

# Package ‘creditmodel’

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**Title** Toolkit for Credit Modeling, Analysis and Visualization

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**Description** Provides a highly efficient R tool suite for Credit Modeling, Analysis and Visualization. Contains infrastructure functionalities such as data exploration and preparation, missing values treatment, outliers treatment, variable derivation, variable selection, dimensionality reduction, grid search for hyper parameters, data mining and visualization, model evaluation, strategy analysis etc. This package is designed to make the development of binary classification models (machine learning based models as well as credit scorecard) simpler and faster.

**Depends** R(>= 3.2)

**Imports** data.table,dplyr,ggplot2,foreach,doParallel,glmnet,rpart,cli,xgboost

**Suggests** pdp,pmml,XML,knitr,gbm,randomForest,rmarkdown

**VignetteBuilder** knitr

**Encoding** UTF-8

**ByteCompile** yes

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**LazyLoad** yes

**License** AGPL-3

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creditmodel-package     *creditmodel: toolkit for credit modeling and data analysis*

---

## Description

creditmodel provides a highly efficient R tool suite for Credit Modeling, Analysis and Visualization. Contains infrastructure functionalities such as data exploration and preparation, missing values treatment, outliers treatment, variable derivation, variable selection, dimensionality reduction, grid search for hyper parameters, data mining and visualization, model evaluation, strategy analysis etc. This package is designed to make the development of binary classification models (machine learning based models as well as credit scorecard) simpler and faster.

## Details

It has three main goals:

- creditmodel is a free and open source automated modeling R package designed to help model developers improve model development efficiency and enable many people with no background in data science to complete the modeling work in a short time. Let them focus more on the problem itself and allocate more time to decision-making.
- creditmodel covers various tools such as data preprocessing, variable processing/derivation, variable screening/dimensionality reduction, modeling, data analysis, data visualization, model evaluation, strategy analysis, etc. It is a set of customized "core" tool kit for model developers.
- 'creditmodel' is suitable for machine learning automated modeling of classification targets, and is more suitable for the risk and marketing data of financial credit, e-commerce, and insurance with relatively high noise and low information content.

To learn more about creditmodel, start with the WeChat Platform: [hansenmode](#)

## Author(s)

**Maintainer:** Dongping Fan <fdp@pku.edu.cn>

---

address\_varieble     *address\_varieble*

---

## Description

This function is not intended to be used by end user.

## Usage

```
address_varieble(df, address_cols = NULL, address_pattern = NULL,  
parallel = TRUE)
```

**Arguments**

df	A data.frame.
address_cols	Variables of address,
address_pattern	Regular expressions, used to match address variable names.
parallel	Logical, parallel computing. Default is TRUE.

---

add\_variable\_process    *add\_variable\_process*

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
add_variable_process(add)
```

**Arguments**

add	A data.frame contained address variables.
-----	---

---

analysis\_nas            *missing Analysis*

---

**Description**

#' analysis\_nas is for understanding the reason for missing data and understand distribution of missing data so we can categorise it as:

- missing completely at random(MCAR)
- Mmissing at random(MAR), or
- missing not at random, also known as IM.

**Usage**

```
analysis_nas(dat, class_var = FALSE, nas_rate = NULL, na_vars = NULL,
  mat_nas_shadow = NULL, dt_nas_random = NULL, ...)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
class_var	Logical, nas analysis of the nominal variables. Default is TRUE.
nas_rate	A list contains nas rate of each variable.
na_vars	Names of variables which contain nas.
mat_nas_shadow	A shadow matrix of variables which contain nas.
dt_nas_random	A data.frame with random nas imputation.
...	Other parameters.

**Value**

A data.frame with outliers analysis for each variable.

---

analysis\_outliers      *Outliers Analysis*

---

**Description**

#' analysis\_outliers is the function for outliers analysis.

**Usage**

```
analysis_outliers(dat, target, x, lof = NULL)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
target	The name of target variable.
x	The name of variable to process.
lof	Outliers of each variable detected by outliers_detection.

**Value**

A data.frame with outliers analysis for each variable.

---

as_percent	<i>Percent Format</i>
------------	-----------------------

---

**Description**

as\_percent is a small function for making percent format..

**Usage**

```
as_percent(x, digits = 2)
```

**Arguments**

x	A numeric vector or list.
digits	Number of digits.Default: 2.

**Value**

x with percent format.

**Examples**

```
as_percent(0.2363, digits = 2)
as_percent(1)
```

---

auc_value	<i>auc_value auc_value is for get best lambda required in lasso_filter. This function required in lasso_filter</i>
-----------	--

---

**Description**

auc\_value auc\_value is for get best lambda required in lasso\_filter. This function required in lasso\_filter

**Usage**

