, correlation and p-value results for each feature, arranged by descending absolute correlation value.

See Also

Other Exploratory: [corr_cross\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)
 Other Correlations: [corr\(\)](#), [corr_cross\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

corr_var(dft, Survived, method = "spearman", plot = FALSE, top = 10)

# With plots, results are easier to compare:

# Correlate Survived with everything else and show only significant results
dft %>% corr_var(Survived_TRUE, max_pvalue = 0.05)

# Top 15 with less than 50% correlation and show ranks
dft %>% corr_var(Survived_TRUE, ceiling = .6, top = 15, ranks = TRUE)
```

cran_logs*Download and plot daily downloads of CRAN packages*

Description

Download daily downloads stats from CRAN for any package, and plot. It can also be used as an auxiliary function to plot (`cranlogs::cran_downloads`) results.

Usage

```
cran_logs(  
  input = "lares",  
  from = Sys.Date() - 31,  
  to = Sys.Date() - 1,  
  type = "daily",  
  plot = TRUE  
)
```

Arguments

<code>input</code>	Character vector with package names or data.frame product of <code>cranlogs::cran_downloads</code> .
<code>from, to</code>	Dates. Range of dates to fetch downloads metrics.
<code>type</code>	Character. Any of: "daily" or "total".
<code>plot</code>	Boolean. Create a plot?

Value

List with data.frame and plot if `plot=TRUE`.

Examples

```
cran_logs(c("lares", "dplyr"), from = "2021-05-31")
```

crosstab*Weighted Cross Tabulation*

Description

A cross-tabulation function with output similar to STATA, tidy friendly, with weighting possibility.

Usage

```
crosstab(
  df,
  ...,
  wt = NULL,
  prow = FALSE,
  pcol = FALSE,
  pall = FALSE,
  decimals = 2,
  rm.na = FALSE,
  total = TRUE,
  order = TRUE
)
```

Arguments

<code>df</code>	Data.frame.
<code>...</code>	Variables. Dependent and independent variables.
<code>wt</code>	Variable, numeric. Weights.
<code>prow, pcol, pall</code>	Boolean. Calculate percent values for rows, columns, or the whole table, respectively.
<code>decimals</code>	Integer. How many decimals should be returned?
<code>rm.na</code>	Boolean. Remove NA values?
<code>total</code>	Boolean. Return total values column?
<code>order</code>	Boolean. Sort columns and rows by frequencies? Else, will be sorted alphabetically

Value

`data.frame`. Result of crossing the variables provided in `...` and counting how many observations (rows) fall into each criteria.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Examples

```
data(dft) # Titanic dataset
crosstab(dft, Survived, Pclass, total = FALSE)
# Show values in percentages
crosstab(dft, Pclass, Survived, prow = TRUE)
crosstab(dft, Pclass, Survived, pall = TRUE)
# Weighted by another variable
crosstab(dft, Survived, Pclass, wt = Fare, prow = TRUE)
```

dalex_local*DALEX Local*

Description

DALEX function for local interpretations

Usage

```
dalex_local(explainer, observation = NA, row = 1, type = "break_down")
```

Arguments

explainer	Object. Result from h2o_explainer function
observation	Data.frame. If you want to use an observation that was not in the original explainer function, add here. Else, use row
row	Dataframe. Row number from the data.frame used in explainer.
type	Character. The type of variable attributions. Either shap, oscillations, break_down or break_down_interactions.

Value

List. Containing observation, breakdown results, and breakdown plot.

See Also

Other Interpretability: [dalex_residuals\(\)](#), [dalex_variable\(\)](#), [h2o_explainer\(\)](#)

dalex_residuals*DALEX Residuals*

Description

DALEX function for residuals

Usage

```
dalex_residuals(explainer)
```

Arguments

explainer	Object. Result from h2o_explainer function
-----------	--

Value

Plot. Based of explainer residual results.

See Also

Other Interpretability: [dalex_local\(\)](#), [dalex_variable\(\)](#), [h2o_explainer\(\)](#)

dalex_variable

DALEX Partial Dependency Plots (PDP)

Description

DALEX auxiliary function for creating Partial Dependency Plots and study variable's responses vs independent vector.

Usage

```
dalex_variable(explainer, vars, force_class = NA, seed = 123, ...)
```

Arguments

<code>explainer</code>	Object. Result from <code>h2o_explainer</code> function.
<code>vars</code>	Character vector. Which features do you wish to study?
<code>force_class</code>	Character. If you wish to force a class on your vars, which one do you need?
<code>seed</code>	Numeric. Seed for reproducibility
<code>...</code>	Additional parameters passed to <code>model_profile</code> .

Value

List. Containing PDP results, plot and vars input.

See Also

Other Interpretability: [dalex_local\(\)](#), [dalex_residuals\(\)](#), [h2o_explainer\(\)](#)

Examples

```
# You must have "DALEX" library to use this auxiliary function:
## Not run:
# Having an "explainer" object created with \code{h2o_explainer}:
# For numerical variables
dalex_variable(explainer, vars = c("Age", "Fare"))
# For categorical variables
dalex_variable(explainer, vars = c("Pclass", "Sex"))

## End(Not run)
```

date_cuts	<i>Convert Date into Year + Cut</i>
-----------	-------------------------------------

Description

This function returns categorical values for any date(s) using year cuts such as bimonths, quarters, terms, and halves.

Usage

```
date_cuts(date = Sys.Date(), type = "Q")
```

Arguments

date	Date. Date we wish to transform
type	Character. Any of the following: B (2 months), Q (3 months), T (4 months), H (6 months)

Value

Vector with date cut for each date

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
date_cuts(Sys.Date(), type = "Q")
date_cuts(Sys.Date(), type = "H")
```

date_feats	<i>One Hot Encoding for Date/Time Variables (Dummy Variables)</i>
------------	---

Description

This function lets the user automatically create new columns out of a dataframe or vector with date/time variables.

Usage

```
date_feats(
  dates,
  drop = FALSE,
  only = NA,
  append = FALSE,
  holidays = FALSE,
  country = "Venezuela",
  currency_pair = NA,
  quiet = FALSE
)
```

Arguments

<code>dates</code>	Vector or dataframe. Non-date/time columns will be automatically ignored/extracted.
<code>drop</code>	Boolean. Should the original date/time columns be kept in the results? Only valid when input is a dataframe.
<code>only</code>	Character or vector. Which columns do you wish to process? If non are explicitly defined, all will be processed
<code>append</code>	Boolean. Append results to existing data.frame? If FALSE, only calculated values will be returned.
<code>holidays</code>	Boolean. Include holidays as new columns?
<code>country</code>	Character or vector. For which countries should the holidays be included?
<code>currency_pair</code>	Character. Which currency exchange do you wish to get the history from? i.e, USD/COP, EUR/USD...
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

data.frame with additional features calculated out of time or date vectors.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Feature Engineering: [holidays\(\)](#), [ohse\(\)](#)

Other One Hot Encoding: [holidays\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#)

Examples

```
df <- data.frame(
  dates = sample(seq(Sys.Date() - 365, Sys.Date(), by = 1), 50),
  times = sample(seq(Sys.time() - 1e+07, Sys.time(), by = 1), 50)
)
```

```

# Input as a vector or dataframe
date_feats(df, drop = TRUE, quiet = TRUE) %>% head(10)

# Holidays given a date range and country
## Not run:
hol <- date_feats(
  seq(Sys.Date() - 365, Sys.Date(), by = 1),
  holidays = TRUE,
  country = "Venezuela"
)
head(hol[!is.na(hol$holiday_name), ])

## End(Not run)

```

db_download*Download/Import Dropbox File by File's Name***Description**

This function lets the user download a file from Dropbox, specifying its name, using a previously created token or with interactive window.

Usage

```

db_download(
  query,
  local_path = NULL,
  xlsx = TRUE,
  token_dir = NA,
  token_name = "token_pers.rds",
  quiet = FALSE
)

```

Arguments

<code>query</code>	Search string. This string is split (on spaces) into individual words. Files will be used if they contain all words in the search string.
<code>local_path</code>	Character. Path to save file to. If <code>NULL</code> (the default), saves file to working directory with same name. If not, but a valid folder, file will be saved in this folder with same basename as path. If not <code>NULL</code> and not a folder, file will be saved to this path exactly.
<code>xlsx</code>	Boolean. Is it an Excel file? Can be returned as a list for each tab and not as a file if needed. Will delete downloaded file.
<code>token_dir</code>	Character. RDS with token local directory. You may set to <code>NA</code> if you already set your credentials (see <code>get_creds()</code>)
<code>token_name</code>	Character. RDS file name with your token's data.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

If query returns a .xlsx file and xlsx=TRUE, will return a data.frame. Else, local_path string.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Other Credentials: [db_upload\(\)](#), [encrypt_file\(\)](#), [get_credentials\(\)](#), [get_tweets\(\)](#), [mail_send\(\)](#), [queryDB\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#), [stocks_file\(\)](#)

Other Dropbox: [db_upload\(\)](#)

Examples

```
## Not run:
# Download a specific file
db_download("stocksReport.Rmd", local_path = "~/Desktop/generic.Rmd")
# Import an Excel file from Dropbox into a data.frame
df <- db_download("Portfolio LC.xlsx", xlsx = FALSE)

## End(Not run)
```

db_upload*Upload Local Files to Dropbox***Description**

This function lets the user upload a local file to Dropbox, using a previously created token or with interactive window.

Usage

```
db_upload(
  filename,
  dir,
  delete_file = FALSE,
  token_dir = NA,
  token_name = "token_pers.rds"
)
```

Arguments

<code>filename</code>	String. Local file's name to upload.
<code>dir</code>	String. Directory you wish to upload the file to.
<code>delete_file</code>	Boolean. Delete local file after uploading?
<code>token_dir</code>	Character. RDS with token local directory. You may set to NA if you already set your credentials (see <code>get_creds()</code>)
<code>token_name</code>	Character. RDS file name with your token's data.

Value

TRUE when successfully uploads file.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Other Credentials: `db_download()`, `encrypt_file()`, `get_credentials()`, `get_tweets()`, `mail_send()`, `queryDB()`, `queryGA()`, `slackSend()`, `stocks_file()`

Other Dropbox: `db_download()`

Description

List with categorical (2 and 3 classes) and continuous predictions, generated with `h2o_automl()` and the `dft`. Note that the models per se won't work to predict.

Usage

```
data(dfr)
```

Format

An object of class "list" with 3 "data.frame"

class2 Predictions for a Binomial Classification Model

class3 Predictions for a Multi-Categorical Classification Model

regr Predictions for a Continuous Regression Model

Value

List

See Also

Other Dataset: [dft](#)

Examples

```
data(dfr)
lapply(dfr, head)
```

dft

Titanic Dataset

Description

The sinking of the Titanic is one of the most infamous shipwrecks in history. On April 15, 1912, during her maiden voyage, the widely considered "unsinkable" RMS Titanic sank after colliding with an iceberg. Unfortunately, there weren't enough lifeboats for everyone onboard, resulting in the death of 1502 out of 2224 passengers and crew. While there was some element of luck involved in surviving, it seems some groups of people were more likely to survive than others. This dataset contains the details of a subset of the passengers on board (891 to be exact) taken from Kaggle's Titanic [Train.csv](#).

Usage

```
data(dft)
```

Format

An object of class "data.frame"

PassengerId Unique ID for each passenger (1-891)

Survived Did the passenger survive? (TRUE, FALSE)

Pclass Ticket class, from first to third (1, 2, 3)

Sex Gender (female, male)

Age Age for each passenger in years (0.42-80)

SibSp Amount of siblings / spouses aboard the Titanic (0-8)

Parch Amount of parents / children aboard the Titanic (0-6)

Ticket Ticket IDs

Fare Amount paid for passenger's ticket (0-512.3292)

Cabin width of top of diamond relative to widest point (43-95)

Embarked Port of Embarkation (43-95)

Value

```
data.frame
```

See Also

Other Dataset: [dfr](#)

Examples

```
data(dft)
head(dft)
```

df_str*Dataset columns and rows structure*

Description

This function lets the user to check quickly the structure of a dataset (data.frame). It returns multiple counters for useful metrics, a plot, and a list of column names for each of the column metrics.

Usage

```
df_str(df, return = "plot", subtitle = NA, quiet = FALSE)
```

Arguments

df	Dataframe
return	Character. Return "skimr" for skim report, "numbers" for stats and numbers, "names" for a list with the column names of each of the class types, "plot" for a nice plot with "numbers" output, "distr" for an overall summary plot showing categorical, numeric, and missing values by using <code>plot_df</code> distributions
subtitle	Character. Add subtitle to plot
quiet	Boolean. Keep quiet or show other options available?

Value

Depending on `return` input and based on your `df` structure:

- list with the names of the columns classified by class
- `data.frame` with numbers: total values, row, columns, complete rows
- `plot` with visualizations

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset
df_str(dft, "names")
df_str(dft, "numbers", quiet = TRUE)
df_str(dft, "plot", quiet = TRUE)
```

dist2d

Distance from specific point to line

Description

This function lets the user calculate the mathematical linear distance Between a specific point and a line (given geometrical 3 points)

Usage

```
dist2d(x, a = c(0, 0), b = c(1, 1))
```

Arguments

- x Vector. Coordinates of the point from which we want to measure the distance
- a Vector. Coordinates of 1st point over the line
- b Vector. Coordinates of 2st point over the line

Value

Numeric value result

See Also

Other Calculus: [corr\(\)](#), [model_metrics\(\)](#), [quants\(\)](#)

Examples

```
dist2d(x = c(5, 2))
dist2d(x = c(5, 2), a = c(0, 0), b = c(0, 1))
dist2d(x = c(5, 2), a = c(0, 0), b = c(1, 0))
```

distr	<i>Compare Variables with their Distributions</i>
--------------	---

Description

Compare the distribution of a target variable vs another variable. This function automatically splits into quantiles for numerical variables. Custom and tidyverse friendly.

Usage

```
distr(
  data,
  ...,
  type = 1,
  ref = TRUE,
  note = NA,
  top = 10,
  breaks = 10,
  na.rm = FALSE,
  force = "none",
  trim = 0,
  clean = FALSE,
  abc = FALSE,
  custom_colours = FALSE,
  plot = TRUE,
  chords = FALSE,
  save = FALSE,
  subdir = NA
)
```

Arguments

data	Dataframe
...	Variables. Main (target variable) and secondary (values variable) to group by (if needed).
type	Integer. 1 for both plots, 2 for counter plot only, 3 for percentages plot only.
ref	Boolean. Show a reference line if levels = 2? Quite useful when data is unbalanced (not 50/50) because a reference line is drawn.
note	Character. Caption for the plot.
top	Integer. Filter and plot the most n frequent for categorical values.
breaks	Integer. Number of splits for numerical values.
na.rm	Boolean. Ignore NAs if needed.
force	Character. Force class on the values data. Choose between 'none', 'character', 'numeric', 'date'

<code>trim</code>	Integer. Trim labels until the nth character for categorical values (applies for both, target and values)
<code>clean</code>	Boolean. Use <code>cleanText()</code> for categorical values (applies for both, target and values)
<code>abc</code>	Boolean. Do you wish to sort by alphabetical order?
<code>custom_colours</code>	Boolean. Use custom colours function?
<code>plot</code>	Boolean. Return a plot? Otherwise, a table with results
<code>chords</code>	Boolean. Use a chords plot?
<code>save</code>	Boolean. Save the output plot in our working directory
<code>subdir</code>	Character. Into which subdirectory do you wish to save the plot to?

Value

Plot when `plot=TRUE` with two plots in one: counter distribution grouped by cuts, and proportions distribution grouped by same cuts. `data.frame` when `plot=FALSE` with counting, percentages, and cumulative percentages results. When `type` argument is used, single plots will be returned.

See Also

Other Exploratory: `corr_cross()`, `corr_var()`, `crosstab()`, `df_str()`, `freqs()`, `freqs_df()`, `freqs_list()`, `freqs_plot()`, `lasso_vars()`, `missingness()`, `plot_cats()`, `plot_df()`, `plot_nums()`, `tree_var()`
 Other Visualization: `freqs()`, `freqs_df()`, `freqs_list()`, `freqs_plot()`, `noPlot()`, `plot_chord()`, `plot_survey()`, `plot_timeline()`, `tree_var()`

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

# Relation for categorical/categorical values
distr(dft, Survived, Sex)

# Relation for categorical/numeric values
dft %>%
  distr(Survived, Fare, plot = FALSE) %>%
  head(10)
# Sort values
dft %>% distr(Survived, Fare, abc = TRUE)
# Less splits/breaks
dft %>% distr(Survived, Fare, abc = TRUE, breaks = 5)

# Distribution of numerical only
dft[dft$Fare < 20, ] %>% distr(Fare)

# Distribution of numerical/numerical
dft %>% distr(Fare, Age)

# Select only one of the two default plots of distr()
```

```
dft %>% distr(Survived, Age, type = 2)
dft %>% distr(Survived, Age, type = 3)
```

dont_sleep*Prevent Computer from Sleeping by Simulating Mouse Activity*

Description

Keeps the computer awake by moving the mouse cursor and clicking periodically with a human-like behavior. It automatically detects the operating system (Windows or macOS) and runs the appropriate underlying logic.

Usage

```
dont_sleep(sec_range = c(20, 60), off_time = 18.5, quiet = FALSE)

dont_sleep_time(quiet = FALSE)
```

Arguments

sec_range	Numeric vector of length 2. Range (in seconds) of random intervals between mouse movements and clicks. Default is c(20, 60).
off_time	Numeric. Decimal hour (24h format) to stop the function automatically, e.g. 18.5 means 18:30 (6:30 PM). Default is 18.5.
quiet	Logical. If TRUE, suppresses progress messages. Default is FALSE.

Details

The cursor initially moves to the upper-left corner (0,0), then performs small random movements and clicks periodically to avoid system sleep. The function exits if the user moves the mouse away from the corner or when the specified off-time is reached.

Requires `cliclick` installed on macOS (install via Homebrew: `brew install cliclick`), and the `KeyboardSimulator` package on Windows.

Value

Invisibly returns NULL when the function exits.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
## Not run:
# Keep the computer awake until 8:00 PM, with mouse actions every 30 to 60 seconds
dont_sleep(sec_range = c(30, 60), off_time = 20)
# Summary of amount of daily time running function
dont_sleep_time()

## End(Not run)
```

encrypt_file

File Encryption and Decryption (AES-256-CBC)

Description

This set of functions provides utilities for encrypting and decrypting files using AES-256 in CBC mode, primarily for handling key-value pair "secrets" files.

Usage

```
encrypt_file(input_file, output_file, key)

read_encrypted(input_file, key)

write_encrypted(data, output_file = "encrypted.enc", key = NULL, quiet = FALSE)

hex_to_raw(x)

raw_to_hex(x)
```

Arguments

<code>input_file</code>	Character string. The path to the file to be encrypted.
<code>output_file</code>	Character string. The path where the encrypted file will be saved.
<code>key</code>	A raw vector of 32 bytes (256 bits) to be used as the encryption key.
<code>data</code>	List, vector or any json-able object to write into encrypted file.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>x</code>	Character string or hex vector to convert.

Details

The encryption process pads the data, generates a random Initialization Vector (IV), and encrypts the content. The IV is prepended to the encrypted data in the output file. Decryption involves extracting the IV, decrypting, and unpadding the data. A specialized decryption function `read_encrypted` is provided to directly parse decrypted content (assumed to be key=value pairs) into a named R list. If no "=" sign found in the first line, text will be imported as character vector.

Value

No return value, called for side effects (writes to output_file).

See Also

Other Credentials: [db_download\(\)](#), [db_upload\(\)](#), [get_credentials\(\)](#), [get_tweets\(\)](#), [mail_send\(\)](#), [queryDB\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#), [stocks_file\(\)](#)

Examples

```
## Not run:  
# Create dummy files for demonstration  
temp_input <- tempfile(fileext = ".txt")  
temp_output <- tempfile(fileext = ".enc")  
writeLines("library=lares \nversion=5.3.0 \npass=123", temp_input)  
readLines(temp_input)  
  
# Generate a random 32-byte key (and save it in a secure place)  
raw_key <- openssl::rand_bytes(32)  
# You can convert from hex to raw and viceversa  
raw_to_hex(raw_key)  
hex_to_raw(raw_to_hex(raw_key))  
  
# Encrypt the file  
encrypt_file(temp_input, temp_output, raw_key)  
  
# Import the data from encrypted file to list or vector  
secrets <- read_encrypted(temp_output, raw_key)  
unlist(secrets)  
  
# Example using a string (JSON in this case)  
writeLines(jsonlite:: toJSON(list(a = 1, list(b = 2, c = 1:3))), temp_input)  
encrypt_file(temp_input, temp_output, raw_key)  
read_encrypted(temp_output, raw_key)  
  
# Example writing a file from a list or vector directly from R  
my_secrets <- list(  
  api_key = "some_secret_api_key_123",  
  username = "data_analyst",  
  server = "production.server.com"  
)  
raw_key <- write_encrypted(my_secrets, temp_output, quiet = FALSE)  
fromJSON(read_encrypted(temp_output, key = raw_key))  
  
## End(Not run)
```

Description

This function lets the user calculate all errors and R squared simultaneously.
 This function lets the user calculate Root Mean Squared Error
 This function lets the user calculate Mean Absolute Error
 This function lets the user calculate Mean Squared Error
 This function lets the user calculate Mean Squared Error
 This function lets the user calculate R Squared
 This function lets the user calculate Adjusted R Squared

Usage

```
errors(tag, score)

rmse(tag, score)

mae(tag, score)

mse(tag, score)

mape(tag, score)

rsq(tag, score)

rsqa(tag, score)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result

Value

data.frame or numeric values results for multiple error metrics on continuous numerical vectors inputs.

See Also

Other Model metrics: [ROC\(\)](#), [conf_mat\(\)](#), [gain_lift\(\)](#), [loglossBinary\(\)](#), [model_metrics\(\)](#)

Examples

```
data(dfr) # Results for AutoML Predictions
head(dfr$regr)
df <- errors(dfr$regr$tag, dfr$regr$score)
head(df)
```

etf_sector	<i>ETF's Sectors Breakdown</i>
------------	--------------------------------

Description

`etf_sector()` scraps etf.com data for sector breakdown on ETFs. Use `splot_etf()` for visualization.

Usage

```
etf_sector(etf = "VTI", quiet = FALSE, cache = TRUE)
```

Arguments

etf	Character Vector. Which ETFs you wish to scrap?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
cache	Boolean. Use daily cache if available?

Value

data.frame with ETF break.down data by sector

See Also

Other Investment: [splot_summary\(\)](#), [stocks_file\(\)](#), [stocks_quote\(\)](#)

Examples

```
etf_sector(etf = "VTI")
```

export_plot	<i>Export ggplot2, gridExtra, or any plot object into rendered file</i>
-------------	---

Description

Export any ggplot2, gridExtra, or any plot object created with R into rendered png or jpg file.

Usage

```
export_plot(
  p,
  name = "plot",
  vars = NA,
  sep = ".vs.",
  width = 8,
  height = 6,
  format = "png",
  res = 300,
  dir = getwd(),
  subdir = NA,
  quiet = FALSE
)
```

Arguments

<code>p</code>	Plot object. Plot to render and export.
<code>name</code>	Character. File's name or suffix if vars is not NA. No need to include file format on file name.
<code>vars</code>	Vector. Variable names to identify by filename.
<code>sep</code>	Character. Separator for vars.
<code>width, height, res</code>	Numeric. Plot's width, height, and res (for grids).
<code>format</code>	Character. One of: png or jpeg.
<code>dir, subdir</code>	Character. In which directory/subdirectory do you wish to save the plot? Working directory as default dir.
<code>quiet</code>	Boolean. Display successful message with filename when saved?

Value

No return value, called for side effects.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
p <- noPlot()
export_plot(p, name = "noplott", width = 10, height = 8, res = 300, dir = tempdir())
export_plot(p, name = "noplott2", subdir = "newplots", dir = tempdir())
```

export_results	<i>Export h2o_automl's Results</i>
----------------	------------------------------------

Description

Export RDS, TXT, POJO, MOJO and all results from `h2o_automl()`.

Usage

```
export_results(  
  results,  
  thresh = 10,  
  which = c("txt", "csv", "rds", "binary", "mojo", "plots", "dev", "production"),  
  note = NA,  
  subdir = NA,  
  save = TRUE,  
  seed = 0  
)
```

Arguments

results	<code>h2o_automl</code> or <code>h2o</code> model
thresh	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
which	Character vector. Select which file format to export: Possible values: txt, csv, rds, binary, mojo, plots. You might also use dev (txt, csv, rds) or production (binary, mojo) or simply don't use parameter to export everything
note	Character. Add a note to the txt file. Useful when lots of models are trained and saved to remember which one is which one
subdir	Character. In which directory do you wish to save the results?
save	Boolean. Do you wish to save/export results?
seed	Numeric. For reproducible results and random splits.

Value

No return value, called for side effects.

See Also

Other Machine Learning: `ROC()`, `conf_mat()`, `gain_lift()`, `h2o_automl()`, `h2o_predict_MOJO()`, `h2o_selectmodel()`, `impute()`, `iter_seeds()`, `lasso_vars()`, `model_metrics()`, `model_preprocess()`, `msplit()`

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`,

```
glued(), grepM(), h2o_selectmodel(), haveInternet(), image_metadata(), importxlsx(),
ip_data(), json2vector(), list_cats(), listfiles(), mail_send(), markdown2df(), move_files(),
msplit(), myip(), quiet(), read.file(), statusbar(), tic(), try_require(), updateLares(),
warnifnot(), what_size()
```

fb_accounts	<i>Facebook Ad Accounts</i>
-------------	-----------------------------

Description

This returns all ad accounts for a FB Business Account FB. For more information on Ad Insights' API, go to the [original documentaion](#)

Usage

```
fb_accounts(
  token,
  business_id = "904189322962915",
  type = c("owned", "client"),
  limit = 100,
  api_version = NULL,
  ...
)
```

Arguments

<code>token</code>	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
<code>business_id</code>	Character. Business ID.
<code>type</code>	Character vector. Values: owned, client.
<code>limit</code>	Integer. Query limit by pagination.
<code>api_version</code>	Character. Facebook API version.
<code>...</code>	Additional parameters.

Value

`data.frame` with un-nested processed results fetched with API.

See Also

Other API: `bring_api()`, `fb_ads()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`, `gemini_ask()`, `gpt_ask()`, `queryGA()`, `slackSend()`

Other Meta: `fb_ads()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`

Examples

```
## Not run:  
# Query all accounts (owned and with permissions) of a Business ID  
accounts <- fb_accounts(YOURTOKEN, YOURBUSINESS)  
  
## End(Not run)
```

fb_ads

Facebook Ads API

Description

This returns all available FB ads for any account, campaign, or ad set id. For more information on Ad' API, go to the [original documentaion](#)

Usage

```
fb_ads(  
  token,  
  which,  
  start_date = Sys.Date() - 31,  
  end_date = Sys.Date(),  
  fields = NA,  
  api_version = NULL,  
  process = TRUE,  
  ...  
)
```

Arguments

token	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
which	Character vector. This is the accounts, campaigns, adsets, or ads IDs to be queried. Remember: if report_level = "account", you must start the ID with act_.
start_date, end_date	Character. The first and last full day to report, in the format "YYYY-MM-DD".
fields	Character, json format. Leave NA for default fields OR NULL to ignore.
api_version	Character. Facebook API version.
process	Boolean. Process GET results to a more friendly format?
...	Additional parameters.

Details

This function was based on FBinsightsR.

Value

`data.frame` with un-nested processed results if `process=TRUE` or raw API results as list when `process=FALSE`.

See Also

Other API: `bring_api()`, `fb_accounts()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`, `gemini_ask()`, `gpt_ask()`, `queryGA()`, `slackSend()`

Other Meta: `fb_accounts()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`

Examples

```
## Not run:
token <- YOURTOKEN
account <- act_ADACCOUNT

# Query all ads for "which" (account) with results in the last 10 days
ads <- fb_ads(token, account, start_date = Sys.Date() - 10)

## End(Not run)
```

fb_creatives

Facebook Creatives API

Description

For more information: [Marketing API](#)

Usage

```
fb_creatives(token, which, api_version = NULL, process = TRUE, ...)
```

Arguments

<code>token</code>	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
<code>which</code>	Character vector. This is the accounts, campaigns, adsets, or ads IDs to be queried. Remember: if <code>report_level = "account"</code> , you must start the ID with <code>act_</code> .
<code>api_version</code>	Character. Facebook API version.
<code>process</code>	Boolean. Process GET results to a more friendly format?
<code>...</code>	Additional parameters.

Value

`data.frame` with un-nested processed results if `process=TRUE` or raw API results as list when `process=FALSE`.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#)

Examples

```
## Not run:  
token <- YOURTOKEN  
account <- act_ADACCOUNT  
  
# Query all creatives for "which" (account in this case)  
creatives <- fb_creatives(token, account)  
  
## End(Not run)
```

fb_insights

Facebook Insights API

Description

This returns all available FB insights per day including any given breakdown to the specified report level, and place into a data frame. For more information on Ad Insights' API, go to the original [documentaion](#).

Usage

```
fb_insights(  
  token,  
  which,  
  start_date = Sys.Date() - 7,  
  end_date = Sys.Date(),  
  time_increment = "1",  
  report_level = "campaign",  
  ad_object = "insights",  
  breakdowns = NA,  
  fields = NA,  
  filtering = NULL,  
  limit = 100,  
  api_version = NULL,  
  process = TRUE,  
  async = FALSE,  
  ...  
)
```

Arguments

token	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
which	Character vector. This is the accounts, campaigns, adsets, or ads IDs to be queried. Remember: if <code>report_level = "account"</code> , you must start the ID with <code>act_</code> .
start_date, end_date	Character. The first and last full day to report, in the format "YYYY-MM-DD".
time_increment	Character. Group by months ("monthly"), everything together ("all_days") or an integer per days [1-90]. Default: each day separately (i.e. "1").
report_level	Character. One of "ad", "adset", "campaign", or "account"
ad_object	Character. One of: "insights" (default), "adsets", ...
breakdowns	Character Vector. One or more of breakdowns for segmentation results. Set to NA for no breakdowns
fields	Character, json format. Leave NA for default fields OR NULL to ignore.
filtering	List. Each filter will be a list containing "field", "operator", and "value". Read more about the operators in the official docs . Example: <code>dplyr::tibble(field = "country", operator = "IN", value = list("PE"))</code> .
limit	Integer. Query limit by pagination.
api_version	Character. Facebook API version.
process	Boolean. Process GET results to a more friendly format?
async	Boolean. Run an async query. When set to TRUE, instead of making a GET query, it'll run a POST query and will return a report run ID.
...	Additional parameters.

Value

data.frame with un-nested processed results if process=TRUE or raw API results as list when process=FALSE.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)
 Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#)

Examples

```
## Not run:
token <- "YOURTOKEN"
which <- "act_20846447"

# Platforms' Insights: all ad-sets platforms of "which" account,
# aggregated, for the last 30 days
```

```
platforms <- fb_insights(
  token, which,
  start_date = Sys.Date() - 30,
  time_increment = "all_days",
  report_level = "adset",
  fields = c(
    "account_name",
    "adset_id",
    "adset_start",
    "adset_end"
  ),
  breakdowns = c(
    "publisher_platform",
    "platform_position",
    "device_platform"
  )
)

# Daily results for all campaigns of "which" account,
# with custom performance fields with no breakdowns.
insights_adset <- fb_insights(
  token, which,
  time_increment = "1",
  report_level = "campaign",
  fields = c(
    "adset_id",
    "reach",
    "frequency",
    "spend",
    "cpm",
    "objective",
    "optimization_goal"
  )
)

## End(Not run)
```

fb_process*Paginate and Process Facebook's API Results*

Description

Process and paginate raw results from Facebook's API, result of querying the API with `httr::GET` or by passing an API link.

Usage

```
fb_process(input, paginate = TRUE, sleep = 0, quiet = FALSE, ...)
```

Arguments

input	GET's output object (response) or link (character).
paginate	Boolean or integer. Run through all paginations? If set to FALSE, only the first one will be processed. If set to any other integer value, will process the first N paginations.
sleep	Numeric value. How much should each loop wait until running the next pagination query?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.

Value

data.frame with un-nested processed results or NULL if no results found.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#)

fb_report_check

Facebook API Report Status Check

Description

This returns all available FB insights per day including any given breakdown to the specified report level, and place into a data frame. For more information on Ad Insights' API, go to the original [documentaion](#).

Usage

```
fb_report_check(
  token,
  report_run_id,
  api_version = NULL,
  live = FALSE,
  sleep = 10,
  quiet = FALSE
)
```

Arguments

token	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
report_run_id	Integer. Report ID to check status.
api_version	Character. Facebook API version.
live	Boolean. Run until status report is finished?
sleep	Boolean. If live=TRUE, then how many seconds should we wait until next check?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

List with API status results.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#)

Examples

```
## Not run:
token <- "YOURTOKEN"
report_run_id <- "123456789"
fb_report_check(token, report_run_id, live = TRUE, quiet = FALSE)

## End(Not run)
```

fb_rf

Facebook Reach and Frequency API

Description

Create or query reach and frequency predictions using Facebook's Reach and Frequency API. For more information on the API and its parameters, go to the [original documentaion](#).

Usage

```
fb_rf(
  token,
  ad_account = NA,
  prediction = NA,
  objective = "REACH",
  days = 28,
```

```

budget = 2e+06,
destination_ids = NA,
countries = "MX",
frequency_cap = 8,
prediction_mode = 1,
curve = TRUE,
api_version = NULL,
process = TRUE,
...
)

```

Arguments

token	Character. Valid access token with sufficient privileges. Visit the Facebook API Graph Explorer to acquire one.
ad_account	Character. Ad Account. Remember to start with act_. If you use the prediction argument, no need to provide this parameter.
prediction	Integer. Prediction ID if you already created the prediction and wish to query the curve's data. As this prediction already exists, the rest of arguments of this function will be ignored.
objective	Character. Any of: "BRAND_AWARENESS", "LINK_CLICKS", "POST_ENGAGEMENT", "MOBILE_APP_INSTALLS", "CONVERSIONS", "REACH", or "VIDEO_VIEWS".
days	Integer. Amount of days for your campaign's predictions.
budget	Integer. The budget in the Ad Account currency in cents.
destination_ids	Integer vector. Page ID and/or Instagram Account ID.
countries	Character vector. Country's acronyms.
frequency_cap	Integer. Frequency cap over all the campaign duration.
prediction_mode	Integer. "1" for predicting Reach by providing budget, "2" is for predicting Budget given a specific Reach.
curve	Boolean. Return curve data? If not, only prediction will be created.
api_version	Character. Facebook API version.
process	Boolean. Process GET results to a more friendly format?
...	Additional parameters passed to target specs.

Value

data.frame with un-nested processed results if process=TRUE or raw API results as list when process=FALSE.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)
 Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_token\(\)](#)

Examples

```
## Not run:  
token <- "YOURTOKEN"  
account_id <- "act_20846447"  
  
# BASIC 1: Create and return data for a new prediction  
basic1 <- fb_rf(token, account_id, destination_ids = 187071108930, countries = "AR")  
  
# BASIC 2: Fetch data for an existing prediction ID  
basic2 <- fb_rf(token, account_id, prediction = 6317720998974)  
  
# ADVANCED (Fully custom prediction)  
advanced <- fb_rf(token, account_id,  
objective = "REACH",  
days = 28,  
budget = 2000000,  
destination_ids = c(187071108930, 1142958119078556),  
age_min = 15,  
age_max = 65,  
genders = 2,  
countries = "MX",  
publisher_platforms = c(  
    "facebook",  
    "instagram",  
    #' audience_network',  
    "messenger"  
,  
    # interests_ids = NA,  
facebook_positions = c(  
    "feed",  
    #' instant_article',  
    "marketplace",  
    "video_feeds",  
    "story",  
    "search",  
    "instream_video"  
,  
    instagram_positions = c(  
        "stream",  
        "story",  
        "explore"  
,  
    # audience_network_positions = c(  
    # 'classic',  
    # 'instream_video')  
messenger_positions = c(  
    "messenger_home",  
    "sponsored_messages",  
    "story"  
,  
device_platforms = c(  
    "mobile",
```

```

    "desktop"
)
)

## End(Not run)

```

fb_token*Facebook's Long-Life User API Token***Description**

Using a 1-hour generic user token you can generate a 60 day token. You will need to have an App ID and App secret, and a valid token. Generate a new valid User Token with the [API Graph](#).

Usage

```
fb_token(app_id, app_secret, token, api_version = NULL)
```

Arguments

app_id, app_secret	Character. Application ID and Secret.
token	Character. User token, created with API Graph or with this same fb_token()'s token.
api_version	Character. Facebook API version.

Details

More info: [Long-Lived Access Tokens](#)

Value

Character. String with token requested. If successful, it'll contain an attribute called "expiration" with date and time of expiration.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [gemini_ask\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Other Meta: [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#)

filesGD*Google Drive Files (API v4)*

Description

Authenticate and find Google Drive files and IDs by name.

Usage

```
filesGD(title, server = FALSE, json = NULL, api_key = NULL, email = NULL)
```

Arguments

<code>title</code>	Character. Title of Google Drive file. Uses regular expressions so you may fetch with patterns instead of names.
<code>server</code>	Boolean. Force interacting auth process?
<code>json</code>	Character. JSON filename with service auth
<code>email, api_key</code>	Character. If you have multiple pre-authorized accounts in your machine, you may non-interactively select which one you wish to use by email and/or api_key.

Value

Vector with found file names based on `title` on Google Drive.

See Also

Other Scrapper: [get_mp3\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)
Other Google: [gtrends_related\(\)](#), [queryGA\(\)](#), [readGS\(\)](#)

files_functions*List all functions used in R script files by package*

Description

Parses all functions called by an R script and then lists them by package. Wrapper for 'getParseData'. May be of great use for those developing a package to help see what namespace 'imports-From' calls will be required.

Usage

```
files_functions(filename, abc = TRUE, quiet = FALSE)
```

Arguments

- `filename` Character. Path to an R file (or directory) containing R code files.
`abc` Boolean. List functions alphabetically. If FALSE, will list in order of frequency.
`quiet` Boolean. Keep quiet? If not, print messages and statusbar.

Value

`data.frame`. Each row is a function and columns stating number of appearances, percentage, packages, and files searched.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepm()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
## Not run:  

# Choose an R script file with functions  

rfile <- file.choose()  

files_functions(rfile)  

  
## End(Not run)
```

*file_name**Extract file raw name and type from file names***Description**

Extract file raw name and type from file names

Get file extensions without file names

Usage

```
file_name(filepath)  
  
file_type(filepath)
```

Arguments

- `filepath` Character vector. File path(s) to get file raw names without extension nor path OR extension without path nor raw name.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
file_name("file.aux")
file_name("temp/file.R")
file_name("/temp/temp3/music.mp3")
file_type("file.aux")
file_type("temp/file.R")
file_type("/temp/temp3/music.mp3")
```

font_exists *Check if Font is Installed*

Description

This function checks if a font is installed in your machine. To list all available fonts, set `font = NULL`.

Usage

```
font_exists(font = "Arial Narrow", font_dirs = NULL, quiet = FALSE, ...)
```

Arguments

<code>font</code>	Character. Which font to check. No need to add .TFF.
<code>font_dirs</code>	Character vector. Additional directories to check for fonts.
<code>quiet</code>	Boolean. Keep quiet? If not, show message
<code>...</code>	Additional parameters.

Value

Boolean result of the existing fonts check.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepm()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
font_exists(font = "Arial")
font_exists(font = "arial")
font_exists(font = "")
font_exists(font = NULL)
```

`forecast_arima`

ARIMA Forecast

Description

This function automates the ARIMA iterations and modeling for time forecasting. For the moment, units can only be days.

Usage

```
forecast_arima(
  time,
  values,
  n_future = 30,
  ARMA = 8,
  ARMA_min = 5,
  AR = NA,
  MA = NA,
  wd_excluded = NA,
  plot = TRUE,
  plot_days = 90,
  project = NA
)
```

Arguments

<code>time</code>	POSIX. Vector with date values
<code>values</code>	Numeric. Vector with numerical values
<code>n_future</code>	Integer. How many steps do you wish to forecast?
<code>ARMA</code>	Integer. How many days should the model look back for ARMA? Between 5 and 10 days recommended. If set to 0 then it will forecast until the end of max date's month; if set to -1, until the end of max date's following month

ARMA_min	Integer. How many days should the model look back for ARMA? Between 5 and 10 days recommended. If set to 0 then it will forecast until the end of max date's month; if set to -1, until the end of max date's following month
AR	Integer. Force AR value if known
MA	Integer. Force MA value if known
wd_excluded	Character vector. Which weekdays are excluded in your training set. If there are, please define know which ones. Example: c('Sunday','Thursday'). If set to 'auto' then it will detect automatically which weekdays have no data and forcast without these days.
plot	Boolean. If you wish to plot your results
plot_days	Integer. How many days back you wish to plot?
project	Character. Name of your forecast project

Details

The ARIMA method is appropriate only for a time series that is stationary (i.e., its mean, variance, and autocorrelation should be approximately constant through time) and it is recommended that there are at least 50 observations in the input data.

The model consists of two parts, an autoregressive (AR) part and a moving average (MA) part. The AR part involves regressing the variable on its own lagged (i.e., past) values. The MA part involves modeling the error term as a linear combination of error terms occurring contemporaneously and at various times in the past.

One thing to keep in mind when we think about ARIMA models is given by the great power to capture very complex patters of temporal correlation (Cochrane, 1997: 25)

Value

List. Containing the trained model, forecast accuracy results, data.frame for forecast (test) and train, and if `plot=TRUE`, a plot.

See Also

Other Forecast: [prophesize\(\)](#)

Description

Print Coloured Messages

Usage

```
formatColoured(
  txt,
  colour = c("yellow", "blue", "grey"),
  bold = FALSE,
  cat = TRUE
)
```

Arguments

<code>txt</code>	Character. Text to print or transform.
<code>colour</code>	Character. Any of: grey, red, green, yellow, blue, or purple.
<code>bold</code>	Boolean. Set bold text?
<code>cat</code>	Boolean. Print with cat? If not, raw string

Value

Depends on `cat`: NULL if TRUE or character string if FALSE.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
opts <- c("GREY", "RED", "GREEN", "YELLOW", "BLUE", "PURPLE")
for (colour in opts) formatColoured(paste("Colour:", colour, "\n"), colour)
formatColoured("my bold coloured text", bold = TRUE, cat = TRUE)
```

Description

Format any character string to HTML or markdown format. We recommend using this format with the `ggtext::geom_richtext` function to format text in ggplot2 objects.

This function lets the user format numerical values nicely

Usage

```
formatHTML(text, color = "black", size = 20, bold = FALSE)

formatNum(
  x,
  decimals = 2,
  signif = NULL,
  type = Sys.getenv("LARES_NUMFORMAT"),
  pre = "",
  pos = "",
  sign = FALSE,
  abbr = FALSE,
  ...
)
```

Arguments

<code>text</code>	Character. Strings to format.
<code>color</code>	Character. Hex colour code.
<code>size</code>	Numeric. Text size.
<code>bold</code>	Boolean. Should the text be bold?
<code>x</code>	Numerical Vector
<code>decimals</code>	Integer. Amount of decimals to display. If set to NULL, then <code>getOption("digits")</code> will be used.
<code>signif</code>	Integer. Rounds the values in its first argument to the specified number of significant digits.
<code>type</code>	Integer. 1 for International standards. 2 for American Standards. Use <code>Sys.setenv("LARES_NUMFORMAT" = 2)</code> to set this parameter globally.
<code>pre, pos</code>	Character. Add string before or after number.
<code>sign</code>	Boolean. Add + sign to positive values.
<code>abbr</code>	Boolean. Abbreviate using <code>num_abbr()</code> ? You can use the 'decimals' parameter to set <code>abbr</code> 's <code>n(-1)</code> parameter.
<code>...</code>	Additional lazy eval parameters.

Value

String with format characters included.

Character. String vector with reformatted continuous numbers

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`,

```
msplit(), myip(), quiet(), read.file(), statusbar(), tic(), try_require(), updateLares(),
warnifnot(), what_size()

Other Data Wrangling: balance_data(), categ_reducer(), cleanText(), date_cuts(), date_feats(),
file_name(), holidays(), impute(), left(), normalize(), num_abbr(), ohe_commas(), ohse(),
quants(), removenocols(), replaceall(), replacefactor(), textFeats(), textTokenizer(),
vector2text(), year_month(), zerovar()
```

Examples

```
formatHTML("Text test", color = "#000000")
formatHTML(c(123, 456), color = "orange", size = 120, bold = TRUE)

# If you want to use it with \code{ggtext}:
## Not run:
col1 <- "grey"
col2 <- "orange"
pt <- data.frame(
  label = paste0(
    formatHTML(123, color = col2, size = 120, bold = TRUE), "<br/>",
    formatHTML("of children had a", col1), "<br/>",
    formatHTML("traditional stay-at-home mom", color = col2, bold = TRUE), "<br/>",
    formatHTML(paste0("in 2012, compared to ", 321, " in 1970"), color = col1)
  )
)
ggplot(pt, aes(x = 0, y = 0)) +
  ggtext::geom_richtext(
    aes(label = label),
    hjust = 0,
    label.color = NA,
    lineheight = 1.5
  ) +
  xlim(0, 0.01) +
  theme_void()

## End(Not run)
formatNum(1.23456, decimals = 3)
formatNum(1.23456, type = 1)
formatNum(1.23456, pre = "$", pos = "/person")
formatNum(123456, abbr = TRUE)
formatNum(c(123123, 123.123, 0.123123), signif = 2)
formatNum(1234567890, abbr = TRUE, signif = 3)
formatNum(1234567890, decimals = 0, abbr = TRUE)
formatNum(c(-3:3), sign = TRUE)
```

Description

This function lets the user group, count, calculate percentages and cumulatives. It also plots results if needed. Tidyverse friendly.

Usage

```
freqs(  
  df,  
  ...,  
  wt = NULL,  
  rel = FALSE,  
  results = TRUE,  
  variable_name = NA,  
  plot = FALSE,  
  rm.na = FALSE,  
  title = NA,  
  subtitle = NA,  
  top = 20,  
  abc = FALSE,  
  save = FALSE,  
  subdir = NA,  
  quiet = FALSE  
)
```

Arguments

df	Data.frame
...	Variables. Variables you wish to process. Order matters. If no variables are passed, the whole data.frame will be considered
wt	Variable, numeric. Weights.
rel	Boolean. Relative percentages (or absolute)?
results	Boolean. Return results in a datafram?
variable_name	Character. Overwrite the main variable's name
plot	Boolean. Do you want to see a plot? Three variables tops.
rm.na	Boolean. Remove NA values in the plot? (not filtered for numerical output; use na.omit() or filter() if needed)
title	Character. Overwrite plot's title with.
subtitle	Character. Overwrite plot's subtitle with.
top	Integer. Filter and plot the most n frequent for categorical values. Set to NA to return all values
abc	Boolean. Do you wish to sort by alphabetical order?
save	Boolean. Save the output plot in our working directory
subdir	Character. Into which subdirectory do you wish to save the plot to?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

Plot when `plot=TRUE` and data.frame with grouped frequency results when `plot=FALSE`.

See Also

Other Frequency: [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#)

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Other Visualization: [distr\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

# How many survived?
dft %>% freqs(Survived)

# How many survived per Class?
dft %>% freqs(Pclass, Survived, abc = TRUE)

# How many survived per Class with relative percentages?
dft %>% freqs(Pclass, Survived, abc = TRUE, rel = TRUE)

# Using a weighted feature
dft %>% freqs(Pclass, Survived, wt = Fare / 100)

### Let's check the results with plots:

# How many survived and see plot?
dft %>% freqs(Survived, plot = TRUE)

# How many survived per class?
dft %>% freqs(Survived, Pclass, plot = TRUE)

# Per class, how many survived?
dft %>% freqs(Pclass, Survived, plot = TRUE)

# Per sex and class, how many survived?
dft %>% freqs(Sex, Pclass, Survived, plot = TRUE)

# Frequency of tickets + Survived
dft %>% freqs(Survived, Ticket, plot = TRUE)

# Frequency of tickets: top 10 only and order them alphabetically
dft %>% freqs(Ticket, plot = TRUE, top = 10, abc = TRUE)
```

Description

This function lets the user analize data by visualizing the frequency of each value of each column from a whole data frame.

Usage

```
freqs_df(
  df,
  max = 0.9,
  min = 0,
  novar = TRUE,
  plot = FALSE,
  top = 30,
  quiet = FALSE,
  save = FALSE,
  subdir = NA
)
```

Arguments

df	Data.frame
max	Numeric. Top variance threshold. Range: (0-1]. These variables will be excluded
min	Numeric. Minimum variance threshold. Range: [0-1). These values will be grouped into a high frequency (HF) value
novar	Boolean. Remove no variance columns?
plot	Boolean. Do you want to see a plot? Three variables tops
top	Integer. Plot most relevant (less categories) variables
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
save	Boolean. Save the output plot in our working directory
subdir	Character. Into which subdirectory do you wish to save the plot to?

Value

Plot when `plot=TRUE` and data.frame with grouped frequency results when `plot=FALSE`.

See Also

Other Frequency: [freqs\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#)

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```
data(dft) # Titanic dataset
freqs_df(dft)
freqs_df(dft, plot = TRUE)
```

freqs_list

Frequencies on Lists and UpSet Plot

Description

Visualize frequency of elements on a list, list vector, or vector with comma separated values. Detect which combinations and elements are the most frequent and how much they represent of your total observations. This is similar to the [UpSet Plots](#) which may be used as an alternative to Venn diagrams.

Usage

```
freqs_list(
  df,
  var = NULL,
  wt = NULL,
  fx = "mean",
  rm.na = FALSE,
  min_elements = 1,
  limit = 10,
  limit_x = NA,
  limit_y = NA,
  tail = TRUE,
  size = 10,
  unique = TRUE,
  abc = FALSE,
  title = "",
  plot = TRUE
)
```

Arguments

<code>df</code>	Data.frame
<code>var</code>	Variable. Variables you wish to process.
<code>wt</code>	Variable, numeric. Select a numeric column to use in the colour scale, used as sum, mean... of those values for each of the combinations.
<code>fx</code>	Character. Set operation: mean, sum
<code>rm.na</code>	Boolean. Remove NA value from <code>wt</code> ?
<code>min_elements</code>	Integer. Exclude combinations with less than n elements

limit, limit_x, limit_y	Integer. Show top n combinations (x) and/or elements (y). The rest will be grouped into a single element. Set argument to 0 to ignore. limit_x/limit_y answer to limit's argument.
tail	Boolean. Show tail grouped into "..." on the plots?
size	Numeric. Text base size
unique	Boolean. a,b = b,a?
abc	Boolean. Do you wish to sort by alphabetical order?
title	Character. Overwrite plot's title with.
plot	Boolean. Plot viz? Will be generated anyways in the output object

Value

List. data.frame with the data results, elements and combinations.

See Also

Other Frequency: [freqs\(\)](#), [freqs_df\(\)](#), [freqs_plot\(\)](#)

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```

## Not run:
df <- dplyr::starwars
head(df[, c(1, 4, 5, 12)], 10)

# Characters per movies combinations in a list column
head(df$films, 2)
freqs_list(df, films)

# Skin colours in a comma-separated column
head(df$skin_color)
x <- freqs_list(df, skin_color, min_elements = 2, limit = 5, plot = FALSE)
# Inside "x" we'll have:
names(x)

# Using the 'wt' argument to add a continuous value metric
# into an already one-hot encoded columns dataset (and hide tail)
csv <- "https://raw.githubusercontent.com/hms-dbmi/UpSetR/master/inst/extdata/movies.csv"
movies <- read.csv(csv, sep = ";")
head(movies)
freqs_list(movies,
           wt = AvgRating, min_elements = 2, tail = FALSE,
           title = "Movies\nMixed Genres\nRanking")
)
# So, please: no more Comedy+SciFi and more Drama+Horror films (based on ~50 movies)!
```

```
## End(Not run)
```

freqs_plot*Combined Frequencies Plot for Categorical Features***Description**

Plot frequencies of multiple categories within a data.frame in a new fancy way. Tidyverse friendly, based on `lares::freqs()`, no limits on amount of features to evaluate.

Usage

```
freqs_plot(
  df,
  ...,
  top = 10,
  rm.na = FALSE,
  abc = FALSE,
  title = NA,
  subtitle = NA,
  quiet = FALSE
)
```

Arguments

<code>df</code>	Data.frame
<code>...</code>	Variables. Variables you wish to process. Order matters. If no variables are passed, the whole data.frame will be considered
<code>top</code>	Integer. Filter and plot the most n frequent for categorical values. Set to NA to return all values
<code>rm.na</code>	Boolean. Remove NA values in the plot? (not filtered for numerical output; use <code>na.omit()</code> or <code>filter()</code> if needed)
<code>abc</code>	Boolean. Do you wish to sort by alphabetical order?
<code>title</code>	Character. Overwrite plot's title with.
<code>subtitle</code>	Character. Overwrite plot's subtitle with.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

Plot. Result of the frequency of combined variables.

See Also

Other Frequency: [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#)

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

x <- freqs_plot(dft, Pclass, Survived)
x$data
plot(x)

freqs_plot(dft, Pclass, Survived, Sex, Embarked)

freqs_plot(dft, Pclass, Survived, Sex, Embarked, top = 15)
```

gain_lift*Cumulative Gain, Lift and Response***Description**

This function calculates cumulative gain, lift, and response values for a predictive score of a specific target. You can use the [mplot_gain\(\)](#) function to create a plot.

Usage

```
gain_lift(
  tag,
  score,
  target = "auto",
  splits = 10,
  plot = FALSE,
  quiet = FALSE
)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result
target	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only used when binary categorical model.

splits	Integer. Number of percentiles to split the data
plot	Boolean. Plot results? Uses <code>mplot_gain()</code>
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

`data.frame` when `plot=FALSE` or `plot` when `plot=TRUE`.

See Also

Other Machine Learning: `ROC()`, `conf_mat()`, `export_results()`, `h2o_automl()`, `h2o_predict_MOJO()`, `h2o_selectmodel()`, `impute()`, `iter_seeds()`, `lasso_vars()`, `model_metrics()`, `model_preprocess()`, `msplit()`

Other Model metrics: `ROC()`, `conf_mat()`, `errors()`, `loglossBinary()`, `model_metrics()`

Examples

```
data(dfr) # Results for AutoML Predictions
head(dfr$class2)

# Results for Binomial Model
gain_lift(dfr$class2$tag, dfr$class2$scores, target = "FALSE")
gain_lift(dfr$class2$tag, dfr$class2$scores, target = "TRUE", splits = 5)
```

Description

This function lets the user interact with Google's Gemini LLM Model using its API, and returns the rendered reply.

Usage

```
gemini_ask(
  ask,
  secret_key = get_creds("gemini")$api_key,
  url = Sys.getenv("LARES_GEMINI_API"),
  temperature = 0.5,
  max_tokens = 1024,
  quiet = FALSE,
  ...
)
gemini_image(
  ask,
  image,
```

```

secret_key = get_creds("gemini")$api_key,
url = Sys.getenv("LARES_GEMINI_API"),
temperature = 0.5,
max_tokens = 1024,
quiet = FALSE,
...
)

```

Arguments

ask	Character. Redacted prompt to ask. If multiple asks are requested, they will be concatenated with "+" into a single request.
secret_key	Character. Secret Key. Get yours in: platform.openai.com for OpenAI or makersuite.google.com/app/apikey for Gemini.
url	Character. Base API URL.
temperature	Numeric. The temperature to use for generating the response. Default is 0.5. The lower the temperature, the more deterministic the results in the sense that the highest probable next token is always picked. Increasing temperature could lead to more randomness, which encourages more diverse or creative outputs. You are essentially increasing the weights of the other possible tokens. In terms of application, you might want to use a lower temperature value for tasks like fact-based QA to encourage more factual and concise responses. For poem generation or other creative tasks, it might be beneficial to increase the temperature value.
max_tokens	Integer. The maximum number of tokens in the response.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.
image	Character. Data to be encoded/decoded. It can be a raw vector, text connection or file name.

Value

(Invisible) list. Content returned from API POST and processed.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gpt_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)
 Other LLM: [gpt_ask\(\)](#), [gpt_prompt\(\)](#)

Examples

```

## Not run:
api_key <- get_credentials()$gemini$api_key
# Open question:
gemini_ask("Can you write an R function to plot a dummy histogram?", api_key)
# Image question

```

```

image <- "man/figures/automl_map.png"
gemini_image("Can you explain this flow with more detail?", image, api_key)

## End(Not run)

```

get_credentials *Load Credentials from a YML File*

Description

Load credentials from a local YML file. You can set your `.Renviron` and the `LARES_CREDS` parameter to remember (forever) the directory of your credentials' file. To use it later, you may leave `dir = NA`. You may also use this function for external (non-lares) code/use.

Usage

```

get_credentials(
  from = NA,
  dir = NA,
  filename = "config.yml",
  env = "LARES_CREDS",
  ...
)

get_creds(
  from = NA,
  dir = NA,
  filename = "config.yml",
  env = "LARES_CREDS",
  ...
)

```

Arguments

<code>from</code>	Character. Family of values to import from the YML file. If you don't know these names, set <code>from = NA</code> and a warning will display all possible values, depending on your YML file.
<code>dir</code>	Character. Credentials directory where your YML file is. If used frequently, set your directory by using the <code>.Renviron</code> file. To do so, leave <code>dir</code> as <code>NA</code> and follow the steps. If <code>dir</code> is a list, it'll return <code>dir</code> (manual credentials input).
<code>filename</code>	Character. YML filename with your credentials.
<code>env</code>	Character. Environment variable name. No need to set differently for any function that uses this library. Only for external use.
<code>...</code>	Additional parameters.

Value

List. Result of reading your credential's YML file, filtered by your from input if provided.

Set the default directory

The first time you use any function that has the creds parameter, if the dir parameter is set to NA, this function will ask you to set the directory where you save your YML local file with your credentials. This will be asked once and will be set for further R sessions. Remember to reset your session for this setup to start working properly.

YML file format

A YML file is a text file, with .yml file format. You may start from the dummy YML file shared which shows the structure you must follow to set your credentials file. Check it out [here](#) or find it locally using system.file("docs", "config.yml", package = "lares").

See Also

Other Credentials: [db_download\(\)](#), [db_upload\(\)](#), [encrypt_file\(\)](#), [get_tweets\(\)](#), [mail_send\(\)](#), [queryDB\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#), [stocks_file\(\)](#)

Examples

```
## Not run:  
# Load dummy config.yml file from the library  
# Recommendation: set dir with NA (read documentation)  
# We need the directory, not the file  
yml <- dirname(system.file("docs", "config.yml", package = "lares"))  
  
# Let's see which credentials we have in our file  
get_credentials(dir = yml)  
# Warning message: No credentials for NA found in your YML file.  
# Try any of the following: 'service1', 'service2', 'service3'  
  
# Get credentials for service2  
get_credentials("service2", dir = yml)  
  
## End(Not run)
```

Description

This function lets the user download historical currency exchange rate between two currencies.

Usage

```
get_currency(
  currency_pair,
  from = Sys.Date() - 99,
  to = Sys.Date(),
  fill = FALSE,
  ...
)
```

Arguments

<code>currency_pair</code>	Character. Which currency exchange do you wish to get the history from? i.e, USD/COP, EUR/USD...
<code>from</code>	Date. From date
<code>to</code>	Date. To date
<code>fill</code>	Boolean. Fill weekends and non-quoted dates with previous values?
...	Additional parameters.

Value

`data.frame`. Result of fetching online data for `currency_pair` grouped by date.

Examples

```
# For today (or any one single date)
get_currency("USD/COP", from = Sys.Date())
# For multiple dates
get_currency("EUR/USD", from = Sys.Date() - 7, fill = TRUE)
```

get_mp3*Download MP3 from URL***Description**

This function downloads YouTube videos or Soundcloud or any other platform supported by the youtube-dl library, and converts them into high quality MP3 files. The URL can be for a single video or a whole playlist. It also returns metadata into an (invisible) list.

Usage

```
get_mp3(
  id,
  mp3 = TRUE,
  repo = "yt-dlp",
  params = "--no-check-certificate",
```

```

    start_time = 0,
    end_time = NA,
    overwrite = TRUE,
    open = FALSE,
    delete = open,
    info = TRUE,
    cover = FALSE,
    quiet = FALSE
)

```

Arguments

id	Character. YouTube URL or ID to search for.
mp3	Boolean. Add mp3 optimal parameters?
repo	Character. Chose repository you installed youtube-dl from. Any of: "youtube-dl" (latest stable version), "yt-dlp" (latest dev version).
params	Character. Additional parameters.
start_time, end_time	Numeric. Start and end time to trim the audio output in seconds.
overwrite	Boolean. Overwrite original file?
open, delete	Boolean. After everything's done, should the file be opened? Should it be deleted?
info	Boolean. Import and return metadata?
cover	Boolean. Google Search its squared cover?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

(Invisible) list with id's meta-data.

youtube-dl

More info from the original developers and its code: [youtube-dl's Github](#)

See Also

Other Scrapper: [filesGD\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)

Other Audio: [trim_mp3\(\)](#)

Examples

```

# You must have "youtube-dl" installed in your OS:
## Not run:
# Download video from YouTube and convert to MP3
get_mp3("https://www.youtube.com/watch?v=lrlKcCdVw9Q")
# OR simply

```

```
get_mp3("lrlKcCdVw9Q")
# For dev version, use:
get_mp3("m3RX4LJh0ii", repo = "yt-dlp")

## End(Not run)
```

get_tweets*Get Tweets***Description**

This function downloads tweets with personal credentials

Usage

```
get_tweets(q, n = 10000, creds = NA)
```

Arguments

- | | |
|--------------------|--|
| <code>q</code> | Query. Check for ?rtweet::search_tweets() |
| <code>n</code> | Integer. Total of tweets to return |
| <code>creds</code> | Character. Credential's user (see <code>get_creds()</code>) |

Value

`data.frame` with API response results.

See Also

Other Credentials: [db_download\(\)](#), [db_upload\(\)](#), [encrypt_file\(\)](#), [get_credentials\(\)](#), [mail_send\(\)](#), [queryDB\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#), [stocks_file\(\)](#)

gg_fill_customs*Custom fill, colour and text colours for ggplot2***Description**

This function lets the user use pre-defined default colours. Check your `lares_pal()$labels` scale. Feel free to use `gg_vals()` to debug colours used in latest plot.

Usage

```
gg_fill_customs(column = "fill", ...)

gg_colour_customs(column = "colour", ...)

gg_text_customs(column = "colour", ...)

gg_vals(layer = "fill", column = layer, cols = NULL, ...)
```

Arguments

column	Character. Select any of "fill" or "colour" to use on your <code>lares_pal()\$labels</code> palette.
...	Allow additional parameters not used.
layer	Character. Select any of "fill", "colour", or "label" to get the layer containing the colours labels you wish to colour.
cols	Data.frame. Customize colour palette with a data.frame. Must contain values, fill, and colour columns.

Details

Notice that when the layer defined is any of GeomPoint, GeomLine, GeomText or GeomLabel, `gg_colour_customs()` will force `column = "fill"` parameter.

Value

Same as `scale_fill_manual` or `scale_colour_manual` but with custom palette applied.

See Also

Other Themes: [lares_pal\(\)](#), [plot_palette\(\)](#), [theme_lares\(\)](#)

Examples

```
library("ggplot2")
# Generic plot function to run examples to
run_plot <- function(add_fxs = TRUE) {
  p <- data.frame(station = c("spring", "summer", "fall", "winter"), num = 1:4) %>%
    ggplot(aes(x = station, y = num, fill = station)) +
    geom_col() +
    geom_text(aes(y = 0.5, label = num, colour = station), size = 6)
  if (add_fxs) p <- p + gg_fill_customs() + gg_colour_customs()
  return(p)
}
# Default colours
run_plot()
# Check last colours used
gg_vals("fill", "fill")
gg_vals("colour", "colour")
```

```

# Change any default colour
options("lares.colours.custom" = data.frame(
  values = c("summer", "winter"),
  fill = c("pink", "black"),
  colour = c("black", "white")
))
run_plot()
# Check last colours used
gg_vals("fill", "fill")
gg_vals("colour", "colour")
# Reset to default colours
options("lares.colours.custom" = NULL)
# Notice you can use 'pal = 4' argument on theme_lares() too
run_plot(add_fxs = FALSE) + theme_lares(pal = 4)

```

glued*Interpolate a string [glue wrapper]***Description**

Format and interpolate a string using a glue wrapper. Allows simple operations, `NULL` values as input, and interactions with internal (created within `glued`) and external (environment) objects.

Usage

```
glued(..., .sep = "", empty_lines = "keep", .envir = parent.frame())
```

Arguments

<code>...</code>	[expressions]
	Unnamed arguments are taken to be expression string(s) to format. Multiple inputs are concatenated together before formatting. Named arguments are taken to be temporary variables available for substitution.
<code>.sep</code>	[character(1): ""]
	Separator used to separate elements.
<code>empty_lines</code>	Character. Set to "keep" to keep or "drop" to drop empty lines.
<code>.envir</code>	[environment: parent.frame()]
	Environment to evaluate each expression in. Expressions are evaluated from left to right. If <code>.x</code> is an environment, the expressions are evaluated in that environment and <code>.envir</code> is ignored. If <code>NULL</code> is passed, it is equivalent to <code>emptyenv()</code> .

Value

Same as input but transformed (`glued`).

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
name <- "Bernardo"
age <- 29
anniversary <- as.Date("2016-04-30")
glued(
  My name is {name},
  my age next year will be {age + 1},
  and I got married on {format(anniversary, '%A, %B %d, %Y')}.")

# Single braces can be inserted by doubling them
glued("My name is {name}, not {{name}}.")

# You can also used named arguments
glued(
  "Her name is {name}, ",
  "and her age next year will be {age + 1}.",
  name = "Maru",
  age = 6
)

# And run operations with memories (beware!)
glued("My name, {name}, has {n <- nchar(name)}; n} characters.
      If we multiply by ten, we'll have {10 * n} characters!")

# If you pass a vector, the operation will be repeated for each element
glued("Here's the value #{1:3}")
```

Description

This function lets the user ask ChatGPT via its API, and returns the rendered reply. There are a couple of specific verbs (functions) with a preset prompt to help fetch the data in specific formats. We also store the prompts and replies in current session with their respective time-stamps so user can gather historical results.

Usage

```

gpt_ask(
  ask,
  secret_key = get_credentials()$openai$secret_key,
  url = Sys.getenv("LARES_GPT_URL"),
  model = Sys.getenv("LARES_GPT_MODEL"),
  num_retries = 3,
  temperature = 0.5,
  max_tokens = NULL,
  pause_base = 1,
  quiet = FALSE,
  ...
)

gpt_history(quiet = TRUE, ...)

gpt_table(x, cols = NULL, quiet = TRUE, ...)

gpt_classify(x, categories, quiet = TRUE, ...)

gpt_tag(x, tags, quiet = TRUE, ...)

gpt_extract(x, extract, quiet = TRUE, ...)

gpt_format(x, format, quiet = TRUE, ...)

gpt_convert(x, unit, quiet = TRUE, ...)

gpt_translate(x, language, quiet = TRUE, ...)

```

Arguments

<code>ask</code>	Character. Redacted prompt to ask. If multiple asks are requested, they will be concatenated with "+" into a single request.
<code>secret_key</code>	Character. Secret Key. Get yours in: platform.openai.com for OpenAI or makersuite.google.com/app/apikey for Gemini.
<code>url</code>	Character. Base API URL.
<code>model</code>	Character. OpenAI model to use. This can be adjusted according to the available models in the OpenAI API (such as "gpt-4").
<code>num_retries</code>	Integer. Number of times to retry the request in case of failure. Default is 3.
<code>temperature</code>	Numeric. The temperature to use for generating the response. Default is 0.5. The lower the temperature, the more deterministic the results in the sense that the highest probable next token is always picked. Increasing temperature could lead to more randomness, which encourages more diverse or creative outputs. You are essentially increasing the weights of the other possible tokens. In terms of application, you might want to use a lower temperature value for tasks like

fact-based QA to encourage more factual and concise responses. For poem generation or other creative tasks, it might be beneficial to increase the temperature value.

max_tokens	Integer. The maximum number of tokens in the response.
pause_base	Numeric. The number of seconds to wait between retries. Default is 1.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.
x	Vector. List items you wish to process in your instruction
cols	Vector. Force column names for your table results.
categories, tags	Vector. List of possible categories/tags to consider.
extract, format, unit	Character. Length 1 or same as x to extract/format/unit information from x. For example: email, country of phone number, country, amount as number, currency ISO code, ISO, Fahrenheit, etc.
language	Character. Language to translate to

Value

(Invisible) list. Content returned from API POST and processed.

See Also

Other API: [bring_api\(\)](#), [fb_accounts\(\)](#), [fb_ads\(\)](#), [fb_creatives\(\)](#), [fb_insights\(\)](#), [fb_process\(\)](#), [fb_report_check\(\)](#), [fb_rf\(\)](#), [fb_token\(\)](#), [gemini_ask\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Other ChatGPT: [gpt_promptr\(\)](#)

Other LLM: [gemini_ask\(\)](#), [gpt_promptr\(\)](#)

Examples

```
## Not run:
api_key <- get_credentials()$openai$secret_key
# Open question:
gpt_ask("Can you write an R function to plot a dummy histogram?", api_key)

##### The following examples return dataframes:
# Classify each element based on categories:
gpt_classify(1:10, c("odd", "even"))

# Add all tags that apply to each element based on tags:
gpt_tag(
  c("I love chocolate", "I hate chocolate", "I like Coke"),
  c("food", "positive", "negative", "beverage")
)

# Extract specific information:
gpt_extract()
```

```

c("My mail is 123@test.com", "30 Main Street, Brooklyn, NY, USA", "+82 2-312-3456", "$1.5M"),
  c("email", "full state name", "country of phone number", "amount as number")
)

# Format values
gpt_format(
  c("March 27th, 2021", "12-25-2023 3:45PM", "01.01.2000", "29 Feb 92"),
  format = "ISO Date getting rid of time stamps"
)

# Convert temperature units
gpt_convert(c("50C", "300K"), "Fahrenheit")

# Create a table with data
gpt_table("5 random people's address in South America, email, phone, age between 18-30")
gpt_table(
  ask = "5 largest cities, their countries, and population",
  cols = c("city_name", "where", "POP")
)

# Translate text to any language
gpt_translate(
  rep("I love you with all my heart", 5),
  language = c("spanish", "chinese", "japanese", "russian", "german")
)

# Now let's read the historical prompts, replies, and more from current session
gpt_history()

## End(Not run)

```

Description

Build standard prompts to get the right outcomes using the four theoretical elements required to build standard and cleaner asks: instruction, input, context, and output. Inspired by the [Prompt Engineering Guide](#) free guide. Remember to start simple and be very specific to get exactly what you need.

Usage

```

gpt_promter(
  instruction = NULL,
  input = NULL,
  context = NULL,
  output = NULL,
  quiet = TRUE,
  ...
)

```

Arguments

instruction, input, context, output	Character or vector. You do not need all the four elements for a prompt and the format depends on the task at hand.
quiet	Boolean. Should the written prompt be printed or not?
...	Additional parameters. You can pass <code>cols</code> parameter to explicitly set column names.

Value

(Invisible) list with written prompt and elements provided.

Elements of a Prompt

- Instruction** a specific task or instruction you want the model to perform. It usually starts with a verb given its an instruction.
- Input** input data, elements or question that we are interested to find a response for,
- Context** external information, additional context or references to steer the model to better responses.
- Output** type or format of the output.

See Also

- Other ChatGPT: [gpt_ask\(\)](#)
- Other LLM: [gemini_ask\(\)](#), [gpt_ask\(\)](#)

Examples

```
# A simple formatted table with data
# Note: I mostly use output = "table" and enabled an auxiliary enrichment prompt
(p <- gpt_promter(instruction = "Capitals of the world", output = "table"))

# Classify
p <- gpt_promter(
  instruction = "For each of the inputs, classify using only the options in context",
  input = c("Molecule", "Elephant", "Milky Way", "Cat", "Planet Earth"),
  context = c("Big", "Medium", "Small"),
  output = "table",
  # This cols parameter is auxiliary
  cols = c("Input", "Category"),
  quiet = FALSE
)

# Tag all categories that apply
p <- gpt_promter(
  instruction = paste(
    "For each of the inputs, provide which of the",
    "context values apply as correct tags using TRUE/FALSE"
  ),
)
```

```

input = c("I love chocolate", "I hate chocolate", "I like Coke", "Who am I?", "T-REX"),
context = c("food", "positive", "negative", "beverage"),
output = "table",
quiet = FALSE
)

# Extract information from strings
p <- gpt_promptr(
  instruction = "For each of the inputs, extract each of the information asked in context",
  input = c("My mail is 123@test.com", "30 Main St, NY, USA", "+82 2-312-3456", "$1.5M"),
  context = c("email", "full state name", "country of phone", "full non-abbreviated number"),
  output = "table",
  cols = c("Input", "Element_to_extract", "Value"),
  quiet = FALSE
)

# Translate to several languages
p <- gpt_promptr(
  instruction = "For each of the inputs, translate to the respective languages in context",
  input = rep("I love you with all my heart", 5),
  context = c("spanish", "chinese", "japanese", "russian", "german"),
  output = "table",
  cols = c("Input", "Language", "Translation"),
  quiet = FALSE
)

# Format date values
p <- gpt_promptr(
  instruction = paste(
    "For each of the inputs,",
    "standardize and format all values to the format in context"
  ),
  input = c("March 27th, 2021", "12-25-2023 3:45PM", "01.01.2000", "29 Feb 92"),
  context = "ISO Date getting rid of time stamps",
  output = "table",
  cols = c("Input", "Formatted"),
  quiet = FALSE
)

# Convert units
p <- gpt_promptr(
  instruction = paste(
    "For each of the inputs,",
    "provide new converted values using the units in context"
  ),
  input = c("50C", "300K", "100F", "0F", "32C", "0K"),
  context = "Fahrenheit",
  output = "table",
  cols = c("Input", "Original_Unit", "Total_Value", "Converted_Value", "New_Unit"),
  quiet = FALSE
)

# Read a text and answer a question related to it

```

```
gpt_promptr(  
  instruction = "read",  
  context = "Long text here",  
  input = "Question here"  
)$prompt
```

grep1_letters*Pattern Matching for Letters considering Blanks*

Description

Match pattern of letters considering blanks within each element of a character vector, allowing counted characters between and around each letter. Used as an auxiliary function for the Scrabble family of functions.

Usage

```
grep1_letters(x, pattern, blank = "_")
```

Arguments

x	Character vector
pattern	Character. Character string containing a semi-regular expression which uses the following logic: "a_b" means any character that contains "a" followed by something followed by "b", anywhere in the string.
blank	Character. String to use between letters.

Value

Boolean check for each value on x.

Examples

```
x <- c("aaaa", "bbbb", "baba", "aabb", "a", "ab")  
grep1_letters(x, "ab")  
grep1_letters(x, "_ab")  
grep1_letters(x, "a_a")  
grep1_letters(x, "c")
```

grepml*Pattern Matching for Any or All Multiple Matches*

Description

This function returns a boolean vector of the same length as ‘x’, each element of which is the result of applying the ‘type’ of matches to the corresponding element of ‘x’, using regular expressions.

Usage

```
grepml(pattern, x, type = "all", ...)
```

Arguments

pattern	character string containing a regular expression (or character string for fixed = TRUE) to be matched in the given character vector. Coerced by as.character to a character string if possible. If a character vector of length 2 or more is supplied, the first element is used with a warning. Missing values are allowed except for <code>regexpr</code> , <code>gregexpr</code> and <code>regexec</code> .
x	Character vector. Text where matches are sought, or an object which can be coerced by <code>as.character</code> to a character vector. Long vectors are supported.
type	Character. Type of match. Choose one of: <code>any</code> , <code>all</code>
...	Additional arguments to pass to <code>grep</code>

Value

Boolean of same length as x

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Examples

```
x <- c(123, 876, 18761)
patterns <- c(1, 2)
grepml(patterns, x, type = "any")
grepml(patterns, x, type = "all")
```

gtrends_related *Google Trends: Related and Time Plots*

Description

This function creates a plot with Google Trend's related topics and queries, and let the user compare different keywords.

This function creates a plot with google trend's data on timelines and let the user compare different keywords.

Usage

```
gtrends_related(gtrend, top = NA, title = NA, note = NA, exclude = NULL)  
gtrends_time(gtrend, title = NA)
```

Arguments

gtrend	List. Result from gtrendsR::gtrends(keyword, geo, time)
top	Integer. Filter top n results only.
title	Character. Custom title for the plot.
note	Character. Add a note to the plot if needed.
exclude	Character vector. Which observations do you wish to exclude?

Value

plot for Google Trend's results input gtrend.

Plot for Google Trend's results input gtrend.

See Also

Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)

Other Google: [filesGD\(\)](#), [queryGA\(\)](#), [readGS\(\)](#)

`h2o_automl`*Automated H2O's AutoML*

Description

This function lets the user create a robust and fast model, using H2O's AutoML function. The result is a list with the best model, its parameters, datasets, performance metrics, variables importance, and plots. Read more about the `h2o_automl()` pipeline [here](#).

Usage

```
h2o_automl(  
  df,  
  y = "tag",  
  ignore = NULL,  
  train_test = NA,  
  split = 0.7,  
  weight = NULL,  
  target = "auto",  
  balance = FALSE,  
  impute = FALSE,  
  no_outliers = TRUE,  
  unique_train = TRUE,  
  center = FALSE,  
  scale = FALSE,  
  thresh = 10,  
  seed = 0,  
  nfolds = 5,  
  max_models = 3,  
  max_time = 10 * 60,  
  start_clean = FALSE,  
  exclude_algos = c("StackedEnsemble", "DeepLearning"),  
  include_algos = NULL,  
  plots = TRUE,  
  alarm = TRUE,  
  quiet = FALSE,  
  print = TRUE,  
  save = FALSE,  
  subdir = NA,  
  project = "AutoML Results",  
  verbosity = NULL,  
  ...  
)  
  
## S3 method for class 'h2o_automl'  
plot(x, ...)
```

```
## S3 method for class 'h2o_automl'
print(x, importance = TRUE, ...)
```

Arguments

df	Dataframe. Dataframe containing all your data, including the dependent variable labeled as 'tag'. If you want to define which variable should be used instead, use the y parameter.
y	Variable or Character. Name of the dependent variable or response.
ignore	Character vector. Force columns for the model to ignore
train_test	Character. If needed, df's column name with 'test' and 'train' values to split
split	Numeric. Value between 0 and 1 to split as train/test datasets. Value is for training set. Set value to 1 to train with all available data and test with same data (cross-validation will still be used when training). If train_test is set, value will be overwritten with its real split rate.
weight	Column with observation weights. Giving some observation a weight of zero is equivalent to excluding it from the dataset; giving an observation a relative weight of 2 is equivalent to repeating that row twice. Negative weights are not allowed.
target	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only used when binary categorical model.
balance	Boolean. Auto-balance train dataset with under-sampling?
impute	Boolean. Fill NA values with MICE?
no_outliers	Boolean/Numeric. Remove y's outliers from the dataset? Will remove those values that are farther than n standard deviations from the dependent variable's mean (Z-score). Set to TRUE for default (3) or numeric to set a different multiplier.
unique_train	Boolean. Keep only unique row observations for training data?
center, scale	Boolean. Using the base function scale, do you wish to center and/or scale all numerical values?
thresh	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
seed	Integer. Set a seed for reproducibility. AutoML can only guarantee reproducibility if max_models is used because max_time is resource limited.
nfolds	Number of folds for k-fold cross-validation. Must be >= 2; defaults to 5. Use 0 to disable cross-validation; this will also disable Stacked Ensemble (thus decreasing the overall model performance).
max_models, max_time	Numeric. Max number of models and seconds you wish for the function to iterate. Note that max_models guarantees reproducibility and max_time not (because it depends entirely on your machine's computational characteristics)

<code>start_clean</code>	Boolean. Erase everything in the current h2o instance before we start to train models? You may want to keep other models or not. To group results into a custom common AutoML project, you may use <code>project_name</code> argument.
<code>exclude_algos, include_algos</code>	Vector of character strings. Algorithms to skip or include during the model-building phase. Set NULL to ignore. When both are defined, only <code>include_algos</code> will be valid.
<code>plots</code>	Boolean. Create plots objects?
<code>alarm</code>	Boolean. Ping (sound) when done. Requires beep.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>print</code>	Boolean. Print summary when process ends?
<code>save</code>	Boolean. Do you wish to save/export results into your working directory?
<code>subdir</code>	Character. In which directory do you wish to save the results? Working directory as default.
<code>project</code>	Character. Your project's name
<code>verbosity</code>	Verbosity of the backend messages printed during training; Optional. Must be one of NULL (live log disabled), "debug", "info", "warn", "error". Defaults to "warn".
<code>...</code>	Additional parameters on <code>h2o::h2o.automl</code>
<code>x</code>	<code>h2o_automl</code> object
<code>importance</code>	Boolean. Print important variables?

Value

List. Trained model, predicted scores and datasets used, performance metrics, parameters, importance data.frame, seed, and plots when `plots=TRUE`.

List of algorithms

[-> Read more here](#)

DRF Distributed Random Forest, including Random Forest (RF) and Extremely-Randomized Trees (XRT)

GLM Generalized Linear Model

XGBoost eXtreme Grading Boosting

GBM Gradient Boosting Machine

DeepLearning Fully-connected multi-layer artificial neural network

StackedEnsemble Stacked Ensemble

Methods

print Use `print` method to print models stats and summary

plot Use `plot` method to plot results using `mplot_full()`

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

Examples

```
## Not run:
# CRAN
data(dft) # Titanic dataset
dft <- subset(dft, select = -c(Ticket, PassengerId, Cabin))

# Classification: Binomial - 2 Classes
r <- h2o_automl(dft, y = Survived, max_models = 1, impute = FALSE, target = "TRUE", alarm = FALSE)

# Let's see all the stuff we have inside:
lapply(r, names)

# Classification: Multi-Categorical - 3 Classes
r <- h2o_automl(dft, Pclass, ignore = c("Fare", "Cabin"), max_time = 30, plots = FALSE)

# Regression: Continuous Values
r <- h2o_automl(dft, y = "Fare", ignore = c("Pclass"), exclude_algos = NULL, quiet = TRUE)
print(r)

# WITH PRE-DEFINED TRAIN/TEST DATAFRAMES
splits <- msplit(dft, size = 0.8)
splits$train$split <- "train"
splits$test$split <- "test"
df <- rbind(splits$train, splits$test)
r <- h2o_automl(df, "Survived", max_models = 1, train_test = "split")

## End(Not run)
```

Description

DALEX helper function to create an explainer object using a h2o trained model.

Usage

```
h2o_explainer(df, model, y = "tag", ignore = NULL, ...)
```

Arguments

df	Dataframe. Must contain all columns and predictions
model	Model object (H2O)
y	Character or Variable name. Variable's column name.
ignore	Character vector. Which columns should be ignored?
...	Additional parameters to pass to h2o_predict_model or h2o_predict_MOJO.

Value

List; explainer. Containing the model, data, y, predict_function, y_hat, residuals, class, label, model_info, residual_function, and weights.

See Also

Other Interpretability: [dalex_local\(\)](#), [dalex_residuals\(\)](#), [dalex_variable\(\)](#)

Examples

```
# You must have "DALEX" library to use this auxiliary function:
## Not run:
data(dft) # Titanic dataset

# TRAIN A SIMPLE MODEL
dfm <- h2o_automl(dft,
  y = "Survived",
  ignore = c("Ticket", "PassengerId", "Cabin"),
  max_models = 1
)

# EXPLAINER
explainer <- h2o_explainer(df = dfm$datasets$test, model = dfm$model, y = "Survived")
explainer$data <- na.omit(explainer$data)

# CATEGORICAL EXAMPLE
class <- dalex_variable(explainer, vars = c("Pclass", "Sex"))
class$plot

# NUMERICAL EXAMPLE
num <- dalex_variable(explainer, vars = c("Fare", "Age"))
num$plot

# LOCAL EXAMPLE
local <- dalex_local(explainer, row = 1)
# OR YOU COULD MANUALLY INPUT THE OBSERVATION
local <- dalex_local(explainer, observation = explainer$data[1, ])
local$plot

# xai2shiny's UI (needs to be installed from ModelOriented/xai2shiny)
xai2shiny(explainer, run = TRUE)

## End(Not run)
```

h2o_predict_MOJO*Calculate predictions of h2o Models*

Description

h2o_predict_MOJO lets the user predict using the h2o .zip file containing the MOJO files. Note that it works with the files generated when using the function export_results()

h2o_predict_binary lets the user predict using the h2o binary file. Note that it works with the files generated when using the function export_results(). Recommendation: use the h2o_predict_MOJO() function when possible - it let's you change h2o's version without problem.

h2o_predict_model lets the user get scores from a H2O Model Object.

h2o_predict_API lets the user get the score from an API service

Usage

```
h2o_predict_MOJO(df, model_path, method = "mojo", batch = 300)

h2o_predict_binary(df, model_path, sample = NA)

h2o_predict_model(df, model)

h2o_predict_API(df, api, exclude = "tag")
```

Arguments

df	Dataframe/Vector. Data to insert into the model.
model_path	Character. Relative model path directory or zip file.
method	Character. One of "mojo" or "json".
batch	Integer. Run n batches at a time for "json" method.
sample	Integer. How many rows should the function predict?
model	h2o model Object
api	Character. API URL.
exclude	Character. Name of the variables to exclude.

Value

data.frame with predicted results.

vector with predicted results.

data.frame with predicted results.

vector with predicted results.

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

h2o_results*Automated H2O's AutoML Results***Description**

This is an auxiliary function to calculate predictions and results when using the `h2o_automl()` function.

Usage

```
h2o_results(
  h2o_object,
  test,
  train,
  y = "tag",
  which = 1,
  model_type,
  target = "auto",
  split = 0.7,
  ignore = NULL,
  quiet = FALSE,
  project = "ML Project",
  seed = 0,
  leaderboard = list(),
  plots = TRUE,
  ...
)
```

Arguments

<code>h2o_object</code>	H2O Leaderboard (H2OFrame/H2OAutoML) or Model (h2o)
<code>test, train</code>	Dataframe. Must have the same columns
<code>y</code>	Variable or Character. Name of the dependent variable or response.
<code>which</code>	Integer. Which model to select from leaderboard
<code>model_type</code>	Character. Select "Classification" or "Regression"
<code>target</code>	Value. Which is your target positive value? If set to 'auto', the target with largest <code>mean(score)</code> will be selected. Change the value to overwrite. Only used when binary categorical model.

split	Numeric. Value between 0 and 1 to split as train/test datasets. Value is for training set. Set value to 1 to train with all available data and test with same data (cross-validation will still be used when training). If <code>train_test</code> is set, value will be overwritten with its real split rate.
ignore	Character vector. Columns too ignore
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
project	Character. Your project's name
seed	Integer. Set a seed for reproducibility. AutoML can only guarantee reproducibility if <code>max_models</code> is used because <code>max_time</code> is resource limited.
leaderboard	H2O's Leaderboard. Passed when using <code>h2o_selectmodel</code> as it contains plain model and no leader board.
plots	Boolean. Create plots objects?
...	Additional parameters on <code>h2o::h2o.automl</code>

Value

List. Trained model, predicted scores and datasets used, performance metrics, parameters, importance data.frame, seed, and plots when `plots=TRUE`.

h2o_selectmodel*Select Model from h2o_automl's Leaderboard***Description**

Select which model from the `h2o_automl` function to use

Usage

```
h2o_selectmodel(results, which_model = 1, quiet = FALSE, ...)
```

Arguments

results	<code>h2o_automl()</code> object.
which_model	Integer. Which model from the leaderboard you wish to use?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters on <code>h2o::h2o.automl</code>

Value

H2O processed model

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read_file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

h2o_shap

*SHAP values for H2O Models***Description**

SHAP (SHapley Additive exPlanations) by Lundberg and Lee (2016) is a method to explain individual predictions. SHAP is based on the game theoretically optimal Shapley Values. Calculate SHAP values for h2o models in which each row is an observation and each column a feature. Use `plot` method to visualize features importance and distributions.

Usage

```
h2o_shap(model, test = "auto", scores = "auto", y = "y", ...)
## S3 method for class 'h2o_shap'
plot(x, relevant = TRUE, top = 15, quiet = FALSE, ...)
```

Arguments

<code>model</code>	<code>h2o_automl</code> object or <code>h2o</code> model.
<code>test</code>	String or Dataframe. Leave "auto" to use <code>h2o_automl</code> 's test dataset or pass a valid dataframe.
<code>scores</code>	Numeric vector. If <code>test != "auto"</code> , you must provide predicted values
<code>y</code>	Character. If <code>test != "auto"</code> , you must provide <code>y</code> variable's name
<code>...</code>	Additional argument for <code>predict_contributions.H2OModel</code>
<code>x</code>	<code>h2o_shap</code> object
<code>relevant</code>	Boolean. Keep only relevant non-trivial (>0) features
<code>top</code>	Integer. Plot only top n values (as in importance)
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

H2OFrame with shap values for every observation and feature.

See Also

Other SHAP: [shap_var\(\)](#)

Examples

```
## Not run:
# Train a h2o_automl model
model <- h2o_automl(dft, Survived,
  max_models = 1, target = TRUE,
  ignore = c("Ticket", "Cabin", "PassengerId"),
  quiet = TRUE
)

# Calculate SHAP values
SHAP_values <- h2o_shap(model)
# Equivalent to:
# SHAP_values <- h2o_shap(
#   model = model$model,
#   test = model$datasets$test,
#   scores = model$scores_test$scores)

# Check SHAP results
head(SHAP_values)

# You must have "ggbeeswarm" library to use this auxiliary function:
# Plot SHAP values (feature importance)
plot(SHAP_values)

# Plot some of the variables (categorical)
shap_var(SHAP_values, Pclass)

# Plot some of the variables (numerical)
shap_var(SHAP_values, Fare)

## End(Not run)
```

haveInternet

*Internet Connection Check***Description**

This function checks if your R session currently have Wifi or Internet connection.

Usage

```
haveInternet(thresh = 3, url = "http://www.google.com")
```

Arguments

thresh	Numeric. How many seconds to consider a slow connection?
url	Character. URL to test the readLines 1 command

Value

Boolean. Result of checking if device has internet connection.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

holidays

Holidays in your Country

Description

This function lets the user automatically scrap holiday dates from any country and year within +- 5 years. Thanks to timeanddate.com!

Usage

```
holidays(
  countries = "Venezuela",
  years = year(Sys.Date()),
  quiet = FALSE,
  include_regions = FALSE
)
```

Arguments

<code>countries</code>	Character or vector. For which country(ies) should the holidays be imported?
<code>years</code>	Character or vector. For which year(s) do you wish to import holiday dates?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>include_regions</code>	Boolean. Default FALSE. If TRUE, for countries with internal subdivisions, it will provide details on which sub-state the found holidays apply.

Value

`data.frame` with holidays data for given countries and years.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zeroVar\(\)](#)
 Other Feature Engineering: [date_feats\(\)](#), [ohse\(\)](#)
 Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [gtrends_related\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)
 Other One Hot Encoding: [date_feats\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#)

Examples

```
holidays(countries = "Argentina")
year <- as.integer(format(Sys.Date(), format = "%Y"))
holidays(countries = c("Spain", "Venezuela"), years = year)
holidays(countries = "Germany", include_regions = TRUE)
```

image_metadata*Get Meta Data from Image Files***Description**

This function lets the user get meta data from image files or directory.

Usage

```
image_metadata(files)
```

Arguments

files	Character vector. Files or directory which contains files.
-------	--

Value

data.frame with meta-data for each image file.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read_file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

importxlsx*Import Excel File with All Its Tabs***Description**

This function lets the user import an Excel file's tabs into a list

Usage

```
importxlsx(file)
```

Arguments

file	String. Local Excel file name
-------------	-------------------------------

Value

List or data.frame. If single tab is found, a data.frame; if multiple tabs are found on file, a list of data.frames.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

impute*Impute Missing Values (using MICE)***Description**

This function uses the MICE methodology to impute missing values.

Usage

```
impute(df, m = 5, iters = 5, seed = 0, quiet = FALSE)
```

Arguments

df	Dataframe. Dataframe to transform.
m	Integer. Number of multiple imputations.
iters	Integer. Number of iterations.
seed	Integer. Set a seed for reproducibility.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

data.frame with imputed values.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

Other Missing Values: [missingness\(\)](#)

install_recommended *Install/Update Additional Recommended Libraries*

Description

All needed libraries to use (most) lares are already a dependency. There are some functions that many people won't even know exist that will require other additional libraries. Also, this may be used as a Docker way of installing useful libraries on a new instance.

Usage

```
install_recommended(progress = TRUE, all = FALSE)
```

Arguments

progress	Boolean. Show status bar?
all	Boolean. All packages? If not, only the ones not installed yet.

ip_data *Scrap data based on IP address*

Description

This function lets the user scrap <https://db-ip.com/> given IP address(es) to get their associated address type, ASN, ISP, organization, country, state or region, county, city, ZIP postal code, weather station, coordinates, Timezone, local time, languages, and currency.

Usage

```
ip_data(ip = myip(), quiet = FALSE)
```

Arguments

- ip** Vector. Vector with all IP's we wish to search.
quiet Boolean. Do not show the loading statusbar?

Value

data.frame. Each row is an unique ip address, and columns will bee created for all the additional information found.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)
 Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)

Examples

```
ip_data("163.114.132.0")
ip_data(ip = c(myip(), "201.244.197.199"), quiet = TRUE)
```

Description

This functions lets the user iterate and search for best seed. Note that if the results change a lot, you are having a high variance in your data.

Usage

```
iter_seeds(df, y, tries = 10, ...)
```

Arguments

- df** Dataframe. Dataframe containing all your data, including the dependent variable labeled as 'tag'. If you want to define which variable should be used instead, use the y parameter.
y Variable or Character. Name of the dependent variable or response.
tries Integer. Number of iterations
... Additional arguments passed to [h2o_automl](#)

Value

data.frame with performance results by seed tried on every row.

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#), [msplit\(\)](#)

json2vector

Convert Python JSON string to R vector (data.frame with 1 row)

Description

This function lets the user transform a JSON string into vector (data.frame with 1 row). You can also pass a Python's dictionary. For any other JSON transformation, jsonlite is recommended.

Usage

```
json2vector(json)
```

Arguments

json Character. JSON string.

Value

List, data.frame, or vector. Depends on the json string.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read_file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
json2vector('{"id": 1, "nodata": null, "gender": "M"}')
```

lares

*Analytics, Data Mining & Machine Learning Sidekick***Description**

R library for better/faster analytics, visualization, data mining, and machine learning tasks.

Author(s)

Bernardo Lares (laresbernardo@gmail.com)

See Also

Useful links:

- <https://github.com/laresbernardo/lares>
- <https://laresbernardo.github.io/lares/>
- Report bugs at <https://github.com/laresbernardo/lares/issues>

lares-exports

*Pipe operator***Description**

Pipe operator

lares_logo

*Print lares R library Logo***Description**

Used "ASCII Art Generator" from manytools.org to convert logo to ASCII. [Visit](#).

Usage

```
lares_logo(version = TRUE)
```

Arguments

version Boolean. Include R and lares version?

Examples

```
lares_logo()
```

lares_pal*Personal Colours Palette*

Description

Fetch customizable palettes for the library's usage. The package has its own default colour-blind friendly colours but can be customized using R internal options (i.e. `options("lares.palette" = c("#FF8303" = "#000", "#40A4D8" = "#FFF", ...))`). There are 3 options you can use to customize all colour pallettes: "lares.palette" (vector, will be used in the same order as passed, and must have a counter colour defined), "lares.colours" (vector, simple colour names and their HEX codes), and "lares.colours.custom" (data.frame, containing "values" to use dynamically, "fill" for main colour, and "colour" (not obligatory) for counter colour).

Usage

```
lares_pal(return = "list")
```

Arguments

<code>return</code>	Character. Get only what you need. Select any of: "all" or "list" (list), "colors" or "colours" (vector), "pal" or "palette" (named vector), "simple" (named vector), "custom" or "personal" (data.frame)
---------------------	---

Value

Depending on the `return` input, we get a:

- vector with palette results vector
- vector with palette results vector's names
- list with palette results vector, labels results data.frame, and simple results named vector

See Also

Other Themes: [gg_fill_customs\(\)](#), [plot_palette\(\)](#), [theme_lares\(\)](#)

Examples

```
# Simple colour-named palette
lares_pal("simple")

# Raw colours and counter-colours
# OR simply: lares_pal("palette")
nice_palette <- lares_pal("colours")
nice_palette_ctr <- as.vector(lares_pal()$palette)
lapply(list(nice_palette, nice_palette_ctr), head)

# Personal colours by name
df <- lares_pal("custom")
df[sample(nrow(df), 5), ]
```

lasso_vars*Most Relevant Features Using Lasso Regression*

Description

Use Lasso regression to identify the most relevant variables that can predict/identify another variable. You might want to compare with `corr_var()` and/or `x2y()` results to compliment the analysis
No need to standardize, center or scale your data. Tidyverse friendly.

Usage

```
lasso_vars(
  df,
  variable,
  ignore = NULL,
  nlambdas = 100,
  nfolds = 10,
  top = 20,
  quiet = FALSE,
  seed = 123,
  ...
)
```

Arguments

<code>df</code>	Dataframe. Any dataframe is valid as ohse will be applied to process categorical values, and values will be standardize automatically.
<code>variable</code>	Variable. Dependent variable or response.
<code>ignore</code>	Character vector. Variables to exclude from study.
<code>nlambdas</code>	Integer. Number of lambdas to be used in a search.
<code>nfolds</code>	Integer. Number of folds for K-fold cross-validation (≥ 2).
<code>top</code>	Integer. Plot top n results only.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>seed</code>	Numeric.
<code>...</code>	Additional parameters passed to <code>ohse()</code> .

Value

List. Contains lasso model coefficients, performance metrics, the actual model fitted and a plot.

See Also

Other Machine Learning: `ROC()`, `conf_mat()`, `export_results()`, `gain_lift()`, `h2o_automl()`, `h2o_predict_MOJO()`, `h2o_selectmodel()`, `impute()`, `iter_seeds()`, `model_metrics()`, `model_preprocess()`, `msplit()`

Other Exploratory: `corr_cross()`, `corr_var()`, `crosstab()`, `df_str()`, `distr()`, `freqs()`, `freqs_df()`, `freqs_list()`, `freqs_plot()`, `missingness()`, `plot_cats()`, `plot_df()`, `plot_nums()`, `tree_var()`

Examples

```
## Not run:
# CRAN
Sys.unsetenv("LARES_FONT") # Temporal
data(dft) # Titanic dataset

m <- lasso_vars(dft, Survived, ignore = c("Cabin"))
print(m$coef)
print(m$metrics)
plot(m$plot)

## End(Not run)
```

left

Left or Right N characters of a string

Description

This functions lets the user extract the first or last n characters of a string or vector of strings.

Usage

```
left(string, n = 1)

right(string, n = 1)
```

Arguments

string	String or Vector.
n	Integer. How many characters starting on right/left?

Value

Character. Trimmed strings.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
left("Bernardo", 3)
right(c("Bernardo", "Lares", "V"), 3)
```

listfiles	<i>List files in a directory</i>
------------------	----------------------------------

Description

This function lets the user list all files on a given directory. It also lets filter files which contains a string.

Usage

```
listfiles(folder = getwd(), recursive = TRUE, regex = NA, images = FALSE)
```

Arguments

folder	Character. Directory which contains files
recursive	Boolean. Should the listing recurse into directories?
regex	Character. String to use for filtering files
images	Boolean. Bring only image files?

Value

data.frame with relevant data for each file on folder directory.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
# All files in current directory (without recursive files)
df <- listfiles(recursive = TRUE)
head(df, 3)

# All files in current directory (with recursive files)
df <- listfiles(recursive = TRUE)
tail(df, 3)

# Check R files using regex
df <- listfiles(regex = "\\.R$")
```

list_cats*List categorical values for data.frame*

Description

Make a list with all categorical values and

Usage

```
list_cats(df, ..., abc = TRUE)
```

Arguments

df	data.frame
...	Variables to segment counters
abc	Boolean. Sort alphabetically?

Value

List. Length same as number of categorical columns, each with a frequency data.frame using freqs().

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
data(dft) # Titanic dataset
df <- dft[, 1:5]
head(df)
list_cats(df)
```

loglossBinary

*Logarithmic Loss Function for Binary Models***Description**

This function calculates log loss/cross-entropy loss for binary models. NOTE: when result is 0.69315, the classification is neutral; it assigns equal probability to both classes.

Usage

```
loglossBinary(tag, score, eps = 0.001)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result
eps	Numeric. Epsilon value

See Also

Other Model metrics: [ROC\(\)](#), [conf_mat\(\)](#), [errors\(\)](#), [gain_lift\(\)](#), [model_metrics\(\)](#)

mail_send

*Send Emails with Attachments (POST)***Description**

This function lets the user send Emails with Attachments using MailGun's API service.

Usage

```
mail_send(
  from = "RMail <laresbernardo@gmail.com>",
  to = "laresbernardo@gmail.com",
  cc = NULL,
  bcc = NULL,
  subject = "Mail from R",
  text = "\n",
  html = NULL,
  attachment = NULL,
  service = "mailgun",
  creds = NULL,
  quiet = FALSE,
  ...
)
```

Arguments

<code>from, to, cc, bcc</code>	Character. Emails
<code>subject</code>	Character. Subject for the email.
<code>text, html</code>	Character. Text or HTML to send in the body.
<code>attachment</code>	Character, plot or data.frame. Will send the file, plot as PNG or data.frame as CSV, respectively.
<code>service</code>	Character. Service platform to search on creds.
<code>creds</code>	Character. Credential's user (see <code>get_creds()</code>). Must contain: url (POST address), api (API key).
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters.

Value

No return value, called for side effects.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

Other Credentials: `db_download()`, `db_upload()`, `encrypt_file()`, `get_credentials()`, `get_tweets()`, `queryDB()`, `queryGA()`, `slackSend()`, `stocks_file()`

Examples

```
## Not run:
myPlot <- noPlot("My plot")
mail_send(
  from = "BLV <myuser@mail.com>",
  to = "youruser@mail.com",
  cc = "myuser@mail.com",
  subject = paste("Daily report:", Sys.Date()),
  attachment = myPlot
)
## End(Not run)
```

markdown2df*Convert markdown string tables to data.frame***Description**

Convert markdown string tables to data.frame

Usage

```
markdown2df(text, autoformat = TRUE)
```

Arguments

- | | |
|-------------------------|--|
| <code>text</code> | Character. Markdown text representing a table. |
| <code>autoformat</code> | Boolean. Automatically format numerical, logical and date values to their classes? |

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
txt <- "| Item | Value |\n|-----|-----|\n| 50C | 122F |\n| 300K | 80.33F |"
markdown2df(txt)
```

maze_solve*Maze Solver; inspired by Micromouse competitions***Description**

Modified recursive depth-first search (DFS) algorithm to solve mazes. It explores the maze by recursively moving to adjacent cells until it finds a path from the starting point to the destination. Contains options to maximize paths by trying to turn less, allowing diagonal turns, prioritizing turns that chooses next step pointing towards the end point, and a grid search combining parameters to find best route.

Usage

```

maze_solve(
  maze,
  start = c(1, 1),
  end = dim(maze),
  inertia = FALSE,
  aim = TRUE,
  diagonal = TRUE,
  random = FALSE,
  timeout = 4,
  quiet = FALSE,
  seed = NULL,
  ...
)

## S3 method for class 'maze_solve'
print(x, ...)

maze_gridsearch(
  maze,
  start = c(2, 2),
  end = round(dim(maze)/2),
  quiet = TRUE,
  seed = 123,
  ...
)

```

Arguments

<code>maze</code>	Matrix. Using 0 for open space and 1 for walls.
<code>start, end</code>	Integer vector, length 2. Start and end coordinates.
<code>inertia</code>	Boolean. When enabled, algorithm will check for new directions only when impossible to continue in a straight line.
<code>aim</code>	Boolean. When enabled, algorithm will try first the directions closer to the end point, ranked and sorted by shorter distances.
<code>diagonal</code>	Boolean. When enabled, algorithm will have 8 degrees of freedom to move, if not, only 4 (up, down, left, right).
<code>random</code>	Boolean. When enabled, algorithm will pick next direction randomly.
<code>timeout</code>	Numeric. How many seconds set for timeout to force algorithm to stop trying new paths?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>seed</code>	Numeric. Seed to replicate random results.
<code>...</code>	Additional parameters passed to <code>corr</code>
<code>x</code>	<code>maze_solve</code> object

Value

List with data.frame containing solved solution, data.frame with path coordinates and directions, steps counter and turns counter.

Examples

```
micromouse <- matrix(c(
  1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
  1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1,
  1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1,
  1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1,
  1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 1, 1,
  1, 0, 0, 0, 1, 1, 1, 1, 0, 1, 0, 1,
  1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 0, 1,
  1, 0, 0, 0, 1, 0, 1, 1, 0, 1, 0, 1,
  1, 0, 1, 1, 1, 1, 0, 1, 1, 1, 0, 1,
  1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 1,
  1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1,
  1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1
), nrow = 12, byrow = TRUE)
maze_solve(micromouse, start = c(2, 2), end = c(7, 7))
```

Description

This function lets the user calculate the percentage of NAs or missingness in a data.frame. It also plots the results if needed.

Usage

```
missingness(df, plot = FALSE, full = FALSE, subtitle = NA, summary = TRUE)
```

Arguments

<code>df</code>	Dataframe. Dataframe to study
<code>plot</code>	Boolean. Do you wish to plot results?
<code>full</code>	Boolean. Return all variables (or only with missings)?
<code>subtitle</code>	Character. Subtitle to show in plot
<code>summary</code>	Boolean. Show numerical summary text?

Value

data.frame with each variable, number of missing values and percentage. If `plot=TRUE`, a plot with the same information reflected.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

Other Missing Values: [impute\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal

# Dummy data
df <- data.frame(
  A = c(1:5),
  B = c(NA, NA, 1, 1, 1),
  C = rep(NA, 5),
  D = c(NA, LETTERS[1:4])
)

# Missing values summary
missingness(df)

# Visual results

missingness(df, plot = TRUE)

# Show all variables (including those with no missing values)
missingness(df, plot = TRUE, full = TRUE)
```

Description

This function lets the user get a confusion matrix and accuracy, and for for binary classification models: AUC, Precision, Sensitivity, and Specificity, given the expected (tags) values and predicted values (scores).

Usage

```
model_metrics(
  tag,
  score,
  multis = NA,
  abc = TRUE,
  thresh = 10,
  auto_n = TRUE,
  thresh_cm = 0.5,
  target = "auto",
  type = "test",
```

```

  model_name = NA,
  plots = TRUE,
  quiet = FALSE,
  subtitle = NA
)

```

Arguments

<code>tag</code>	Vector. Real known label
<code>score</code>	Vector. Predicted value or model's result
<code>multis</code>	Data.frame. Containing columns with each category score (only used when more than 2 categories coexist)
<code>abc</code>	Boolean. Arrange columns and rows alphabetically when categorical values?
<code>thresh</code>	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
<code>auto_n</code>	Add n_ before digits when it's categorical and not numerical, even though seems numerical?
<code>thresh_cm</code>	Numeric. Value to splits the results for the confusion matrix. Range of values: (0-1)
<code>target</code>	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only used when binary categorical model.
<code>type</code>	Character. One of: "train", "test".
<code>model_name</code>	Character. Model's name for reference.
<code>plots</code>	Boolean. Create plots objects?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>subtitle</code>	Character. Subtitle for plots

Value

List. Multiple performance metrics that vary depending on the type of model (classification or regression). If `plot=TRUE`, multiple plots are also returned.

See Also

Other Machine Learning: `ROC()`, `conf_mat()`, `export_results()`, `gain_lift()`, `h2o_automl()`, `h2o_predict_MOJO()`, `h2o_selectmodel()`, `impute()`, `iter_seeds()`, `lasso_vars()`, `model_preprocess()`, `msplit()`

Other Model metrics: `ROC()`, `conf_mat()`, `errors()`, `gain_lift()`, `loglossBinary()`

Other Calculus: `corr()`, `dist2d()`, `quants()`

Examples

```
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# Metrics for Binomial Model
met1 <- model_metrics(dfr$class2$tag, dfr$class2$scores,
  model_name = "Titanic Survived Model",
  plots = FALSE
)
print(met1)

# Metrics for Multi-Categorical Model
met2 <- model_metrics(dfr$class3$tag, dfr$class3$score,
  multis = subset(dfr$class3, select = -c(tag, score)),
  model_name = "Titanic Class Model",
  plots = FALSE
)
print(met2)

# Metrics for Regression Model
met3 <- model_metrics(dfr$regr$tag, dfr$regr$score,
  model_name = "Titanic Fare Model",
  plots = FALSE
)
print(met3)
```

Description

Pre-process your data before training a model. This is the prior step on the `h2o_automl()` function's pipeline. Enabling for other use cases when wanting too use any other framework, library, or custom algorithm.

Usage

```
model_preprocess(
  df,
  y = "tag",
  ignore = NULL,
  train_test = NA,
  split = 0.7,
  weight = NULL,
  target = "auto",
  balance = FALSE,
  impute = FALSE,
  no_outliers = TRUE,
```

```

unique_train = TRUE,
center = FALSE,
scale = FALSE,
thresh = 10,
seed = 0,
quiet = FALSE
)

```

Arguments

df	Dataframe. Dataframe containing all your data, including the dependent variable labeled as 'tag'. If you want to define which variable should be used instead, use the y parameter.
y	Character. Column name for dependent variable or response.
ignore	Character vector. Force columns for the model to ignore
train_test	Character. If needed, df's column name with 'test' and 'train' values to split
split	Numeric. Value between 0 and 1 to split as train/test datasets. Value is for training set. Set value to 1 to train with all available data and test with same data (cross-validation will still be used when training). If train_test is set, value will be overwritten with its real split rate.
weight	Column with observation weights. Giving some observation a weight of zero is equivalent to excluding it from the dataset; giving an observation a relative weight of 2 is equivalent to repeating that row twice. Negative weights are not allowed.
target	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only used when binary categorical model.
balance	Boolean. Auto-balance train dataset with under-sampling?
impute	Boolean. Fill NA values with MICE?
no_outliers	Boolean/Numeric. Remove y's outliers from the dataset? Will remove those values that are farther than n standard deviations from the dependent variable's mean (Z-score). Set to TRUE for default (3) or numeric to set a different multiplier.
unique_train	Boolean. Keep only unique row observations for training data?
center, scale	Boolean. Using the base function scale, do you wish to center and/or scale all numerical values?
thresh	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
seed	Integer. Set a seed for reproducibility. AutoML can only guarantee reproducibility if max_models is used because max_time is resource limited.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

List. Contains original data.frame df, an index to identify which observations will be part of the train dataset train_index, and which model type should be model_type.

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [msplit\(\)](#)

Examples

```
data(dft) # Titanic dataset

model_preprocess(dft, "Survived", balance = TRUE)

model_preprocess(dft, "Fare", split = 0.5, scale = TRUE)

model_preprocess(dft, "Pclass", ignore = c("Fare", "Cabin"))

model_preprocess(dft, "Pclass", quiet = TRUE)
```

move_files*Move files from A to B*

Description

Move one or more files from a directory to another using R.

Usage

```
move_files(from, to)
```

Arguments

- | | |
|-------------------|--|
| <code>from</code> | Character. File names and directories. All files will be moved recursively. |
| <code>to</code> | Character. File names for each <code>from</code> file or directory. If directory does not exist, it will be created. |

Value

No return value, called for side effects.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepml()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

mplot_conf*Confussion Matrix Plot***Description**

This function plots a confussion matrix.

Usage

```
mplot_conf(
  tag,
  score,
  thresh = 0.5,
  abc = TRUE,
  squared = FALSE,
  diagonal = TRUE,
  top = 20,
  subtitle = NA,
  model_name = NULL,
  save = FALSE,
  subdir = NA,
  file_name = "viz_conf_mat.png"
)
```

Arguments

<code>tag</code>	Vector. Real known label.
<code>score</code>	Vector. Predicted value or model's result.
<code>thresh</code>	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
<code>abc</code>	Boolean. Arrange columns and rows alphabetically?
<code>squared</code>	Boolean. Force plot to be squared?
<code>diagonal</code>	Boolean. FALSE to convert diagonal numbers to zeroes. Ideal to detect must confusing categories.
<code>top</code>	Integer. Plot only the most n frequent variables. Set to NA to plot all.

subtitle	Character. Subtitle to show in plot
model_name	Character. Model's name
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Details

You may use `conf_mat()` to get calculate values.

Value

Plot with confusion matrix results.

See Also

Other ML Visualization: `mplot_cuts()`, `mplot_cuts_error()`, `mplot_density()`, `mplot_full()`, `mplot_gain()`, `mplot_importance()`, `mplot_lineal()`, `mplot_metrics()`, `mplot_response()`, `mplot_roc()`, `mplot_splits()`, `mplot_topcats()`

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# Plot for Binomial Model
mplot_conf(dfr$class2$tag, dfr$class2$scores,
  model_name = "Titanic Survived Model"
)

# Plot for Multi-Categorical Model
mplot_conf(dfr$class3$tag, dfr$class3$score,
  model_name = "Titanic Class Model"
)
```

<code>mplot_cuts</code>	<i>Cuts by quantiles for score plot</i>
-------------------------	---

Description

This function cuts by quantiles any score or prediction.

Usage

```
mplot_cuts(
  score,
  splits = 10,
  model_name = NA,
  subtitle = NA,
  table = FALSE,
  save = FALSE,
  subdir = NA,
  file_name = "viz_ncuts.png"
)
```

Arguments

score	Vector. Predicted value or model's result.
splits	Integer. Numer of separations to plot
model_name	Character. Model's name
subtitle	Character. Subtitle to show in plot
table	Boolean. Do you wish to return a table with results?
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Value

Plot with performance results by cuts.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
head(dfr$class2)

# Data
mplot_cuts(dfr$class2$scores, splits = 5, table = TRUE)

# Plot
mplot_cuts(dfr$class2$scores, model_name = "Titanic Survived Model")
```

mplot_cuts_error *Cuts by quantiles on absolute and percentual errors plot*

Description

This function cuts by quantiles on absolute and percentual errors

Usage

```
mplot_cuts_error(  
  tag,  
  score,  
  splits = 10,  
  title = NA,  
  model_name = NA,  
  save = FALSE,  
  subdir = NA,  
  file_name = "viz_ncuts_error.png"  
)
```

Arguments

tag	Vector. Real known label.
score	Vector. Predicted value or model's result.
splits	Integer. Number of separations to plot
title	Character. Title to show in plot
model_name	Character. Model's name
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Value

Plot with error results by cuts.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
head(dfr$regr)
mplot_cuts_error(dfr$regr$tag, dfr$regr$score,
  model_name = "Titanic Fare Model"
)
```

mplot_density

Density plot for discrete and continuous values

Description

This function plots discrete and continuous values results

Usage

```
mplot_density(
  tag,
  score,
  thresh = 6,
  model_name = NA,
  subtitle = NA,
  save = FALSE,
  subdir = NA,
  file_name = "viz_distribution.png"
)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result
thresh	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
model_name	Character. Model's name
subtitle	Character. Subtitle to show in plot
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Value

Plot with distribution and performance results.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr[c(1, 3)], head)

# Plot for binomial results
mplot_density(dfr$class2$tag, dfr$class2$scores, subtitle = "Titanic Survived Model")

# Plot for regression results
mplot_density(dfr$regr$tag, dfr$regr$score, model_name = "Titanic Fare Model")
```

mplot_full

MPLOTS Score Full Report Plots

Description

This function plots a whole dashboard with a model's results. It will automatically detect if it's a categorical or regression's model by checking how many different unique values the dependent variable (tag) has.

Usage

```
mplot_full(
  tag,
  score,
  multis = NA,
  splits = 8,
  thresh = 6,
  subtitle = NA,
  model_name = NA,
  plot = TRUE,
  save = FALSE,
  subdir = NA,
  file_name = "viz_full.png"
)
```

Arguments

tag	Vector. Real known label.
score	Vector. Predicted value or model's result.

<code>multis</code>	Data.frame. Containing columns with each category probability or score (only used when more than 2 categories coexist).
<code>splits</code>	Integer. Number of separations to plot
<code>thresh</code>	Integer. Threshold for selecting binary or regression models: this number is the threshold of unique values we should have in 'tag' (more than: regression; less than: classification)
<code>subtitle</code>	Character. Subtitle to show in plot
<code>model_name</code>	Character. Model's name
<code>plot</code>	Boolean. Plot results? If not, plot grid object returned
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot

Value

Multiple plots gathered into one, showing tag vs score performance results.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# Dasboard for Binomial Model
mplot_full(dfr$class2$tag, dfr$class2$scores,
  model_name = "Titanic Survived Model"
)

# Dasboard for Multi-Categorical Model
mplot_full(dfr$class3$tag, dfr$class3$score,
  multis = subset(dfr$class3, select = -c(tag, score)),
  model_name = "Titanic Class Model"
)

# Dasboard for Regression Model
mplot_full(dfr$regr$tag, dfr$regr$score,
  model_name = "Titanic Fare Model"
)
```

<code>mplot_gain</code>	<i>Cumulative Gain Plot</i>
-------------------------	-----------------------------

Description

The cumulative gains plot, often named ‘gains plot’, helps us answer the question: When we apply the model and select the best X deciles, what expect to target? The cumulative gains chart shows the percentage of the overall number of cases in a given category “gained” by targeting a percentage of the total number of cases.

Usage

```
mplot_gain(
  tag,
  score,
  multis = NA,
  target = "auto",
  splits = 10,
  highlight = "auto",
  caption = NA,
  save = FALSE,
  subdir = NA,
  file_name = "viz_gain.png",
  quiet = FALSE
)
```

Arguments

<code>tag</code>	Vector. Real known label.
<code>score</code>	Vector. Predicted value or model’s result.
<code>multis</code>	Data.frame. Containing columns with each category probability or score (only used when more than 2 categories coexist).
<code>target</code>	Value. Which is your target positive value? If set to ‘auto’, the target with largest mean(score) will be selected. Change the value to overwrite. Only works for binary classes
<code>splits</code>	Integer. Numer of quantiles to split the data
<code>highlight</code>	Character or Integer. Which split should be used for the automatic conclusion in the plot? Set to “auto” for best value, “none” to turn off or the number of split.
<code>caption</code>	Character. Caption to show in plot
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

Plot with gain and performance results by cuts.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# Plot for Binomial Model
mplot_gain(dfr$class2$tag, dfr$class2$scores,
            caption = "Titanic Survived Model",
            target = "FALSE"
        )
mplot_gain(dfr$class2$tag, dfr$class2$scores,
            caption = "Titanic Survived Model",
            target = "TRUE"
        )

# Plot for Multi-Categorical Model
mplot_gain(dfr$class3$tag, dfr$class3$score,
            multis = subset(dfr$class3, select = -c(tag, score)),
            caption = "Titanic Class Model"
        )
```

mplot_importance *Variables Importances Plot*

Description

This function plots Variable Importances

Usage

```
mplot_importance(
    var,
    imp,
    colours = NA,
    limit = 15,
    model_name = NA,
    subtitle = NA,
    save = FALSE,
```

```

    subdir = NA,
    file_name = "viz_importance.png"
)

```

Arguments

var	Vector. Variable or column's names
imp	Vector. Importance of said variables. Must have same length as var
colours	If positive and negative contribution is known
limit	Integer. Limit how many variables you wish to plot
model_name	Character. Model's name
subtitle	Character. Subtitle to show in plot
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Value

Plot with ranked importance variables results.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```

Sys.unsetenv("LARES_FONT") # Temporal
df <- data.frame(
  variable = LETTERS[1:6],
  importance = c(4, 6, 6.7, 3, 4.8, 6.2) / 100,
  positive = c(TRUE, TRUE, FALSE, TRUE, FALSE, FALSE)
)
head(df)

mplot_importance(
  var = df$variable,
  imp = df$importance,
  model_name = "Random values model"
)

# Add a colour for categories
mplot_importance(
  var = df$variable,
  imp = df$importance,
  colours = df$positive,
  limit = 4
)

```

mplot_lineal*Linear Regression Results Plot***Description**

This function plots a Linear Regression Result

Usage

```
mplot_lineal(
  tag,
  score,
  subtitle = NA,
  model_name = NA,
  save = FALSE,
  subdir = NA,
  file_name = "viz_lineal.png"
)
```

Arguments

<code>tag</code>	Vector. Real known label.
<code>score</code>	Vector. Predicted value or model's result.
<code>subtitle</code>	Character. Subtitle to show in plot
<code>model_name</code>	Character. Model's name
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot

Value

Plot with linear distribution and performance results.

See Also

Other ML Visualization: `mplot_conf()`, `mplot_cuts()`, `mplot_cuts_error()`, `mplot_density()`, `mplot_full()`, `mplot_gain()`, `mplot_importance()`, `mplot_metrics()`, `mplot_response()`, `mplot_roc()`, `mplot_splits()`, `mplot_topcats()`

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)
mplot_lineal(dfr$regr$tag, dfr$regr$score, model_name = "Titanic Fare Model")
```

mplot_metrics*Model Metrics and Performance Plots*

Description

This function generates plots of the metrics of a predictive model. This is an auxiliary function used in `model_metrics()` when the parameter `plot` is set to TRUE.

Usage

```
mplot_metrics(  
  results,  
  subtitle = NA,  
  model_name = NA,  
  save = FALSE,  
  subdir = NA,  
  file_name = "viz_metrics.png"  
)
```

Arguments

<code>results</code>	Object. Results object from <code>h2o_automl</code> function
<code>subtitle</code>	Character. Subtitle to show in plot
<code>model_name</code>	Character. Model's name
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot

Value

Plot with `results` performance.

See Also

Other ML Visualization: `mplot_conf()`, `mplot_cuts()`, `mplot_cuts_error()`, `mplot_density()`, `mplot_full()`, `mplot_gain()`, `mplot_importance()`, `mplot_lineal()`, `mplot_response()`, `mplot_roc()`, `mplot_splits()`, `mplot_topcats()`

mplot_response	<i>Cumulative Response Plot</i>
----------------	---------------------------------

Description

The response gains plot helps us answer the question: When we apply the model and select up until ntile X, what is the expected

Usage

```
mplot_response(
  tag,
  score,
  multis = NA,
  target = "auto",
  splits = 10,
  highlight = "auto",
  caption = NA,
  save = FALSE,
  subdir = NA,
  file_name = "viz_response.png",
  quiet = FALSE
)
```

Arguments

<code>tag</code>	Vector. Real known label.
<code>score</code>	Vector. Predicted value or model's result.
<code>multis</code>	Data.frame. Containing columns with each category probability or score (only used when more than 2 categories coexist).
<code>target</code>	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only works for binary classes
<code>splits</code>	Integer. Numer of quantiles to split the data
<code>highlight</code>	Character or Integer. Which split should be used for the automatic conclusion in the plot? Set to "auto" for best value, "none" to turn off or the number of split.
<code>caption</code>	Character. Caption to show in plot
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

Plot with cumulative response and performance results by cuts.

See Also

Other ML Visualization: `mplot_conf()`, `mplot_cuts()`, `mplot_cuts_error()`, `mplot_density()`, `mplot_full()`, `mplot_gain()`, `mplot_importance()`, `mplot_lineal()`, `mplot_metrics()`, `mplot_roc()`, `mplot_splits()`, `mplot_topcats()`

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# Plot for Binomial Model
mplot_response(dfr$class2$tag, dfr$class2$scores,
  caption = "Titanic Survived Model",
  target = "TRUE"
)
mplot_response(dfr$class2$tag, dfr$class2$scores,
  caption = "Titanic Survived Model",
  target = "FALSE"
)

# Plot for Multi-Categorical Model
mplot_response(dfr$class3$tag, dfr$class3$score,
  multis = subset(dfr$class3, select = -c(tag, score)),
  caption = "Titanic Class Model"
)
```

`mplot_roc`

ROC Curve Plot

Description

This function plots ROC Curves with AUC values with 95% confidence range. It also works for multi-categorical models.

Usage

```
mplot_roc(
  tag,
  score,
  multis = NA,
  sample = 1000,
  model_name = NA,
  subtitle = NA,
  interval = 0.2,
  squared = TRUE,
  plotly = FALSE,
  save = FALSE,
```

```

    subdir = NA,
    file_name = "viz_roc.png"
)

```

Arguments

<code>tag</code>	Vector. Real known label.
<code>score</code>	Vector. Predicted value or model's result.
<code>multis</code>	Data.frame. Containing columns with each category probability or score (only used when more than 2 categories coexist).
<code>sample</code>	Integer. Number of samples to use for rendering plot.
<code>model_name</code>	Character. Model's name
<code>subtitle</code>	Character. Subtitle to show in plot
<code>interval</code>	Numeric. Interval for breaks in plot
<code>squared</code>	Boolean. Keep proportions?
<code>plotly</code>	Boolean. Use plotly for plot's output for an interactive plot
<code>save</code>	Boolean. Save output plot into working directory
<code>subdir</code>	Character. Sub directory on which you wish to save the plot
<code>file_name</code>	Character. File name as you wish to save the plot

Value

Plot with ROC curve and AUC performance results.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_splits\(\)](#), [mplot_topcats\(\)](#)

Examples

```

Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr[c(1, 2)], head)

# ROC Curve for Binomial Model
mplot_roc(dfr$class2$tag, dfr$class2$scores,
           model_name = "Titanic Survived Model"
)

# ROC Curves for Multi-Categorical Model
mplot_roc(dfr$class3$tag, dfr$class3$score,
           multis = subset(dfr$class3, select = -c(tag, score)),
           squared = FALSE,
           model_name = "Titanic Class Model"
)

```

`mplot_splits`*Split and compare quantiles plot*

Description

This function lets us split and compare quantiles on a given prediction to compare different categorical values vs scores grouped by equal sized buckets.

Usage

```
mplot_splits(  
  tag,  
  score,  
  splits = 5,  
  subtitle = NA,  
  model_name = NA,  
  save = FALSE,  
  subdir = NA,  
  file_name = "viz_splits.png"  
)
```

Arguments

tag	Vector. Real known label.
score	Vector. Predicted value or model's result.
splits	Integer. Number of separations to plot
subtitle	Character. Subtitle to show in plot
model_name	Character. Model's name
save	Boolean. Save output plot into working directory
subdir	Character. Sub directory on which you wish to save the plot
file_name	Character. File name as you wish to save the plot

Value

Plot with distribution and performance results by splits.

See Also

Other ML Visualization: `mplot_conf()`, `mplot_cuts()`, `mplot_cuts_error()`, `mplot_density()`, `mplot_full()`, `mplot_gain()`, `mplot_importance()`, `mplot_lineal()`, `mplot_metrics()`, `mplot_response()`, `mplot_roc()`, `mplot_topcats()`

Examples

```

Sys.unsetenv("LARES_FONT") # Temporal
data(dfr) # Results for AutoML Predictions
lapply(dfr, head)

# For categorical (binary) values
mplot_splits(dfr$class2$tag, dfr$class2$scores,
  splits = 4,
  model_name = "Titanic Survived Model"
)

# For categorical (+2) values
mplot_splits(dfr$class3$tag, dfr$class2$scores,
  model_name = "Titanic Class Model"
)

# For continuous values
mplot_splits(dfr$regr$tag, dfr$regr$score,
  splits = 4,
  model_name = "Titanic Fare Model"
)

```

mplot_topcats

Top Hit Ratios for Multi-Classification Models

Description

Calculate and plot a multi-class model's predictions accuracy based on top N predictions and distribution of probabilities.

Usage

```
mplot_topcats(tag, score, multis, model_name = NA)
```

Arguments

tag	Vector. Real known label.
score	Vector. Predicted value or model's result.
multis	Data.frame. Containing columns with each category probability or score (only used when more than 2 categories coexist).
model_name	Character. Model's name

Value

Plot with performance results over most frequent categories.

See Also

Other ML Visualization: [mplot_conf\(\)](#), [mplot_cuts\(\)](#), [mplot_cuts_error\(\)](#), [mplot_density\(\)](#), [mplot_full\(\)](#), [mplot_gain\(\)](#), [mplot_importance\(\)](#), [mplot_lineal\(\)](#), [mplot_metrics\(\)](#), [mplot_response\(\)](#), [mplot_roc\(\)](#), [mplot_splits\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal  
data(dfr) # Results for AutoML Predictions  
mplot_topcats(dfr$class3$tag, dfr$class3$score,  
    multis = subset(dfr$class3, select = -c(tag, score)),  
    model_name = "Titanic Class Model"  
)
```

msplit

Split a dataframe for training and testing sets

Description

This function splits automatically a dataframe into train and test datasets. You can define a seed to get the same results every time, but has a default value. You can prevent it from printing the split counter result.

Usage

```
msplit(df, size = 0.7, seed = 0, print = TRUE)
```

Arguments

df	Dataframe
size	Numeric. Split rate value, between 0 and 1. If set to 1, the train and test set will be the same.
seed	Integer. Seed for random split
print	Boolean. Print summary results?

Value

List with both datasets, summary, and split rate.

See Also

Other Machine Learning: [ROC\(\)](#), [conf_mat\(\)](#), [export_results\(\)](#), [gain_lift\(\)](#), [h2o_automl\(\)](#), [h2o_predict_MOJO\(\)](#), [h2o_selectmodel\(\)](#), [impute\(\)](#), [iter_seeds\(\)](#), [lasso_vars\(\)](#), [model_metrics\(\)](#), [model_preprocess\(\)](#)

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#),

```
formatHTML(), glued(), grepM(), h2o_selectmodel(), haveInternet(), image_metadata(),
importxlsx(), ip_data(), json2vector(), list_cats(), listfiles(), mail_send(), markdown2df(),
move_files(), myip(), quiet(), read.file(), statusbar(), tic(), try_require(), updateLares(),
warnifnot(), what_size()
```

Examples

```
data(dft) # Titanic dataset
splits <- msplit(dft, size = 0.7, seed = 123)
names(splits)
```

`myip`

What's my IP?

Description

Reveal your current IP address.

Usage

`myip()`

Value

Character. Result of your IP address based on ipify.org

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`,
`dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`,
`formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`,
`importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`,
`move_files()`, `msplit()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`,
`warnifnot()`, `what_size()`

Examples

`myip()`

ngrams	<i>Build N-grams and keep most frequent</i>
--------	---

Description

Build out n-grams for multiple text inputs and keep the n most frequent combinations.

Usage

```
ngrams(text, ngram = c(2, 3), top = 10, stop_words = NULL, ...)
```

Arguments

text	Character vector
ngram	Integer vector. Number of continuous n items in text.
top	Integer. Keep n most frequent ngrams only.
stop_words	Character vector. Words to exclude from text. Example: if you want to exclude "a", whenever that word appears it will be excluded, but when the letter "a" appears in a word, it will remain.
...	Additional parameters passed to <code>remove_stopwords</code> .

Value

data.frame with ngrams and counters, sorted by frequency.

See Also

Other Text Mining: [cleanText\(\)](#), [remove_stopwords\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [topics_rake\(\)](#)

Examples

```
# You must have "tidytext" library to use this auxiliary function:  
## Not run:  
women <- read.csv("https://bit.ly/3mXJ00i")  
x <- women$description  
ngrams(x, ngram = c(2, 3), top = 3)  
ngrams(x, ngram = 2, top = 6, stop_words = c("a", "is", "of", "the"))  
## End(Not run)
```

noPlot*Plot Result with Nothing to Plot***Description**

This function lets the user print a plot without plot, with a customizable message. It is quite useful for Shiny renderPlot when using filters and no data is returned.

Usage

```
noPlot(message = "Nothing to show here!", size = 4.5, ...)
```

Arguments

<code>message</code>	Character. What message do you wish to show?
<code>size</code>	Numeric. Font size for message input.
<code>...</code>	Additional parameters passed to <code>theme_lares()</code> .

Value

Empty ggplot2 object (with a message if set).

See Also

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
noPlot(message = "No plot to show!")
noPlot(background = "#FF5500", size = 7)
```

normalize*Normalize Vector***Description**

This function normalizes numerical values into a specified range, defaulting to the 0 to 1 range.

Usage

```
normalize(x, range = c(0, 1), ...)
```

Arguments

- `x` Numeric vector. The numbers to be normalized.
`range` A numeric vector of length 2 specifying the desired range for normalization.
 Default is `c(0, 1)`.
`...` Additional parameters.

Value

A numeric vector with normalized `x` values.

See Also

Other Data Wrangling: `balance_data()`, `categ_reducer()`, `cleanText()`, `date_cuts()`, `date_feats()`, `file_name()`, `formatHTML()`, `holidays()`, `impute()`, `left()`, `num_abbr()`, `ohe_commas()`, `ohse()`, `quants()`, `removenacols()`, `replaceall()`, `replacefactor()`, `textFeats()`, `textTokenizer()`, `vector2text()`, `year_month()`, `zerovar()`

Examples

```
x <- c(0, 1, 4, 7.5, 10)
normalize(x)
normalize(x, range = c(-1, 1))
```

num_abbr

Abbreviate or dis-abbreviate numerical values

Description

This function converts a numeric vector's values into their abbreviated character equivalent, i.e. 100E6 into 100M and viceversa.

Usage

```
num_abbr(x, n = 3, numeric = FALSE, ...)
```

Arguments

- `x` Numeric vector
`n` Integer. Single numeric value, specifying number of significant figures to show.
 Range 1 to 6.
`numeric` Boolean. Transform abbreviated number into numeric?
`...` Additional parameters.

Value

Vector of character or numeric values that contain converted values

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
num_abbr(rnorm(10) * 1e6)
num_abbr(rnorm(10) * 1e6, n = 1)
num_abbr(c("3K", "-58.3M", NA, 1), numeric = TRUE)
```

ohe_commas*One Hot Encoding for a Vector with Comma Separated Values***Description**

This function lets the user do one hot encoding on a variable with comma separated values

Usage

```
ohe_commas(df, ..., sep = ",", noval = "NoVal", remove = FALSE)
```

Arguments

<i>df</i>	Dataframe. May contain one or more columns with comma separated values which will be separated as one hot encoding
<i>...</i>	Variables. Which variables to split into new columns?
<i>sep</i>	Character. Which regular expression separates the elements?
<i>noval</i>	Character. No value text
<i>remove</i>	Boolean. Remove original variables?

Value

data.frame on which all features are numerical by nature or transformed with one hot encoding.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other One Hot Encoding: [date_feats\(\)](#), [holidays\(\)](#), [ohse\(\)](#)

Examples

```
df <- data.frame(
  id = c(1:5),
  x = c("AA, D", "AA,B", "B, D", "A,D,B", NA),
  z = c("AA+BB+AA", "AA", "BB, AA", NA, "BB+AA")
)
ohe_commas(df, x, remove = TRUE)
ohe_commas(df, z, sep = "\\\\+")
ohe_commas(df, x, z)
```

ohse

One Hot Smart Encoding (Dummy Variables)

Description

This function lets the user automatically transform a dataframe with categorical columns into numerical by one hot encoding technic.

Usage

```
ohse(
  df,
  redundant = FALSE,
  drop = TRUE,
  ignore = NULL,
  dates = FALSE,
  holidays = FALSE,
  country = "Venezuela",
  currency_pair = NA,
  trim = 0,
  limit = 10,
  variance = 0.9,
  other_label = "OTHER",
  sep = "_",
  quiet = FALSE,
  ...
)
```

Arguments

df	Dataframe
redundant	Boolean. Should we keep redundant columns? i.e. If the column only has two different values, should we keep both new columns? Is set to NULL, only binary variables will dump redundant columns.
drop	Boolean. Drop automatically some useless features?
ignore	Vector or character. Which column should be ignored?

<code>dates</code>	Boolean. Do you want the function to create more features out of the date/time columns?
<code>holidays</code>	Boolean. Include holidays as new columns?
<code>country</code>	Character or vector. For which countries should the holidays be included?
<code>currency_pair</code>	Character. Which currency exchange do you wish to get the history from? i.e., USD/COP, EUR/USD...
<code>trim</code>	Integer. Trim names until the nth character
<code>limit</code>	Integer. Limit one hot encoding to the n most frequent values of each column. Set to NA to ignore argument.
<code>variance</code>	Numeric. Drop columns with more than n variance. Range: 0-1. For example: if a variable contains 91 unique different values out of 100 observations, this column will be suppressed if value is set to 0.9
<code>other_label</code>	Character. With which text do you wish to replace the filtered values with?
<code>sep</code>	Character. Separator's string
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters.

Value

`data.frame` on which all features are numerical by nature or transformed with one hot encoding.

See Also

Other Data Wrangling: `balance_data()`, `categ_reducer()`, `cleanText()`, `date_cuts()`, `date_feats()`, `file_name()`, `formatHTML()`, `holidays()`, `impute()`, `left()`, `normalize()`, `num_abbr()`, `ohe_commas()`, `quants()`, `removenacols()`, `replaceall()`, `replacefactor()`, `textFeats()`, `textTokenizer()`, `vector2text()`, `year_month()`, `zerovar()`

Other Feature Engineering: `date_feats()`, `holidays()`

Other One Hot Encoding: `date_feats()`, `holidays()`, `ohe_commas()`

Examples

```
data(dft)
dft <- dft[, c(2, 3, 5, 9, 11)]

ohse(dft, limit = 3) %>% head(3)
ohse(dft, limit = 3, redundant = NULL) %>% head(3)

# Getting rid of columns with no (or too much) variance
dft$no_variance1 <- 0
dft$no_variance2 <- c("A", rep("B", nrow(dft) - 1))
dft$no_variance3 <- as.character(rnorm(nrow(dft)))
dft$no_variance4 <- c(rep("A", 20), round(rnorm(nrow(dft) - 20), 4))
ohse(dft, limit = 3) %>% head(3)
```

outlier_tukey *Outliers: Tukey's fences*

Description

Tukey's fences is a technique used in box plots. The non-outlier range is defined with $[Q1 - k(Q3 - Q1), Q3 + k(Q3 - Q1)]$, where $Q1$ and $Q3$ are the lower and upper quartiles respectively, k - some non-negative constant (popular choice is 1.5). A value is an outlier based on Tukey's fences when its value does not lie in non-outlier range.

Usage

```
outlier_tukey(x, k = 1.5)
```

```
outlier_tukey(x, k = 1.5)
```

Arguments

x	Numeric. Distribution
k	Positive Numeric. K-multiplier.

Value

Boolean vector detecting outliers.

See Also

Other Outliers: [outlier_zscore\(\)](#), [outlier_zscore_plot\(\)](#), [winsorize\(\)](#)

outlier_zscore *Outliers: Z-score method*

Description

Z-score, also called a standard score, of an observation is a distance from the population center measured in number of normalization units. The default choice for center is sample mean and for normalization unit is standard deviation. Values are considered outliers based on z-score if its absolute value of default z-score is higher than the threshold (popular choice is 3).

Usage

```
outlier_zscore(x, thresh = 3, mad = FALSE)
```

Arguments

x	Numeric. Distribution
thresh	Numeric. Z-Score threshold for n standard deviations.
mad	Boolean. Use median absolute deviation instead?

Value

`data.frame`. Each row is an `x` observation with its respective std/mean or mad/med calculations depending on `mad` input.

See Also

Other Outliers: [outlier_tukey\(\)](#), [outlier_zscore\(\)](#), [winsorize\(\)](#)

[outlier_zscore](#) *Outliers: Z-score method plot*

Description

Test several Z-score thresholds to visualize outliers. Tidyverse friendly.

Usage

```
outlier_zscore_plot(df, var, group = NULL, thresh = c(2, 3, 5), top = 5)
```

Arguments

df	Dataframe.
var	Numeric variable.
group	Categorical variable. Grouping variable.
thresh	Numeric vector. Z-Score threshold for n standard deviations.
top	Integer. Show only n most frequent categorical values when using the <code>group</code> argument.

Value

`ggplot2` object

See Also

Other Outliers: [outlier_tukey\(\)](#), [outlier_zscore\(\)](#), [winsorize\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal  
data(dft) # Titanic dataset  
outlier_zscore_plot(dft, Fare)  
p <- outlier_zscore_plot(dft, Fare, Pclass, thresh = c(3, 5))  
plot(p)  
attr(p, "z_values")  
head(attr(p, "z_labels"))
```

plot_cats

Plot All Categorical Features (Frequencies)

Description

This function filters categorical columns and plots the frequency for each value on every feature.

Usage

```
plot_cats(df)
```

Arguments

df Dataframe

Value

Plot. Result of df categorical features.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

plot_chord

Chords Plot

Description

This auxiliary function plots discrete and continuous values results

Usage

```
plot_chord(
  origin,
  dest,
  weight = 1,
  mg = 3,
  title = "Chord Diagram",
  subtitle = "",
  pal = NA
)
```

Arguments

<code>origin, dest</code>	Vectors. Origin and destination vectors
<code>weight</code>	Vector. Weight for each chord.
<code>mg</code>	Numeric. Margin adjust for plot in case of need
<code>title</code>	Character. Title for the plot
<code>subtitle</code>	Character. Subtitle for the plot
<code>pal</code>	Vector. Colour palette. Order matters.

Value

chordDiagram object

See Also

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

Examples

```
# You must have "circlize" library to use this auxiliary function:
## Not run:
df <- data.frame(from = c(1, 1, 2, 3, 4, 1, 6), to = c(4, 4, 4, 2, 2, NA, NA))
plot_chord(df$from, df$to)

## End(Not run)
```

Description

This function plots all columns frequencies and boxplots, for categorical and numerical respectively.

Usage

```
plot_df(df)
```

Arguments

df Dataframe

Value

Plot. Result of df categorical and numerical features.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_nums\(\)](#), [tree_var\(\)](#)

plot_nums

Plot All Numerical Features (Boxplots)

Description

This function filters numerical columns and plots boxplots.

Usage

```
plot_nums(df)
```

Arguments

df Dataframe

Value

Plot. Result of df numerical features.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [tree_var\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal  
data(dft) # Titanic dataset  
plot_nums(dft)
```

plot_palette*Plot Palette Colours*

Description

This function plots a list of colours

Usage

```
plot_palette(fill, colour = "black", id = NA, limit = 12, ...)
```

Arguments

fill	Vector. List of colours for fills.
colour	Vector. List of colours for colours.
id	Vector. ID for each color.
limit	Integer. Show only first n values.
...	Additional parameters.

Value

Plot with fill colours and colour counter-colours if provided.

See Also

Other Themes: [gg_fill_customs\(\)](#), [lares_pal\(\)](#), [theme_lares\(\)](#)

Examples

```
# Simply pass a vector
pal <- lares_pal("simple")
plot_palette(pal)
# Or fill + color named vector
pal <- lares_pal("pal")
plot_palette(fill = names(pal), colour = as.vector(pal))
```

plot_survey	<i>Visualize Survey Results</i>
-------------	---------------------------------

Description

This function lets the user plot a survey's result.

Usage

```
plot_survey(answers, ignore = 1, title = NA, subtitle = NA)
```

Arguments

answers	Dataframe. Answers. Each row a different person. Each column a different answer.
ignore	Numeric Vector. Which columns are NOT answers?
title	Character. Title for your plot
subtitle	Character. Subtitle for your plot.

Value

ggplot2 object

See Also

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_timeline\(\)](#), [tree_var\(\)](#)

plot_timeline	<i>Plot timeline as Gantt Plot</i>
---------------	------------------------------------

Description

This function plots groups of observations with timelines in a Gantt Plot way. Only works if start and end are date format values.

Usage

```
plot_timeline(  
  event,  
  start,  
  end = start + 1,  
  label = NA,  
  group = NA,  
  title = "Curriculum Vitae Timeline",
```

```

subtitle = "Bernardo Lares",
interactive = FALSE,
save = FALSE,
subdir = NA
)

```

Arguments

event	Vector. Event, role, label, or row.
start	Vector. Start date.
end	Vector. End date. Only one day by default if not defined
label	Vector. Place, institution, or label.
group	Vector. Academic, Work, Extracurricular... Pass as factor to keep a specific order
title	Character. Title for the plot
subtitle	Character. Subtitle for the plot
interactive	Boolean. Run with plotly?
save	Boolean. Save the output plot in our working directory
subdir	Character. Into which subdirectory do you wish to save the plot to?

Value

ggplot2 object

See Also

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [tree_var\(\)](#)

Examples

```

Sys.unsetenv("LARES_FONT") # Temporal
cols <- c("Role", "Place", "Type", "Start", "End")
today <- as.character(Sys.Date())
cv <- data.frame(rbind(
  c("Marketing Science Partner", "Facebook", "Work Experience", "2019-12-09", today),
  c("Data Scientist Consultant", "MatrixDS", "Work Experience", "2018-09-01", today),
  c("R Community Contributor", "lares library", "Extra", "2018-07-18", today),
  c("Lead Data Scientist", "MEG", "Work Experience", "2019-01-15", "2019-12-09"),
  c("Head of Analytics", "Comparamejor/R5", "Work Experience", "2016-08-01", "2019-01-15"),
  c("Big Data & Data Science Programme", "UdC", "Academic", "2017-09-01", "2018-02-28"),
  c("Project Engineer", "Polytex", "Work Experience", "2016-05-15", "2016-09-01"),
  c("Big Data Analyst", "MEG", "Work Experience", "2016-01-01", "2016-04-30"),
  c("Advanced Excel Instructor", "ARTS", "Work Experience", "2015-11-01", "2016-04-30"),
  c("Continuous Improvement Intern", "PAVCO", "Work Experience", "2015-04-01", "2015-08-30"),
  c("Mechanical Design Intern", "SIGALCA", "Work Experience", "2013-07-01", "2013-09-30"),
  c("DJs Online Community Owner", "LaresDJ.com / SoloParaDJs", "Extra", "2010-01-05", "2020-05-20"),
  c("Mechanical Engineer Degree", "USB", "Academic", "2009-09-15", "2015-11-20"),

```

```

c("DJ and Composer/Producer", "Legacy Discplay", "Extra", "2009-05-01", "2015-04-30")
))
colnames(cv) <- cols
plot_timeline(
  event = cv$Role,
  start = cv$Start,
  end = cv$End,
  label = cv$Place,
  # Simple trick to re-arrange the grids
  group = factor(cv$type, levels = c("Work Experience", "Academic", "Extra"))
)

```

prophesize*Facebook's Prophet Forecast***Description**

Prophet is Facebook's procedure for forecasting time series data based on an additive model where non-linear trends are fit with yearly, weekly, and daily seasonality, plus holiday effects. It works best with time series that have strong seasonal effects and several seasons of historical data. Prophet is robust to missing data and shifts in the trend, and typically handles outliers well.

Usage

```

prophesize(
  df,
  n_future = 60,
  country = NULL,
  trend.param = 0.05,
  logged = FALSE,
  pout = 0.03,
  project = "Prophet Forecast"
)
```

Arguments

<code>df</code>	Data frame. Must contain date/time column and values column, in that order.
<code>n_future</code>	Integer. How many steps do you wish to forecast?
<code>country</code>	Character. Country code for holidays.
<code>trend.param</code>	Numeric. Flexibility of trend component. Default is 0.05, and as this value becomes larger, the trend component will be more flexible.
<code>logged</code>	Boolean. Convert values into logs?
<code>pout</code>	Numeric. Get rid of pout % of outliers.
<code>project</code>	Character. Name of your forecast project for plot title

Details

Official documentation: <https://github.com/facebook/prophet>

Value

List. Containing the forecast results, the prophet model, and a plot.

See Also

Other Forecast: [forecast_arima\(\)](#)

quants

Calculate cuts by quantiles

Description

This function lets the user quickly calculate cuts for quantiles and discretize numerical values into categorical values.

Usage

```
quants(values, splits = 10, return = "labels", n = 2)
```

Arguments

<code>values</code>	Vector. Values to calculate quantile cuts
<code>splits</code>	Integer. How many cuts should split the values?
<code>return</code>	Character. Return "summary" or "labels"
<code>n</code>	Integer. Determines the number of digits used in formatting the break numbers.

Value

Factor vector or data.frame. Depending on `return` input:

- `labels` a factor ordered vector with each observation's quantile
- `summary` a data.frame with information on each quantile cut

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Calculus: [corr\(\)](#), [dist2d\(\)](#), [model_metrics\(\)](#)

Examples

```
data(dft) # Titanic dataset  
quants(dft$Age, splits = 5, "summary")  
quants(dft$Age, splits = 5, "labels")[1:10]
```

queryDB

PostgreSQL Queries on Database (Read)

Description

This function lets the user query a PostgreSQL database. Previously was called `queryDummy` but was replaced and deprecated for a more general function by using the `from` parameter.

Usage

```
queryDB(query, from, creds = NA)
```

Arguments

query	Character. SQL Query
from	Character. Credential's user (see <code>get_creds()</code>)
creds	Character. Credential's directory (see <code>get_creds()</code>)

Value

`data.frame`. Result of fetching the query data.

See Also

Other Credentials: `db_download()`, `db_upload()`, `encrypt_file()`, `get_credentials()`, `get_tweets()`, `mail_send()`, `queryGA()`, `slackSend()`, `stocks_file()`

queryGA

Queries on Google Analytics

Description

This function lets the user query Google Analytics with its API. More about the documentation and parameters in `googleAnalyticsR::google_analytics()` or Google Analytics' API.

Usage

```
queryGA(
  account,
  creds = NA,
  token_dir = NA,
  metrics = "sessions",
  dimensions = "date",
  met_filters = NULL,
  dim_filters = NULL,
  start = lubridate::floor_date(Sys.Date(), "month"),
  end = Sys.Date()
)
```

Arguments

account	Character. Personal named accounts
creds	Character. Credential's user (see <code>get_creds()</code>)
token_dir	Character. Credential's directory (see <code>get_creds()</code>)
metrics	Character. Which metrics we wish to bring
dimensions	Character. Which dimensions we wish to bring
met_filters, dim_filters	A filter_clause_ga4 for filtering metrics/dimensions. Check <code>googleAnalyticsR::google_analytics()</code>
start	Date. Start date for the report
end	Date. End date for the report

Value

data.frame with the API GET request tabulated results.

See Also

Other Credentials: `db_download()`, `db_upload()`, `encrypt_file()`, `get_credentials()`, `get_tweets()`, `mail_send()`, `queryDB()`, `slackSend()`, `stocks_file()`

Other Google: `filesGD()`, `gtrends_related()`, `readGS()`

Other API: `bring_api()`, `fb_accounts()`, `fb_ads()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`, `gemini_ask()`, `gpt_ask()`, `slackSend()`

quiet	<i>Quiet prints and verbose noise</i>
-------	---------------------------------------

Description

This function silences (verbose) output prints. Thanks to Hadley Wickham for bringing the idea.

Usage

```
quiet(fx, quiet = TRUE)
```

Arguments

fx	Function to quiet
quiet	Quiet outputs? If not, skip quietness.

Value

Same as fx but with no messages or prints.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

read.file	<i>Read Files Quickly (Auto-detected)</i>
-----------	---

Description

This function lets the user import csv, xlsx, xls, sav files.

Usage

```
read.file(filename, current_wd = TRUE, sheet = 1, quiet = FALSE)
```

Arguments

filename	Character. File name to import.
current_wd	Boolean. Use current working directory before the file's name? Use this param to NOT get absolute root directory.
sheet	Character. Name or index of the sheet to read data from if file is xlsx or xls.
quiet	Boolean. Quiet summary message?

Value

List or data.frame, depending on `filename`'s data.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepm()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`, `what_size()`

readGS

*Google Sheets Reading and Writing (API v4)***Description**

Read and write data from Google Sheets knowing the file's title. You may use a single value from a cell or a data.frame from a cell range.

Usage

```
readGS(
  title,
  sheet = "Hoja 1",
  range = NULL,
  drop_nas = TRUE,
  json = NULL,
  email = NULL,
  api_key = NULL,
  server = FALSE,
  ...
)

writeGS(
  data,
  title,
  sheet = "Hoja 1",
  range = "A1",
  reformat = FALSE,
  append = FALSE,
  json = NULL,
  email = NULL,
  api_key = NULL,
  server = FALSE,
  ...
)
```

Arguments

title	Character. Title of Google Drive file. Uses regular expressions so you may fetch with patterns instead of names.
sheet	Character. Working sheet to import
range	Character. A cell range to read from
drop_nas	Boolean. Remove columns and rows that contain only NAs?
json	Character. JSON filename with service auth
email, api_key	Character. If you have multiple pre-authorized accounts in your machine, you may non-interactively select which one you wish to use by email and/or api_key.
server	Boolean. Force interacting auth process?
...	Additional parameters passed to <code>read_sheet()</code> .
data	Object (value, vector, data.frame, list).
reformat	Boolean. Reformat the affected cells?
append	Boolean.

Value

For reading, `data.frame` with the results of your Google Sheets file based on its `title`, specifically the `sheet` and `range` requested. For writing, no return value.

See Also

Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)

Other Google: [filesGD\(\)](#), [gtrends_related\(\)](#), [queryGA\(\)](#)

Description

Principal component analysis or (PCA) is a method we can use to reduce high-dimensional data to a low-dimensional space. In other words, we cannot accurately visualize high-dimensional datasets because we cannot visualize anything above 3 features. The main purpose behind PCA is to transform datasets with more than 3 features (high-dimensional) into typically a 2/3 column dataset. Despite the reduction into a lower-dimensional space we still can retain most of the variance or information from our original dataset.

Usage

```
reduce_pca(
  df,
  n = NULL,
  ignore = NULL,
  comb = c(1, 2),
  quiet = FALSE,
  plot = TRUE,
  ...
)
```

Arguments

<code>df</code>	Dataframe
<code>n</code>	Integer. Number of dimensions to reduce to.
<code>ignore</code>	Character vector. Names of columns to ignore.
<code>comb</code>	Vector. Which columns do you wish to plot? Select which two variables by name or column position.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>plot</code>	Boolean. Create plots?
<code>...</code>	Additional parameters passed to <code>stats::prcomp</code>

Value

List with reduced dataframe and possible plots.

See Also

Other Dimensionality: [reduce_tsne\(\)](#)

Other Clusters: [clusterKmeans\(\)](#), [clusterOptimalK\(\)](#), [clusterVisualK\(\)](#), [reduce_tsne\(\)](#)

Examples

```
Sys.unsetenv("LARES_FONT") # Temporal
data("iris")
df <- subset(iris, select = c(-Species))
df$id <- seq_len(nrow(df))
reduce_pca(df, n = 3, ignore = "id")
```

`reduce_tsne`

Reduce Dimensionality with t-SNE

Description

t-SNE takes high-dimensional data and reduces it to a low-dimensional graph (1-3 dimensions). Unlike PCA, t-SNE can reduce dimensions with non-linear relationships. PCA attempts to draw the best fitting line through the distribution. T-SNE calculates a similarity measure based on the distance between points instead of trying to maximize variance.

Usage

```
reduce_tsne(df, n = 2, ignore = NULL, quiet = FALSE, plot = TRUE, ...)
```

Arguments

<code>df</code>	Dataframe
<code>n</code>	Integer. Number of dimensions to reduce to.
<code>ignore</code>	Character vector. Names of columns to ignore.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>plot</code>	Boolean. Create plots?
<code>...</code>	Additional parameters passed to <code>Rtsne</code> :

Value

List with reduced dataframe and possible plots.

See Also

Other Dimensionality: [reduce_pca\(\)](#)

Other Clusters: [clusterKmeans\(\)](#), [clusterOptimalK\(\)](#), [clusterVisualK\(\)](#), [reduce_pca\(\)](#)

Examples

```
## Not run:  
data("iris")  
df <- subset(iris, select = c(-Species))  
df$id <- seq_len(nrow(df))  
reduce_tsne(df, ignore = "id", max_iter = 800, perplexity = 20)  
  
## End(Not run)
```

removenacols*Remove/Drop Columns in which ALL or SOME values are NAs***Description**

This function lets the user remove all columns that have some or all values as NAs

This function lets the user remove all rows that have some or all values as NAs

Usage

```
removenacols(df, all = TRUE, ignore = NULL)

removenarows(df, all = TRUE)

numericalonly(df, dropnacols = TRUE, logs = FALSE, natransform = NA)
```

Arguments

<code>df</code>	Data.frame
<code>all</code>	Boolean. Remove rows which contains ONLY NA values. If set to FALSE, rows which contains at least one NA will be removed
<code>ignore</code>	Character vector. Column names to ignore validation.
<code>dropnacols</code>	Boolean. Drop columns with only NA values?
<code>logs</code>	Boolean. Calculate log(x)+1 for numerical columns?
<code>natransform</code>	String. "mean" or 0 to impute NA values. If set to NA no calculation will run.

Value

`data.frame` with removed columns.

`data.frame` with removed rows.

`data.frame` with all numerical columns selected.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
data(dft) # Titanic dataset
str(dft)
numericalonly(dft) %>% head()
numericalonly(dft, natransform = "mean") %>% head()
```

remove_stopwords

Remove stop-words and patterns from character vector

Description

Remove all stop-words and specific patterns from a character vector

Usage

```
remove_stopwords(text, stop_words, exclude = NULL, sep = " ")
```

Arguments

text	Character vector
stop_words	Character vector. Words to exclude from text. Example: if you want to exclude "a", whenever that word appears it will be excluded, but when the letter "a" appears in a word, it will remain.
exclude	Character. Pattern to exclude using regex.
sep	Character. String that separate the terms.

Value

Character vector with removed texts.

See Also

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [topics_rake\(\)](#)

Examples

```
x <- c("A brown fox jumps over a dog.", "Another brown dog.")
remove_stopwords(x, stop_words = c("dog", "brown", "a"), exclude = "\\\.")
```

replaceall

*Replace Values With***Description**

This function lets the user replace all specific values in a vector or data.frame into another value. If replacing more than one value, order matters so they will be replaced in the same order that you pass them to the function. Factors will be refactored.

Usage

```
replaceall(df, original, change, which = "all", fixclass = TRUE, quiet = TRUE)
```

Arguments

<code>df</code>	Data.frame or Vector
<code>original</code>	String or Vector. Original text you wish to replace
<code>change</code>	String or Vector. Values you wish to replace the originals with
<code>which</code>	Character vector. Name of columns to use. Leave "all" for everything
<code>fixclass</code>	Boolean. Try to detect logical classes after transformations (or leave as default classes as character)?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

data.frame with replaced values based on inputs.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [remove_stopwords\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [topics_rake\(\)](#)

Examples

```
df <- data.frame(
  one = c(1:4, NA),
  two = LETTERS[1:5],
  three = rep("A", 5),
  four = c(NA, "Aaa", 123, "B", "C")
)
print(df)
```

```
replaceall(df, "A", NA)

replaceall(df, "A", "a")

replaceall(df, 1, "*")

replaceall(df, NA, "NotNA")

replaceall(df, NA, 0)

replaceall(df, c("A", "B"), c("'A'", "'B'"))

replaceall(df, "a", "*", which = "four")
```

replacefactor	<i>Replace Factor Values</i>
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Description

This function lets the user replace levels on a factor vector.

Usage

```
replacefactor(x, original, change)
```

Arguments

x	Factor (or Character) Vector
original	String or Vector. Original text you wish to replace
change	String or Vector. Values you wish to replace the originals with

Value

Factor vector with transformed levels.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
library(dplyr)
data(dft)
# Replace a single value
dft <- mutate(dft, Pclass = replacefactor(Pclass, original = "1", change = "First"))
levels(dft$Pclass)
# Replace multiple values
dft <- mutate(dft, Pclass = replacefactor(Pclass, c("2", "3"), c("Second", "Third")))
levels(dft$Pclass)
```

robyn_hypsbuilder

Robyn: Generate default hyperparameters

Description

Generate a list with hyperparameter default values, ready to be passed to Robyn::robyn_inputs().

Usage

```
robyn_hypsbuilder(
  channels,
  media_type = "default",
  adstock = "geometric",
  date_type = "weekly",
  lagged = FALSE
)
```

Arguments

channels	Character vector. Paid media and organic variables names.
media_type	Character vector. Must be length 1 or same as channels. Pick, for every channels value, what type of media it is: "online" or "offline".
adstock	Character. Pick one of: "geometric" or "weibull".
date_type	Character. Pick one of: "daily", "weekly", or "monthly". Only valid to transform thetas when using geometric adstock. Set to "skip" in case you wish to leave default weekly values.
lagged	Boolean vector. Must be length 1 or same as channels. Pick, for every channels value, if you wish to have a lagged effect. Only valid for Weibull adstock.

Value

list with default hyperparameters ranges.

See Also

Other Robyn: [robyn_marginal\(\)](#), [robyn_modelselector\(\)](#), [robyn_performance\(\)](#)

Examples

```
robyn_hypsbuilder(  
  channels = c(  
    "branded_search_spend",  
    "nonbranded_search_spend",  
    "print_spend",  
    "ooh_spend",  
    "tv_spend",  
    "radio_spend"  
  media_type = c(  
    "online", "online", "offline",  
    "offline", "offline", "offline"  
  adstock = "geometric",  
  date_type = "weekly"  
)
```

robyn_marginal

Robyn: Marginal Performance (mROAS & mCPA) [Experimental]

Description

Calculate and plot marginal performance of any spend or organic variable.

Usage

```
robyn_marginal(..., marginal_unit = 1)
```

Arguments

... Additional parameters.
marginal_unit Additional units to calculate the marginal performance.

Value

list with base and marginal response results, marginal performance metric and value, and plot.

See Also

Other Robyn: [robyn_hypsbuilder\(\)](#), [robyn_modelselector\(\)](#), [robyn_performance\(\)](#)

Examples

```
## Not run:
# You may load an exported model to recreate Robyn objects
mod <- Robyn::robyn_recreate(json_file = "your_model.json")
robyn_marginal(
  InputCollect = mod$InputCollect,
  OutputCollect = mod$OutputCollect,
  metric_name = "emails_0",
  metric_value = 100000,
  date_range = "all",
  marginal_unit = 10000000
)
## End(Not run)
```

robyn_modelselector *Robyn: Model Selection by Weighted Criteria Scores*

Description

Consider N best models to select the right ones to study using several criteria/metrics such as potential improvement on budget allocator, how many non-zero coefficients there are, R squared, historical performance, baseline expectation, etc.

Read more about this functionality in Medium post: [here](#).

Usage

```
robyn_modelselector(
  InputCollect,
  OutputCollect,
  metrics = c("rsq_train", "performance", "potential_improvement", "non_zeroes",
             "incluster_models", "cluster_sd", "certainty", "baseline_dist"),
  wt = c(2, 0.1, 0, 1, 0.1, 0, 1.5, 0),
  baseline_ref = 0,
  top = 4,
  n_per_cluster = 5,
  allocator_limits = c(0.5, 2),
  quiet = FALSE,
  cache = TRUE,
  ...
)
## S3 method for class 'robyn_modelselector'
plot(x, ...)
```

Arguments

<code>InputCollect, OutputCollect</code>	Robyn output objects.
<code>metrics</code>	Character vector. Which metrics do you want to consider? Pick any combination from: "rsq_train" for trained R squared, "performance" for ROAS or (inverse) CPA, "potential_improvement" for default budget allocator improvement using <code>allocator_limits</code> , "non_zeroes" for non-zero beta coefficients, "incluster_models" for amount of models per cluster, "baseline_dist" for the difference between the model's baseline and <code>baseline_ref</code> value, "certainty" metric to minimize the channels' distance to their cluster's mean performance, weighted by spends if <code>spend_wt = TRUE</code> , "cluster_sd" metric to score based on the paid channels' performance standard deviations in clusters. Additionally, you can use the standard MOO errors: "nrmse", "decomp.rssd", and "mape" (the lowest the error, the highest the score; same for "baseline_dist" and "cluster_sd").
<code>wt</code>	Vector. Weight for each of the normalized <code>metrics</code> selected, to calculate the score and rank models. Must have the same order and length of <code>metrics</code> parameter input.
<code>baseline_ref</code>	Numeric value. Between 0 and 1. What is the baseline percentage you expect? Baseline in this case are all the sales or conversions from non-media channels (organic & paid). Use with "baseline_dist" metric.
<code>top</code>	Integer. How many ranked models to star? The better the model is, the more stars it will have marked.
<code>n_per_cluster</code>	Integer. How many models per cluster do you want to plot? Default: 5. Keep in mind they will all be considered for the calculations.
<code>allocator_limits</code>	Numeric vector, length 2. How flexible do you want to be with the budget allocator? By default, we'll consider a 0.5X and 2X range to let the budget shift across channels.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>cache</code>	Use cache functionality for allocator's results?
<code>...</code>	Additional parameters.
<code>x</code>	robyn_modelselector object

Value

list with resulting ranked data.frames, weights and plot.

See Also

Other Robyn: [robyn_hypsbuilder\(\)](#), [robyn_marginal\(\)](#), [robyn_performance\(\)](#)

robyn_performance*Robyn: Dynamic Performance and Contribution Report*

Description

Given a date range, calculate specific and total performance and contribution for each of your marketing and non-marketing channels.

Usage

```
robyn_performance(
  InputCollect,
  OutputCollect,
  start_date = NULL,
  end_date = NULL,
  solID = NULL,
  totals = TRUE,
  non_promo = FALSE,
  marginals = FALSE,
  carryovers = FALSE,
  new_version = FALSE,
  quiet = FALSE,
  ...
)
```

Arguments

InputCollect, OutputCollect	Robyn output objects.
start_date, end_date	Date. Start and end date to filter the data to be reported.
solID	Character. Single ID of the model to report. If there's only one available in OutputCollect, no need to define.
totals	Boolean. Add total rows. This includes summary rows (promotional which is paid and organic channels, baseline, grand total).
non_promo	Boolean. Add non-promotional responses as well?
marginals	Boolean. Include mROAS or mCPA marginal performance metric as an additional column called "marginal". Calculations are based on mean spend and mean response with mean carryover results, between start_date and end_date.
carryovers	Boolean. Add mean percentage of carryover response for date range between start_date and end_date on paid channels. Keep in mind organic variables also have carryover but currently not showing.
new_version	Boolean. Use dev version's new function for marginal calculations (if available)?
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.

Value

`data.frame` with results on ROAS/CPA, spend, response, contribution per channel, with or without total rows.

See Also

Other Robyn: [robyn_hypsbuilder\(\)](#), [robyn_marginal\(\)](#), [robyn_modelselector\(\)](#)

Examples

```
## Not run:
# You may load an exported model to recreate Robyn objects
mod <- Robyn::robyn_recreate(json_file = "your_model.json")
robyn_performance(mod$InputCollect, mod$OutputCollect)

## End(Not run)
```

Description

Given a list of recreated Robyn models, this function optimizes budget allocation across MMM with respective constraints by maximizing incremental revenue/conversions. This method assumes each model is independent, that can be compared given its spends were cleanly and properly split, they modeled the same metric (revenue or conversion) and units (currency or type of conversion), and date granularity. For best results, ensure channels have similar granularity across markets to simplify interpretation and application of the outputs.

Given a list of recreated Robyn models, this function optimizes budget allocation across MMM with respective constraints by maximizing response across all channels. This method assumes each model is independent, that can be compared given its spends were cleanly and properly split, they modeled the same metric (revenue or conversion) and units (currency or type of conversion), and date granularity. Recommended to have same channels granularity across markets to simplify results readings and application.

Usage

```
robyn_xmodels(
  models,
  initial_budgets = NULL,
  start_dates = NULL,
  end_dates = NULL,
  budget_constr_low = 0.5,
  budget_constr_up = 1.5,
  channel_constr_low = budget_constr_low,
  channel_constr_up = budget_constr_up,
  cores = NULL,
```

```

quiet = FALSE,
...
)

## S3 method for class 'robyn_crossmmm'
print(x, ...)

robyn_xchannels(
  models,
  initial_budgets = NULL,
  start_dates = NULL,
  end_dates = NULL,
  channel_constr_low = 0.5,
  channel_constr_up = 2,
  quiet = FALSE,
  ...
)

```

Arguments

<code>models</code>	Lists. Recreated Robyn models with <code>robyn_recreate()</code> .
<code>initial_budgets</code>	Numeric vector. Default will use the total spends per model for the specified or total date range. Must be length 1 or same as <code>models</code> .
<code>start_dates, end_dates</code>	Character vector. Start and end dates for each specific model. You can specify a single date and will be used in all models. Default empty value will assume you want all available data and date range. Must be length 1 or same as <code>models</code> .
<code>budget_constr_low, budget_constr_up</code>	Numeric vector. Relative minimum and maximum budgets to consider based on <code>initial_budgets</code> . By default it'll consider 50 Must be length 1 or same as <code>models</code> .
<code>channel_constr_low, channel_constr_up</code>	Numeric vector. Relative lower and upper constraints per channel compared with mean spend during the time period defined. If mean was zero for date range, historical mean spend value will be used. Must have length 1 to replicate for all channels or same length (and order) as <code>paid_media_spends</code> .
<code>cores</code>	Integer. How many cores to use for parallel computations? Set to 1 to not use this option. Default will be the minimum between 10 cores and all available cores - 1.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters to be passed to internal functions.
<code>x</code>	Object to print: <code>robyn_crossmmm()</code> output.

Details

This approach is faster and cleaner compared with previous proposal using `robyn_xmodels()`.

Value

Invisible vector with results by letter.
List. Contains optimized allocation results and plots.

Examples

```
# You must have Robyn installed and some models stored as JSON files
## Not run:
# Import and recreate the models
files <- c("BrandA.json", "BrandB.json", "BrandC.json", "BrandN.json")
models <- lapply(files, function(x) Robyn::robyn_recreate(x))
names(models) <- gsub("\\.json", "", files)

# Calculate cross-brand optimal allocation
res <- robyn_xmodels(
  models,
  cores = 10,
  start_dates = "2023-01-01",
  end_dates = "2023-12-01"
)
print(res)
res$summary

## End(Not run)
## Not run:
# Calculate cross-brand optimal allocation
res <- robyn_xchannels(
  models,
  start_dates = "2023-01-01",
  end_dates = "2023-12-01"
)
## End(Not run)
```

Description

This function calculates ROC Curves and AUC values with 95% confidence range. It also works for multi-categorical models.

Usage

```
ROC(tag, score, multis = NA)
```

Arguments

<code>tag</code>	Vector. Real known label
<code>score</code>	Vector. Predicted value or model's result
<code>multis</code>	Data.frame. Containing columns with each category score (only used when more than 2 categories coexist)

Value

List with ROC's results, area under the curve (AUC) and their CI.

Plot Results

To plot results, use the `mpplot_roc()` function.

See Also

Other Machine Learning: `conf_mat()`, `export_results()`, `gain_lift()`, `h2o_automl()`, `h2o_predict_MOJO()`, `h2o_selectmodel()`, `impute()`, `iter_seeds()`, `lasso_vars()`, `model_metrics()`, `model_preprocess()`, `msplit()`

Other Model metrics: `conf_mat()`, `errors()`, `gain_lift()`, `loglossBinary()`, `model_metrics()`

Examples

```
data(dfr) # Results for AutoML Predictions
lapply(dfr[c(1, 2)], head)

# ROC Data for Binomial Model
roc1 <- ROC(dfr$class2$tag, dfr$class2$scores)
lapply(roc1, head)

# ROC Data for Multi-Categorical Model
roc2 <- ROC(dfr$class3$tag, dfr$class3$score,
             multis = subset(dfr$class3, select = -c(tag, score)))
)
lapply(roc2, head)
```

Description

Generative Art: Sphere XmodY

Usage

```
rtistry_sphere(eye = c(100, 0, 0), pal = "auto", var = 3)
```

Arguments

eye, pal, var Parameters to change aesthetics and calculations

Value

ggplot object

scale_x_comma *Axis scales format*

Description

The _comma ones set comma format for axis text, the _percent ones set percent format for axis text, _dollar for dollar currency, and _abbr for abbreviated format. Lastly, use _formatNum to further customize your numerical scales with formatNum.

Usage

```
scale_x_comma(...)

scale_y_comma(...)

scale_x_percent(...)

scale_y_percent(...)

scale_x_dollar(...)

scale_y_dollar(...)

scale_x_abbr(...)

scale_y_abbr(...)

scale_x_formatNum(
  ...,
  decimals = 2,
  signif = NULL,
  type = Sys.getenv("LARES_NUMFORMAT"),
  pre = "",
  pos = "",
  sign = FALSE,
  abbr = FALSE
)

scale_y_formatNum(
  ...,
```

```

decimals = 2,
signif = NULL,
type = Sys.getenv("LARES_NUMFORMAT"),
pre = "",
pos = "",
sign = FALSE,
abbr = FALSE
)

```

Arguments

...	Arguments passed to ggplot2::continuous_scale or formatNum depending on the function.
decimals	Integer. Amount of decimals to display. If set to NULL, then getOption("digits") will be used.
signif	Integer. Rounds the values in its first argument to the specified number of significant digits.
type	Integer. 1 for International standards. 2 for American Standards. Use Sys.setenv("LARES_NUMFORMAT" = 2) to set this parameter globally.
pre, pos	Character. Add string before or after number.
sign	Boolean. Add + sign to positive values.
abbr	Boolean. Abbreviate using num_abbr()? You can use the 'decimals' parameter to set abbr's n(-1) parameter.

Value

Reformatted scales on ggplot2 object

Examples

```

library(ggplot2)
df <- ggplot2::txhousing %>% removenarows(all = FALSE)

ggplot(df, aes(x = sales, y = volume)) +
  geom_point() +
  scale_x_dollar() +
  scale_y_abbr()

# Use any argument from scale_x/y_continuous
ggplot(df, aes(x = listings, y = log(inventory))) +
  geom_point() +
  scale_x_comma() +
  scale_y_percent(limits = c(0, 3))

# Use any argument from scale_x/y_continuous AND formatNum
ggplot(df, aes(x = median, y = inventory)) +
  geom_point() +
  scale_x_formatNum(n.breaks = 3, pre = "@", abbr = TRUE) +
  scale_y_formatNum(position = "right", decimals = 0, pos = " X")

```

scrabble_dictionary *Scrabble: Dictionaries*

Description

Download words from 4 different languages: English, Spanish, German, and French. Words will be saved into the `temp` directory. This is an auxiliary function. You may want to use `scrabble_words` directly if you are searching for the highest score words!

Get score for any word or list of words. You may set manually depending on the rules and languages you are playing with. Check the examples for Spanish and English values when I played Words With Friends.

Dataframe for every letter and points given a language.

Find highest score words given a set of letters, rules, and language to win at Scrabble! You just have to find the best place to post your tiles.

Usage

```
scrabble_dictionary(lang_dic, quiet = FALSE)

scrabble_score(words, scores.df)

scrabble_points(lang)

scrabble_words(
  tiles = "",
  free = 0,
  force_start = "",
  force_end = "",
  force_str = "",
  force_exclude = "",
  exclude_here = "",
  force_n = 0,
  force_max = 0,
  pattern = "",
  repeated = FALSE,
  language = Sys.getenv("LARES_LANG"),
  scores = language,
  words = NULL,
  quiet = FALSE,
  print = TRUE
)
```

Arguments

`lang_dic` Character. Any of "en","es","de","fr". Set to NULL if you wish to skip this step (and use `words` parameter in `scrabble_words` instead).

<code>quiet</code>	Boolean. Do not print words as they are being searched.
<code>words</code>	Character vector. Use if you wish to manually add words.
<code>scores.df</code>	Dataframe. Must contain two columns: "tiles" with every letter of the alphabet and "scores" for each letter's score.
<code>lang</code>	Character. Any of "en", "es" or "chars". Set to NULL if you wish to skip this step (and use <code>words</code> parameter in <code>scrabble_words()</code> instead). The "chars" parameter will score the number of characters a word has.
<code>tiles</code>	Character. The letters you wish to consider.
<code>free</code>	Integer. How many free blank tiles you have?
<code>force_start, force_end</code>	Character. Force words to start or end with a pattern of letters and position. Examples: "S" or "SO" or "__S_O"... If the string contains tiles that were not specified in <code>tiles</code> , they will automatically be included.
<code>force_str</code>	Character vector. Force words to contain strings. If the string contains tiles that were not specified in <code>tiles</code> , they will automatically be included.
<code>force_exclude, exclude_here</code>	Character vector. Exclude words containing these tiles (and positions). Not very relevant on Scrabble but for Wordle.
<code>force_n, force_max</code>	Integer. Force words to be n or max n characters long. Leave 0 to ignore parameter.
<code>pattern</code>	Character string. Custom regex patterns you'd like to match.
<code>repeated</code>	Boolean. By default, no replacement allowed. When activated, a single tile can be repeated and won't be "used and discarded".
<code>scores, language</code>	Character. Any of "en", "es", "de", "fr". If <code>scores</code> is not any of those languages, must be a data.frame that contains two columns: "tiles" with every letter of the alphabet and "scores" for each letter's score. If you wish to overwrite or complement this dictionary other words you can set to "none" and/or use the <code>words</code> parameter. You might also want to set this parameter globally with <code>Sys.setenv("LARES_LANG" = "en")</code> and forget about it!
<code>print</code>	Boolean. Print how many words are left by step.

Value

- data.frame with words and language columns.
- data.frame with word, scores, and length values for each word.
- data.frame with tiles and scores for each alphabet letter.
- data.frame with matching words found, sorted by higher points.

Examples

```
if (haveInternet()) {
  # For Spanish words (default)
  es_scores <- scrabble_points("es")
```

```

# Custom scores for each letter
cu_scores <- data.frame(
  tiles = tolower(LETTERS),
  scores = c(
    1, 1, 1, 1, 1, 1, 1, 5, 1, 1, 5, 2, 4, 2, 1,
    4, 10, 1, 1, 1, 2, 5, 4, 8, 3, 10
  )
)

# Score values for each set of rules
words <- c("Bernardo", "Whiskey", "R is great")
scrabble_score(words, es_scores)
scrabble_score(words, cu_scores)
}

scrabble_points("es")
scrabble_points("en")
# Not yet available
scrabble_points("fr")

if (haveInternet()) {
  # Automatic use of languages and scores
  Sys.setenv("LARES_LANG" = "es")
  scrabble_words(
    tiles = "hola",
    free = 2,
    force_start = "h",
    force_n = 4,
    force_str = "_o_a",
    exclude_here = "__z|j"
  )

  wordle <- c("board", "tempo", "shoes", "hoard")
  scrabble_words(
    language = NULL,
    words = wordle,
    force_n = 5,
    force_str = "O_R"
  )
}

# Words considered for a language (you can custom it too!)
es_words <- scrabble_dictionary("es")
}

```

Description

This function searches for relevant words in a given text and adds sentiments labels (joy, anticipation, surprise, positive, trust, anger, sadness, fear, negative, disgust) for each of them, using NRC.

Then, makes a summary for all words and plot results.

Usage

```
sentimentBreakdown(
  text,
  lang = "spanish",
  exclude = c("maduro", "que"),
  append_file = NA,
  append_words = NA,
  plot = TRUE,
  subtitle = NA
)
```

Arguments

<code>text</code>	Character vector
<code>lang</code>	Character. Language in text (used for stop words)
<code>exclude</code>	Character vector. Which word do you wish to exclude?
<code>append_file</code>	Character. Add a dictionary to append. This file must contain at least two columns, first with words and second with the sentiment (consider sentiments on description).
<code>append_words</code>	Dataframe. Same as <code>append_file</code> but appending data frame with word and sentiment directly
<code>plot</code>	Boolean. Plot results summary?
<code>subtitle</code>	Character. Add subtitle to the plot

Value

List. Contains data.frame with words and sentiments, summary and plot.

See Also

Other Text Mining: `cleanText()`, `ngrams()`, `remove_stopwords()`, `replaceall()`, `textCloud()`, `textFeats()`, `textTokenizer()`, `topics_rake()`

Description

Generate a sequence of numbers that determines the order in which surnames should be listed based on the number of generations of ancestors you wish to include. This sequence follows the traditional Latin custom of assigning the father's surname first, followed by the mother's surname. The same logic extends systematically to higher generations, ensuring that the order of surnames remains consistent as you move upward through the family tree.

Usage

```
seq_surnames(n = 1)
```

Arguments

- n Integer. Number of generations to include in the sequence. Notice it will generate a vector with $2^{(n-1)}$ values.

Value

Integer vector.

Examples

```
seq_surnames(1)  
seq_surnames(2)  
seq_surnames(3)  
seq_surnames(4)
```

shap_var

SHAP-based dependence plots for categorical/numerical features (PDP)

Description

Having a h2o_shap object, plot a dependence plot for any categorical or numerical feature.

Usage

```
shap_var(x, var, keep_outliers = FALSE)
```

Arguments

- x h2o_shap object
var Variable name
keep_outliers Boolean. Outliers detected with z-score and 3sd may be suppress or kept in your plot. Keep them?

Value

ggplot2 objct with shap values plotted

See Also

Other SHAP: [h2o_shap\(\)](#)

Examples

```

## Not run:
# Train a h2o_automl model
model <- h2o_automl(dft, Survived,
  max_models = 1, target = TRUE,
  ignore = c("Ticket", "Cabin", "PassengerId"),
  quiet = TRUE
)

# Calculate SHAP values
SHAP_values <- h2o_shap(model)
# Equivalent to:
# SHAP_values <- h2o_shap(
#   model = model$model,
#   test = model$datasets$test,
#   scores = model$scores_test$scores)

# Check SHAP results
head(SHAP_values)

# You must have "ggbeeswarm" library to use this auxiliary function:
# Plot SHAP values (feature importance)
plot(SHAP_values)

# Plot some of the variables (categorical)
shap_var(SHAP_values, Pclass)

# Plot some of the variables (numerical)
shap_var(SHAP_values, Fare)

## End(Not run)

```

slackSend

Send Slack Message (Webhook)

Description

This function send a Slack message using its Webhooks.

Usage

```
slackSend(text, title = "", pretext = "", hook = NA, creds = NA)
```

Arguments

text, title, pretext

Character. Content on you Slack message.

hook

Character. Web hook URL. Ths value will be overwritten by creds if correctly used.

creds	Character. Credential's dir (see <code>get_creds()</code>). Set hook URL into the "slack" list in your YML file. Will use first value.
-------	---

Details

For more help, you can follow the [Sending messages using Incoming Webhooks](#) original documentation.

Value

Invisible POST response

See Also

Other API: `bring_api()`, `fb_accounts()`, `fb_ads()`, `fb_creatives()`, `fb_insights()`, `fb_process()`, `fb_report_check()`, `fb_rf()`, `fb_token()`, `gemini_ask()`, `gpt_ask()`, `queryGA()`

Other Credentials: `db_download()`, `db_upload()`, `encrypt_file()`, `get_credentials()`, `get_tweets()`, `mail_send()`, `queryDB()`, `queryGA()`, `stocks_file()`

Examples

```
## Not run:  
slackSend(text = "This is a message", title = "TEST", pretext = Sys.info()["user"])  
  
## End(Not run)
```

Description

`splot_summary()` plots a summary for the whole portfolio, showing how much have you invested, how much has each ticker changed, etc.

`splot_change()` plots each stock's change through history, since inception, with weighted attributions or absolute values.

`splot_growth()` plots your portfolio's growth, in cash and investment, since inception.

`stocks_plots()` plots a portfolio's historical dividends incomes grouped by quarter an year.

`splot_roi()` plots a portfolio's historical ROI since inception or since last n days, with 2 moving average lines.

`splot_types()` lets the user plot types or categories of tickers.

`splot_etf()` lets the user plot his portfolio's distribution, specifically ETF's sectors.

Usage

```

splot_summary(p, s, save = FALSE)

splot_change(
  p,
  s,
  rel = TRUE,
  group = FALSE,
  n_days = 365,
  keep_old = FALSE,
  save = FALSE
)

splot_growth(p, save = FALSE)

splot_divs(p, type = 1)

splot_roi(p, n_days = 365, historical = TRUE, ma = c(12, 50), save = FALSE)

splot_types(s, save = FALSE)

splot_etf(s, keep_all = FALSE, cache = TRUE, save = FALSE)

```

Arguments

p	Dataframe. Result from <code>daily_portfolio()</code>
s	Dataframe. Result from <code>daily_stocks()</code>
save	Boolean. Save plot into a local file?
rel	Boolean. Relative delta values (weighted with portfolio)? If not, absolute monetary delta values.
group	Boolean. Group stocks by stocks type?
n_days	Integer. How many days back you want to see?
keep_old	Boolean. Include sold tickers even though not currently in portfolio?
type	Integer. Typo of plot. 1 for incomes.
historical	Boolean. Historical ROI metric? If not, ROI will be calculated locally for n_days parameter
ma	Numeric Vector. Select 2 values for moving averages. Set to NA to turn this metric off
keep_all	Boolean. Keep "Not Known / Not ETF"?
cache	Boolean. Use daily cache if available?

Value

ggplot object

See Also

Other Investment: [etf_sector\(\)](#), [stocks_file\(\)](#), [stocks_quote\(\)](#)

Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [stocks_quote\(\)](#)

spread_list

Spread list column into new columns

Description

Spread an existing list column into new columns on a data.frame. Note that every element on every observation must have a name for the function to do its work. Original column will be automatically suppressed but you can set the `replace` argument to avoid it.

Usage

```
spread_list(df, col, str = NULL, replace = TRUE)
```

Arguments

df	Dataframe
col	Variable name.
str	Character. Start column names with. If set to NULL, original name of column will be used.
replace	Boolean. Replace original values (delete column)

Value

data.frame. Result of un-nesting named or un-named list columns.

Examples

```
df <- dplyr::starwars
# Un-named list columns
spread_list(df, films, replace = FALSE) %>%
  dplyr::select(name, dplyr::starts_with("films")) %>%
  head(8)
# Named (and un-named) list columns
df <- dplyr::tibble(id = 1:3, platform = list(
  list("fb" = 1, "ig" = 2),
  list("fb" = 3),
  list()
))
spread_list(df, platform, str = "ptf_")
```

statusbar*Progressive Status Bar (Loading)*

Description

This function lets the user view a progressbar for a 'for' loop.

Usage

```
statusbar(
  run = 1,
  max.run = 100,
  label = run,
  msg = "",
  type = Sys.getenv("LARES_STATUSBAR"),
  start_time = NA,
  multiples = 1,
  alarm = FALSE
)
```

Arguments

<code>run</code>	Iterator. for loop or an integer with the current loop number. Start with 1 preferably
<code>max.run</code>	Number. Maximum number of loops
<code>label</code>	String. With additionaly information to be printed at the end of the line. The default is <code>run</code> .
<code>msg</code>	Character. Finish message.
<code>type</code>	Character. Loading type style: equal, domino, sword, filled.
<code>start_time</code>	POSIXct. Start time to consider. If NA, then when first iteration starts will be set as start time. Useful for when first iteration is showed as done but started a few seconds/minutes ago.
<code>multiples</code>	Integer. Only print when multiples of N (to avoid) wasting resources on fast and lots of iterations.
<code>alarm</code>	Boolean. Ping (sound) when done. Requires beepr.

Value

No return value, called for side effects.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepml\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#),

```
importxlsx(), ip_data(), json2vector(), list_cats(), listfiles(), mail_send(), markdown2df(),
move_files(), msplit(), myip(), quiet(), read.file(), tic(), try_require(), updateLares(),
warnifnot(), what_size()
```

Examples

```
for (i in 1:9) {
  statusbar(i, 9, multiples = 2)
  Sys.sleep(0.3)
}
```

stocks_file

Build a Personal Investing Portfolio Report

Description

stocks_file() lets the user download his personal Excel with his Portfolio's data, locally or from Dropbox.

daily_stocks() creates a dataframe with all relevant metrics and values, for each ticker or symbol, for every day since inception.

daily_portfolio() creates a data.frame with all relevant metrics and values, for the overall portfolio, for every day since inception.

stocks_obj() lets the user create his portfolio's calculations and plots for further study.

stocks_report() lets the user create his portfolio's full report with plots and send it to an email with the HTML report attached

Usage

```
stocks_file(
  file = NA,
  creds = NA,
  auto = TRUE,
  sheets = c("Portafolio", "Fondos", "Transacciones"),
  keep_old = TRUE,
  cache = TRUE,
  quiet = FALSE,
  ...
)

daily_stocks(hist, trans, tickers = NA, window = "MAX", ...)

daily_portfolio(hist, trans, cash, cash_fix = 0, window = "MAX")

stocks_obj(
  data = stocks_file(),
  cash_fix = 0,
```

```

tax = 30,
sectors = FALSE,
parg = FALSE,
window = c("1M", "YTD", "1Y", "MAX"),
cache = TRUE,
quiet = FALSE
)

stocks_report(
  data = NA,
  keep_old = TRUE,
  dir = NA,
  mail = FALSE,
  attachment = TRUE,
  to = "laresbernardo@gmail.com",
  sectors = FALSE,
  keep = FALSE,
  creds = NA,
  cache = TRUE
)

```

Arguments

<code>file</code>	Character. Import an Excel file, local or from URL.
<code>creds</code>	Character. Credential's user (see <code>get_creds()</code>) for sending mail and Dropbox interaction.
<code>auto</code>	Boolean. Automatically use my local personal file? You might want to set in into your <code>.Renvironment</code> <code>LARES_PORTFOLIO=~/dir/to/your/file.xlsx</code> so you can leave all other parameters as NA and use it every time.
<code>sheets</code>	Character Vector. Names of each sheet containing Portfolio summary, Cash, and Transactions information. Please, keep the order of these tabs.
<code>keep_old</code>	Boolean. Include sold tickers even though not currently in portfolio?
<code>cache</code>	Boolean. Use daily cache if available?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>...</code>	Additional parameters.
<code>hist</code>	Dataframe. Result from <code>stocks_hist()</code>
<code>trans</code>	Dataframe. Result from <code>stocks_file()\$transactions</code>
<code>tickers</code>	Dataframe. Result from <code>stocks_file()\$portfolio</code>
<code>window</code>	Character. Choose any of: "1W", "1M", "6M", "1Y", "YTD", "5Y", "MAX"
<code>cash</code>	Dataframe. Result from <code>stocks_file()\$cash</code>
<code>cash_fix</code>	Numeric. If, for some reason, you need to fix your cash amount for all reports, set the amount here
<code>data</code>	Character. <code>stocks_obj()</code> output. If NA, automatic parameters and <code>stocks_file()</code> defaults will be used.

tax	Numeric. How much [0-99] of your dividends are gone with taxes?
sectors	Boolean. Return sectors segmentation for ETFs?
parg	Boolean. Personal argument. Used to personalize stuff, in this case, taxes changed from A to B in given date (hard-coded)
dir	Character. Directory for HTML report output. If set to NA, current working directory will be used. If mail sent, file will be erased
mail	Boolean. Do you want to send an email with the report attached? If not, an HTML file will be created in dir
attachment	Boolean. Create and add report as attachment if mail=TRUE? If not, no report will be rendered and only tabulated summaries will be included on email's body.
to	Character. Email to send the report to
keep	Boolean. Keep HTML file when sent by email?

Value

List with portfolio, transactions, and cash data.frames.
 data.frame. Processed at date and symbol level.
 data.frame. Processed at date and portfolio level.
 List. Aggregated results and plots.
 Invisible list. Aggregated results and plots.

See Also

Other Investment: [etf_sector\(\)](#), [splot_summary\(\)](#), [stocks_quote\(\)](#)
 Other Credentials: [db_download\(\)](#), [db_upload\(\)](#), [encrypt_file\(\)](#), [get_credentials\(\)](#), [get_tweets\(\)](#),
[mail_send\(\)](#), [queryDB\(\)](#), [queryGA\(\)](#), [slackSend\(\)](#)

Examples

```
## Not run:
# Load lares dummy portfolio XLSX
file <- system.file("inst/docs", "dummyPortfolio.xlsx", package = "lares")
df <- stocks_file(
  file = file,
  sheets = c("Portafolio", "Fondos", "Transacciones"),
  keep_old = FALSE
)

## End(Not run)
## Not run:
list <- stocks_obj()
stocks_report(list, dir = "~/Desktop")

## End(Not run)
```

`stocks_quote`*Download Stocks Historical and Current Values*

Description

`stocks_quote()` lets the user download stocks live data.
`stocks_hist()` lets the user download stocks historical data.

Usage

```
stocks_quote(symbols, ...)

stocks_hist(
  symbols = c("VTI", "META"),
  from = Sys.Date() - 365,
  to = Sys.Date(),
  today = TRUE,
  tax = 15,
  parg = FALSE,
  cache = TRUE,
  quiet = FALSE,
  ...
)

## S3 method for class 'stocks_hist'
plot(x, type = 1, ...)
```

Arguments

<code>symbols</code>	Character Vector. List of symbols to download historical data.
<code>...</code>	Additional parameters.
<code>from, to</code>	Date. Dates for range. If not set, 1 year will be downloaded. Do use more than 4 days or will be over-written.
<code>today</code>	Boolean. Do you wish to add today's live quote? This will happen only if to value is the same as today's date
<code>tax</code>	Numeric. How much [0-99] of your dividends are gone with taxes?
<code>parg</code>	Boolean. Personal argument. Used to personalize stuff, in this case, taxes changed from A to B in given date (hard-coded)
<code>cache</code>	Boolean. Use daily cache if available?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>x</code>	<code>stocks_hist</code> object
<code>type</code>	Integer. Select type of plot.

Value

data.frame with Symbol, Type of stock, Quote time, current value, Daily Change, Market, and Symbol Name.

See Also

Other Investment: [etf_sector\(\)](#), [splot_summary\(\)](#), [stocks_file\(\)](#)

Other Scrapper: [filesGD\(\)](#), [get_mp3\(\)](#), [gtrends_related\(\)](#), [holidays\(\)](#), [ip_data\(\)](#), [readGS\(\)](#), [splot_summary\(\)](#)

Examples

```
## Not run:
# Multiple quotes at the same time
stocks_quote(c("VTI", "VOO", "TSLA"))

## End(Not run)
## Not run:
df <- stocks_hist(symbols = c("VTI", "META", "FIW"), from = Sys.Date() - 180)
print(head(df))
plot(df)

## End(Not run)
```

Description

Solve a Sudoku puzzle, where empty values are represented by 0s into a matrix object.

Usage

```
sudoku_solver(board, needed_cells = NULL, index = 1, quiet = FALSE)
```

Arguments

board	Matrix. 9x9 matrix or vector length 81, with only digits from 0 to 9.
needed_cells, index	Auxiliary parameters to auto-iterate using this same fx.
quiet	Boolean. Keep quiet? If not, plot results.

Value

Logical output answering of the input board can be solved. The actual solved solution will be created as an object named solved in your .GlobalEnv.

Examples

```
# board <- c(0,0,0,0,0,6,0,0,0,
#           0,9,5,7,0,0,3,0,0,
#           4,0,0,0,9,2,0,0,5,
#           7,6,4,0,0,0,0,0,3,
#           0,0,0,0,0,0,0,0,0,
#           2,0,0,0,0,0,9,7,1,
#           5,0,0,2,1,0,0,0,9,
#           0,0,7,0,0,5,4,8,0,
#           0,0,0,8,0,0,0,0,0)
# sudoku_solver(board)

# Trivial input (everything)
trivial <- matrix(rep(0, 81), byrow = TRUE, ncol = 9)
trivial
sudoku_solver(trivial)

# Wrong / Impossible to solve input
imp <- matrix(c(rep(1, 72), rep(0, 9)), byrow = TRUE, ncol = 9)
imp
sudoku_solver(imp)
```

target_set

Set Target Value in Target Variable

Description

This function detects or forces the target value when predicting a categorical binary model. This is an auxiliary function.

Usage

```
target_set(tag, score, target = "auto", quiet = FALSE)
```

Arguments

tag	Vector. Real known label
score	Vector. Predicted value or model's result
target	Value. Which is your target positive value? If set to 'auto', the target with largest mean(score) will be selected. Change the value to overwrite. Only used when binary categorical model.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.

Value

List. Contains original data.frame df and which with the target variable.

textCloud*Wordcloud Plot*

Description

Study the distribution of a target variable vs another variable. This function is quite similar to the funModeling's corrplot function.

Usage

```
textCloud(  
  text,  
  lang = "english",  
  exclude = NULL,  
  seed = 0,  
  keep_spaces = FALSE,  
  min = 2,  
  pal = NA,  
  print = TRUE  
)
```

Arguments

text	Character vector
lang	Character. Language in text (used for stop words)
exclude	Character vector. Which word do you wish to exclude?
seed	Numeric. Seed for re-producible plots
keep_spaces	Boolean. If you wish to keep spaces in each line to keep unique compound words, separated with spaces, set to TRUE. For example, 'LA ALAMEDA' will be set as 'LA_ALAMEDA' and treated as a single word.
min	Integer. Words with less frequency will not be plotted
pal	Character vector. Which colours do you wish to use
print	Boolean. Plot results as textcloud?

Value

wordcloud plot object

See Also

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [remove_stopwords\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [topics_rake\(\)](#)

textFeats	<i>Create features out of text</i>
-----------	------------------------------------

Description

This function creates a data.frame with features based on a text vector

Usage

```
textFeats(text, auto = TRUE, contains = NA, prc = FALSE)
```

Arguments

<code>text</code>	Character vector
<code>auto</code>	Boolean. Auto create some useful parameters?
<code>contains</code>	Character vector. Which columns do you wish to add with a contains (counter) string validator?
<code>prc</code>	Boolean. Also add percentage of each column compared with length?

Value

data.frame with additional features based on `text`.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [remove_stopwords\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textTokenizer\(\)](#), [topics_rake\(\)](#)

Examples

```
textFeats("Bernardo Lares")
textFeats("Bernardo Lares 123!", prc = TRUE)
textFeats("I'm 100% Lares...", contains = c("Lares", "lares"))
textFeats(c("GREAT library!!", "Have you tried this 2?", "Happy faces :D :-"))
```

textTokenizer *Tokenize Vectors into Words*

Description

This function transforms texts into words, calculate frequencies, supress stop words in a given language.

Usage

```
textTokenizer(  
  text,  
  exclude = NULL,  
  lang = NULL,  
  min_word_freq = 5,  
  min_word_len = 2,  
  keep_spaces = FALSE,  
  lowercase = TRUE,  
  remove_numbers = TRUE,  
  remove_punct = TRUE,  
  remove_leett = TRUE,  
  laughs = TRUE,  
  utf = TRUE,  
  df = FALSE,  
  h2o = FALSE,  
  quiet = FALSE  
)
```

Arguments

text	Character vector. Sentences or texts you wish to tokenize.
exclude	Character vector. Which words do you wish to exclude?
lang	Character. Language in text (used for stop words). Example: "spanish" or "english". Set to NA to ignore.
min_word_freq	Integer. This will discard words that appear less than <int> times. Defaults to 2. Set to NA to ignore.
min_word_len	Integer. This will discard words that have less than <int> characters. Defaults to 5. Set to NA to ignore.
keep_spaces	Boolean. If you wish to keep spaces in each line to keep unique compound words, separated with spaces, set to TRUE. For example, 'one two' will be set as 'one_two' and treated as a single word.
lowercase, remove_numbers, remove_punct	Boolean.
remove_leett	Boolean. Repeated letters (more than 3 consecutive).
laughs	Boolean. Try to unify all laughs texts.

<code>utf</code>	Boolean. Transform all characters to UTF (no accents and crazy symbols)
<code>df</code>	Boolean. Return a dataframe with a one-hot-encoding kind of results? Each word is a column and returns if word is contained.
<code>h2o</code>	Boolean. Return H2OFrame?
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

Value

`data.frame`. Tokenized words with counters.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [vector2text\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [remove_stopwords\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textFeats\(\)](#), [topics_rake\(\)](#)

`theme_lares`

Theme for ggplot2 (lares)

Description

Based on hrbrthemes' `theme_ipsum` and customized for `lares` usage. With this team you can custom the colour and fill palettes, global colour parameters, major and minor grids, legend, font and font size.

Usage

```
theme_lares(
  font = Sys.getenv("LARES_FONT"),
  size = 12,
  main.colour = "darkorange3",
  hard.colour = "black",
  soft.colour = "grey30",
  plot.colour = "transparent",
  panel.colour = "transparent",
  background = "transparent",
  no.facets = FALSE,
  legend = NULL,
  grid = TRUE,
  axis = TRUE,
  clean = FALSE,
  mg = 9,
  pal = 0,
```

```

palette = NULL,
which = "fc",
quiet = TRUE,
...
)

```

Arguments

font, size	Character and numeric. Base font family and base size for texts. Arial Narrow is set by default when the library is loaded; you may change it with Sys.setenv("LARES_FONT" = "X") or by using this parameter manually. If font is "ignore" or NA, it'll be ignored.
main_colour, hard_colour, soft_colour, plot_colour, panel_colour	Character. Main colours for your theme.
background	Character. Main colour for your background. Overwrites plot_colour and panel_colour.
no_facets	Boolean. Suppress facet labels?
legend	Character. Legend position: "top", "right", "bottom", or "left" You can also set to FALSE or "none" to suppress legend.
grid	Character or Boolean. Use TRUE/FALSE or a combination of X, x, Y, y to enable/disable minor and major grids.
axis	Character or Boolean. Use TRUE/FALSE, x or Y to enable X and/or Y axis lines.
clean	Boolean. Suppress grids and axis? Overwrites both parameters.
mg	Numeric. External margins reference.
pal	Integer. 1 for fill and colour palette, 2 for only colour palette, 3 for only fill palette, 4 for personal labels-colour palette. 0 for nothing.
palette	Character vector. Pass a vector with HEX colour codes to use a custom palette. If you pass a named vector, the name values will be used as fill and the values will be used as colour.
which	Character. When pal = 3, select which colours should be added with the custom colours palette: fill, colour, text (fct) - first letters.
quiet	Boolean. Keep quiet? If not, informative messages will be shown.
...	Additional parameters.

Value

Themed ggplot2 object

Why Arial Narrow?

First and foremost, Arial Narrow is generally installed by default or readily available on any modern system, so it's "free"-ish; plus, it is a condensed font with solid default kerning pairs and geometric numbers.

See Also

Other Themes: [gg_fill_customs\(\)](#), [lares_pal\(\)](#), [plot_palette\(\)](#)

Examples

```
data(dft)
library(ggplot2)
p <- ggplot(dft, aes(x = Pclass, y = sum(Fare), fill = Pclass)) +
  geom_col()
p + theme_lares()
p + theme_lares(pal = 1)
p + theme_lares(background = "#999999", mg = 25)
p + theme_lares(legend = "top", grid = "Yy")
p + theme_lares(clean = TRUE)
```

tic*Stopwatch to measure timings in R***Description**

Start a stopwatch.

Stop a stopwatch.

Usage

```
tic(id = 1, start = proc.time()["elapsed"], quiet = TRUE)

toc(
  id = 1,
  msg = "Elapsed time:",
  type = "units",
  signif = 3,
  quiet = FALSE,
  ...
)
```

Arguments

<code>id</code>	Define ID if multiple <code>tic</code> & <code>toc</code> are being used.
<code>start</code>	Start time. Now is default.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.
<code>msg</code>	Character. Custom message shown
<code>type</code>	Character. Output format for time list element. Choose any of: <code>units</code> , <code>clock</code> , <code>seconds</code> .
<code>signif</code>	Integer. Significant digits.
<code>...</code>	Additional parameters.

Value

Invisible list. Contains tic (start time), toc (stop time), elapsed time and message printed.

toc returns an (invisible) list containing the time-stamps tic and toc, time in seconds and the message msg.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
# Basic use (global stopwatch)
tic()
Sys.sleep(0.1)
toc()

# Multiple tic tocs
tic(id = "two", quiet = FALSE)
Sys.sleep(0.2)
toc(id = "two")

# Global is still working (id = 1)
toc(msg = "The function finished its work in")
```

topics_rake*Keyword/Topic identification using RAKE*

Description

RAKE is a basic algorithm which tries to identify keywords in text. Based on udpipe library, model models, and keywords_rake function.

Usage

```
topics_rake(text, file = "english-ewt-ud-2.4-190531.udpipe", lang = "english")
```

Arguments

text	Character vector
file	Character. Name of udpipe model previously downloaded for a specific language
lang	Character. If file does not exist, this language will be downloaded from udpipe's models.

Value

data.frame with topics for each text input.

See Also

Other Text Mining: [cleanText\(\)](#), [ngrams\(\)](#), [remove_stopwords\(\)](#), [replaceall\(\)](#), [sentimentBreakdown\(\)](#), [textCloud\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#)

tree_var

*Recursive Partitioning and Regression Trees***Description**

Fit and plot a rpart model for exploratory purposes using rpart and rpart.plot libraries.

Usage

```
tree_var(
  df,
  y,
  type = 2,
  max = 3,
  min = 20,
  cp = 0,
  ohse = TRUE,
  plot = TRUE,
  explain = TRUE,
  title = NA,
  subtitle = NULL,
  ...
)
```

Arguments

df	Data frame
y	Variable or Character. Name of the dependent variable or response.
type	Type of plot. Possible values: 0 Draw a split label at each split and a node label at each leaf. 1 Label all nodes, not just leaves. Similar to text.rpart's all=TRUE. 2 Default. Like 1 but draw the split labels below the node labels. Similar to the plots in the CART book. 3 Draw separate split labels for the left and right directions. 4 Like 3 but label all nodes, not just leaves. Similar to text.rpart's fancy=TRUE. See also clip.right.labs .
	5 Show the split variable name in the interior nodes.

max	Integer. Maximal depth of the tree.
min	Integer. The minimum number of observations that must exist in a node in order for a split to be attempted.
cp	complexity parameter. Any split that does not decrease the overall lack of fit by a factor of cp is not attempted. For instance, with anova splitting, this means that the overall R-squared must increase by cp 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 cp will likely be pruned off by cross-validation, and that hence the program need not pursue it.
ohse	Boolean. Auto generate One Hot Smart Encoding?
plot	Boolean. Return a plot? If not, rpart object.
explain	Boolean. Include a brief explanation on the bottom part of the plot.
title, subtitle	Character. Title and subtitle to include in plot. Set to NULL to ignore.
...	Additional parameters passed to rpart.plot().

Value

(Invisible) list type 'tree_var' with plot (function), model, predictions, performance metrics, and interpret auxiliary text.

See Also

Other Exploratory: [corr_cross\(\)](#), [corr_var\(\)](#), [crosstab\(\)](#), [df_str\(\)](#), [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [lasso_vars\(\)](#), [missingness\(\)](#), [plot_cats\(\)](#), [plot_df\(\)](#), [plot_nums\(\)](#)

Other Visualization: [distr\(\)](#), [freqs\(\)](#), [freqs_df\(\)](#), [freqs_list\(\)](#), [freqs_plot\(\)](#), [noPlot\(\)](#), [plot_chord\(\)](#), [plot_survey\(\)](#), [plot_timeline\(\)](#)

Examples

```
data(dft)
# Regression Tree
tree <- tree_var(dft, Fare, subtitle = "Titanic dataset")
tree$plot() # tree plot
tree$model # rpart model object
tree$performance # metrics
# Binary Tree
tree_var(dft, Survived_TRUE, explain = FALSE, cex = 0.8)$plot()
# Multiclass tree
tree_var(dft[, c("Pclass", "Fare", "Age")], Pclass, ohse = FALSE)$plot()
```

trim_mp3*Trim MP3 Audio File***Description**

This function trims MP3 files given a start and/or end numeric timestamp. Requires `ffmpeg` installed in your machine.

Usage

```
trim_mp3(
  file,
  start_time = 0,
  end_time = NA,
  overwrite = FALSE,
  ext = "mp3",
  quiet = FALSE
)
```

Arguments

<code>file</code>	Character. File name to trim.
<code>start_time, end_time</code>	Numeric. Start and end time to trim the audio output in seconds.
<code>overwrite</code>	Boolean. Overwrite original file?
<code>ext</code>	Character. File extension/type.
<code>quiet</code>	Boolean. Keep quiet? If not, informative messages will be shown.

See Also

Other Audio: [get_mp3\(\)](#)

try_require*Check if Specific Package is Installed***Description**

This function checks library dependencies

Usage

```
try_require(package, stop = TRUE, load = TRUE, lib.loc = NULL, ...)
```

Arguments

package	Character. Name of the library
stop	Boolean. Stop if not installed. If FALSE and library is not available, warning will be shown.
load	Boolean. Load library?
lib.loc	Character vector. Location of R library trees to search through, or NULL. The default value of NULL corresponds to all libraries currently known to .libPaths(). Non-existent library trees are silently ignored.
...	Additional parameters.

Value

No return value, called for side effects.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [updateLares\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

Examples

```
# Check if library base is installed. If not, stop and show error
try_require("base", stop = TRUE)
# Check if library xxx is installed. If not, show warning
try_require("xxx", stop = FALSE)
```

updateLares

*Update the library (dev or CRAN version)***Description**

This auxiliary function lets the user update lares to latest CRAN or developer version.

Usage

```
updateLares(force = FALSE, dev = TRUE, all = FALSE, local = FALSE, fb = FALSE)
```

Arguments

force	Boolean. Force install.
dev	Boolean. Developer version (Github)? If not, CRAN version.
all	Boolean. Install other recommended libraries? Kinda Docker install!
local	Boolean. Install package with local files? (or Github repo).
fb	Boolean. From FB instance? Personal internal use.

Value

No return value, called for side effects.

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepm\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [warnifnot\(\)](#), [what_size\(\)](#)

vector2text

*Convert a vector into a comma separated text***Description**

Convert a vector into a comma separated text

Usage

```
vector2text(vector, sep = ", ", quotes = TRUE, force_single = FALSE, and = "")  
v2t(vector, sep = ", ", quotes = TRUE, force_single = FALSE, and = "")
```

Arguments

vector	Vector. Vector with more than 1 observation.
sep	Character. String text wished to insert between values.
quotes	Boolean. Bring simple quotes for each observation.
force_single	Boolean. Force single quotes by replacing \".
and	Character. Add 'and' or something before last observation. Not boolean variable so it can be used on other languages. Note that the last comma will be suppressed if Sys.getenv("LARES_NUMFORMAT") is set to 1 and you have less than 3 values.

Value

Vector pasting vector values into a single string

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [year_month\(\)](#), [zerovar\(\)](#)

Examples

```
vector2text(LETTERS[1:5])
vector2text(c(1:5), quotes = FALSE)
vector2text(c(1:5), quotes = FALSE, sep = "-")
vector2text(c(1:5), and = "and also")
vector2text(c("Text", "R's"), force_single = TRUE)
# Shorter function with same purpose
v2t(LETTERS[1:5])
```

warnifnot

Test the Truth of R Expressions and Warn

Description

If the expression in ... is not TRUE, warning is called, producing a warning message indicating the expression which was not true.

Usage

```
warnifnot(...)
```

Arguments

... any R expression, which should evaluate to TRUE

See Also

Other Tools: [autoline\(\)](#), [bind_files\(\)](#), [bring_api\(\)](#), [chr2num\(\)](#), [db_download\(\)](#), [db_upload\(\)](#), [dont_sleep\(\)](#), [export_plot\(\)](#), [export_results\(\)](#), [files_functions\(\)](#), [font_exists\(\)](#), [formatColoured\(\)](#), [formatHTML\(\)](#), [glued\(\)](#), [grepM\(\)](#), [h2o_selectmodel\(\)](#), [haveInternet\(\)](#), [image_metadata\(\)](#), [importxlsx\(\)](#), [ip_data\(\)](#), [json2vector\(\)](#), [list_cats\(\)](#), [listfiles\(\)](#), [mail_send\(\)](#), [markdown2df\(\)](#), [move_files\(\)](#), [msplit\(\)](#), [myip\(\)](#), [quiet\(\)](#), [read.file\(\)](#), [statusbar\(\)](#), [tic\(\)](#), [try_require\(\)](#), [updateLares\(\)](#), [what_size\(\)](#)

Examples

```
warnifnot(TRUE)
warnifnot(FALSE)
warnifnot(1 + 1 == 3)
```

weighted_value*Calculate weighted stock values using FIFO/LIFO***Description**

Calculate weighted stock values using FIFO/LIFO

Usage

```
weighted_value(
  value,
  n = rep(1, length(value)),
  technique = NULL,
  n_stocks = NULL,
  buy_only = TRUE,
  type = 1,
  ...
)
```

Arguments

<code>value</code>	Numeric vector. Representing the values of the stock.
<code>n</code>	Numeric vector. Representing the volume of the operation. Positive for 'Buy' and negative for 'Sale'.
<code>technique</code>	Character. Pick any of FIFO or LIFO, or NULL to skip.
<code>n_stocks</code>	Integer. Specify the number of stocks to consider. By default will sum positive values of <code>n</code> .
<code>buy_only</code>	Boolean. Consider only buy (positive) values?
<code>type</code>	Integer. 1 for returning the value, 2 for returning the data.frame with the details ("df" attribute)
...	Additional parameters.

Value

The calculated weighted mean value.

Examples

```
values <- c(10, 20, 30, 40, 50)
weights <- c(2, 3, -4, 5, 6)
mean(values)
weighted_value(values)
weighted.mean(values, weights)
weighted_value(values, weights, buy_only = FALSE)
# Using FIFO and LIFO
weighted_value(values, weights, "FIFO")
weighted_value(values, weights, "LIFO", n_stocks = 8)
```

what_size*Calculate the size of any R object or directory*

Description

Calculate the size of any R object or directory

Usage

```
what_size(x = NULL, units = "Mb", path = NULL, recursive = TRUE, ...)
```

Arguments

x	Object
units	Character. Specify which unit to use, i.e. "Gb", "Mb", "Kb".
path	a character vector of full path names; the default corresponds to the working directory, <code>getwd()</code> . Tilde expansion (see <code>path.expand</code>) is performed. Missing values will be ignored. Elements with a marked encoding will be converted to the native encoding (and if that fails, considered non-existent).
recursive	logical. Should the listing recurse into directories?
...	further arguments passed to or from other methods.

See Also

Other Tools: `autoline()`, `bind_files()`, `bring_api()`, `chr2num()`, `db_download()`, `db_upload()`, `dont_sleep()`, `export_plot()`, `export_results()`, `files_functions()`, `font_exists()`, `formatColoured()`, `formatHTML()`, `glued()`, `grepM()`, `h2o_selectmodel()`, `haveInternet()`, `image_metadata()`, `importxlsx()`, `ip_data()`, `json2vector()`, `list_cats()`, `listfiles()`, `mail_send()`, `markdown2df()`, `move_files()`, `msplit()`, `myip()`, `quiet()`, `read.file()`, `statusbar()`, `tic()`, `try_require()`, `updateLares()`, `warnifnot()`

Examples

```
what_size(seq(1:1e3), "Kb")
what_size(seq(1:1e6))
what_size(as.character(seq(1:1e6)))
# what_size(path = ".")
```

winsorize*Outliers: Winsorize***Description**

Winsorizing a vector means that a predefined quantum of the smallest and/or the largest values are replaced by less extreme values. Thereby the substitute values are the most extreme retained values.

Usage

```
winsorize(x, thresh = c(0.05, 0.95), na.rm = FALSE)
```

Arguments

- | | |
|--------|--|
| x | Numeric vector. Distribution to be winsorized. |
| thresh | Numeric vector. Lower and upper quantiles thresholds. Set values within [0,1]. |
| na.rm | Boolean. Should NA be omitted to calculate the quantiles? Note that NA in x are preserved and left unchanged anyway. |

Value

Numeric vector transformed.

See Also

Other Outliers: [outlier_tukey\(\)](#), [outlier_zscore\(\)](#), [outlier_zscore_plot\(\)](#)

wordle_check*Wordle Game Validation***Description**

Given and input and a word, validate each letter based on Wordle's rules: correct letter in correct placement (green), correct letter in wrong placement (yellow), letter is not present (red).

Usage

```
wordle_check(
  input,
  word,
  dictionary = NULL,
  lang_dic = "en",
  method = 3,
  print = TRUE
)
```

```
## S3 method for class 'wordle_check'
print(x, print = TRUE, ...)

wordle_dictionary(lang_dic = "en", method = 3, quiet = TRUE)

wordle_simulation(input, word, seed = NULL, quiet = FALSE, ...)

## S3 method for class 'wordle_simulation'
print(x, type = 1, ...)
```

Arguments

input	Character. Word to validate (5-letters)
word	Character. Word actually answer (5-letters).
dictionary	Character vector. List of valid words. If set to NULL then will use modified scrabble_dictionary() to fetch 5 letter words. Use lang_dic param to set language.
lang_dic	Character. Any of: "en", "es". Only used when dictionary parameter is NULL. Requires internet connection the first time. Uses cache.
method	Integer. 1 for scrabble_dictionary(), 3 for scrapping the words taken straight from the game's source code.
print	Boolean. Print validation results?
x	Object to print
...	Additional parameters.
quiet	Boolean. Do not print words as they are being searched.
seed	Numeric. For reproducibility. Accepts more than one: will run as many seeds there are.
type	Integer. 1 for summary and 2 for coloured results.

Value

Invisible vector with results by letter.

Examples

```
word <- "ABBEY"
# Or pick a random one:
# word <- sample(wordle_dictionary("en"), 1)
wordle_check("OPENS", word)
wordle_check("BABES", word)
wordle_check("KEBAB", word, print = FALSE)
wordle_check("ABYSS", word)
wordle_check("ABBEY", word)
# Feel free to use scrabble_words() for hints

x <- wordle_simulation(input = "SAINT", word = "ABBEY", seed = 1:3)
```

```
print(x)
```

x2y

Ranked Predictive Power of Cross-Features (x2y)

Description

The relative reduction in error when we go from a baseline model (average for continuous and most frequent for categorical features) to a predictive model, can measure the strength of the relationship between two features. In other words, x2y measures the ability of x to predict y. We use CART (Classification And Regression Trees) models to be able to 1) compare numerical and non-numerical features, 2) detect non-linear relationships, and 3) because they are easy/quick to train.

Usage

```
x2y(
  df,
  target = NULL,
  symmetric = FALSE,
  target_x = FALSE,
  target_y = FALSE,
  plot = FALSE,
  top = 20,
  quiet = "auto",
  ohse = FALSE,
  corr = FALSE,
  ...
)

x2y_metric(x, y, confidence = FALSE, bootstraps = 20, max_cat = 20)

## S3 method for class 'x2y_preds'
plot(x, corr = FALSE, ...)

## S3 method for class 'x2y'
plot(x, type = 1, ...)

x2y_preds(x, y, max_cat = 10)
```

Arguments

- | | |
|--------|--|
| df | data.frame. Note that variables with no variance will be ignored. |
| target | Character vector. If you are only interested in the x2y values between particular variable(s) in df, set name(s) of the variable(s) you are interested in. Keep NULL to calculate for every variable (column). Check target_x and target_y parameters as well. |

<code>symmetric</code>	Boolean. <code>x2y</code> metric is not symmetric with respect to <code>x</code> and <code>y</code> . The extent to which <code>x</code> can predict <code>y</code> can be different from the extent to which <code>y</code> can predict <code>x</code> . Set <code>symmetric=TRUE</code> if you wish to average both numbers.
<code>target_x, target_y</code>	Boolean. Force target features to be part of <code>x</code> OR <code>y</code> ?
<code>plot</code>	Boolean. Return a plot? If not, only a <code>data.frame</code> with calculated results will be returned.
<code>top</code>	Integer. Show/plot only top N predictive cross-features. Set to <code>NULL</code> to return all.
<code>quiet</code>	Boolean. Keep quiet? If not, show progress bar.
<code>ohse</code>	Boolean. Use <code>lares::ohse()</code> to pre-process the data?
<code>corr</code>	Boolean. Add correlation and pvalue data to compare with? For more custom studies, use <code>lares::corr_cross()</code> directly.
<code>...</code>	Additional parameters passed to <code>x2y_metric()</code>
<code>x, y</code>	Vectors. Categorical or numerical vectors of same length.
<code>confidence</code>	Boolean. Calculate 95% confidence intervals estimated with N bootstraps.
<code>bootstraps</code>	Integer. If <code>confidence=TRUE</code> , how many bootstraps? The more iterations we run the more precise the confidence internal will be.
<code>max_cat</code>	Integer. Maximum number of unique <code>x</code> or <code>y</code> values when categorical. Will select then most frequent values and the rest will be passed as "".
<code>type</code>	Integer. Plot type: 1 for tile plot, 2 for ranked bar plot.

Details

This `x2y` metric is based on Rama Ramakrishnan's [post](#): An Alternative to the Correlation Coefficient That Works For Numeric and Categorical Variables. This analysis complements our `lares::corr_cross()` output.

Value

Depending on `plot` input, a plot or a `data.frame` with `x2y` results.

Examples

```
data(dft) # Titanic dataset
x2y_results <- x2y(dft, quiet = TRUE, max_cat = 10, top = NULL)
head(x2y_results, 10)
plot(x2y_results, type = 2)

# Confidence intervals with 10 bootstrap iterations
x2y(dft,
  target = c("Survived", "Age"),
  confidence = TRUE, bootstraps = 10, top = 8
)

# Compare with mean absolute correlations
x2y(dft, "Fare", corr = TRUE, top = 6, target_x = TRUE)
```

```

# Plot (symmetric) results
symm <- x2y(dft, target = "Survived", symmetric = TRUE)
plot(symm, type = 1)

# Symmetry: x2y vs y2x
on.exit(set.seed(42))
x <- seq(-1, 1, 0.01)
y <- sqrt(1 - x^2) + rnorm(length(x), mean = 0, sd = 0.05)

# Knowing x reduces the uncertainty about the value of y a lot more than
# knowing y reduces the uncertainty about the value of x. Note correlation.
plot(x2y_preds(x, y), corr = TRUE)
plot(x2y_preds(y, x), corr = TRUE)

```

year_month*Convert Date into Year-Month, Year-Quarter or Year-Week Format***Description**

This function lets the user convert a date into YYYY-MM, YYYY-QX, or YYYY-WW format easily.

Usage

```

year_month(date)

year_quarter(date)

year_week(date)

```

Arguments

date Date vector. Date to transform format.

Value

Vector with dates reformatted

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [zerovar\(\)](#)

Examples

```
year_month(Sys.Date())
year_quarter(Sys.Date())
year_week(Sys.Date())
```

zerovar

Zero Variance Columns

Description

This function detects which columns have the same value (whichever) for each column.

Usage

```
zerovar(df)
```

Arguments

df	Dataframe
----	-----------

Value

Character vector with column names on which its values have no variance.

See Also

Other Data Wrangling: [balance_data\(\)](#), [categ_reducer\(\)](#), [cleanText\(\)](#), [date_cuts\(\)](#), [date_feats\(\)](#), [file_name\(\)](#), [formatHTML\(\)](#), [holidays\(\)](#), [impute\(\)](#), [left\(\)](#), [normalize\(\)](#), [num_abbr\(\)](#), [ohe_commas\(\)](#), [ohse\(\)](#), [quants\(\)](#), [removenacols\(\)](#), [replaceall\(\)](#), [replacefactor\(\)](#), [textFeats\(\)](#), [textTokenizer\(\)](#), [vector2text\(\)](#), [year_month\(\)](#)

Examples

```
df <- data.frame(a = c(1, NA, 3), b = rep(NA, 3), c = rep(5, 3))
print(df)
zerovar(df)
```

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