```
auc_value(target, prob)
```

**Arguments**

target	Vector of target.
prob	A list of predict probability or score.

**Value**

Lambda value



---

char_cor_vars	<i>Cramer's V matrix between categorical variables.</i>
---------------	---

---

## Description

char\_cor\_vars is function for calculating Cramer's V matrix between categorical variables. char\_cor is function for calculating the correlation coefficient between variables by cremers 'V

## Usage

```
char_cor_vars(dat, x)
```

```
char_cor(dat, x_list = NULL, ex_cols = "date$", parallel = FALSE,  
note = FALSE)
```

## Arguments

dat	A data frame.
x	The name of variable to process.
x_list	Names of independent variables.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
parallel	Logical, parallel computing. Default is FALSE.
note	Logical. Outputs info. Default is TRUE.

## Value

A list contains correlation index of x with other variables in dat.

## Examples

```
## Not run:  
char_x_list = get_names(dat = UCICreditCard,  
types = c('factor', 'character'),  
ex_cols = "ID$|date$|default.payment.next.month$", get_ex = FALSE)  
char_cor(dat = UCICreditCard[char_x_list])  
  
## End(Not run)
```

---

char_to_num	<i>character to number</i>
-------------	----------------------------

---

### Description

char\_to\_num is for transferring character variables which are actually numerical numbers containing strings to numeric.

### Usage

```
char_to_num(dat, char_list = NULL, note = TRUE, ex_cols = NULL)
```

### Arguments

dat	A data frame
char_list	The list of characteristic variables that need to merge categories, Default is NULL. In case of NULL, merge categories for all variables of string type.
note	Logical, outputs info. Default is TRUE.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.

### Value

A data.frame

### Examples

```
dat_sub = lendingclub[c('dti_joint', 'emp_length')]
str(dat_sub)
#variables that are converted to numbers containing strings
dat_sub = char_to_num(dat_sub)
str(dat_sub)
```

---

checking_data	<i>Checking Data</i>
---------------	----------------------

---

### Description

checking\_data cheking dat before processing.

### Usage

```
checking_data(dat = NULL, target = NULL, occur_time = NULL,
  note = FALSE, pos_flag = NULL)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
target	The name of target variable. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
note	Logical.Outputs info.Default is TRUE.
pos_flag	The value of positive class of target variable, default: "1".

**Value**

data.frame

**Examples**

```
dat = checking_data(dat = UCICreditCard, target = "default.payment.next.month")
```

---

check\_rules

*check rules*

---

**Description**

get\_ctree\_rules This function is used to check rules.

**Usage**

```
check_rules(rules_list, test_dat, target = NULL, value = NULL)
```

**Arguments**

rules_list	A list of rules.
test_dat	A data.frame of test.
target	The name of target variable, default is NULL.
value	The name of value to gather.

**Value**

A data frame with tree rules and percent under each rule.

**See Also**

[get\\_ctree\\_rules](#), [rules\\_filter](#)

**Examples**

```

train_test = train_test_split(UCICreditCard, split_type = "Random", prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
dat_train$default.payment.next.month = as.numeric(dat_train$default.payment.next.month)
rules_list = get_ctree_rules(tree_fit = NULL, train_dat = dat_train[, 8:26],
                             target = "default.payment.next.month", test_dat = dat_test[10:12,2])
check_rules(rules_list = rules_list, target = "default.payment.next.month",
            test_dat = dat_test, value = NULL)

```

---

city_varieble	<i>city_variable</i>
---------------	----------------------

---

**Description**

This function is used for city variables derivation.

**Usage**

```

city_variable(df = df, city_cols = NULL, city_pattern = NULL,
             city_class = city_class, parallel = TRUE)

```

**Arguments**

df	A data.frame.
city_cols	Variables of city,
city_pattern	Regular expressions, used to match city variable names. Default is "city\$".
city_class	Class or levels of cities.
parallel	Logical, parallel computing. Default is TRUE.

---

city_variable_process	<i>Processing of Address Variables</i>
-----------------------	--

---

**Description**

This function is not intended to be used by end user.

**Usage**

```

city_variable_process(df_city, x, city_class)

```

**Arguments**

df_city	A data.frame.
x	Variables of city,
city_class	Class or levels of cities.

---

cohort_analysis	<i>cohort_analysis cohort_analysis is for cohort(vintage) analysis.</i>
-----------------	---

---

**Description**

cohort\_analysis cohort\_analysis is for cohort(vintage) analysis.

cohort\_table

**Usage**

```
cohort_analysis(dat, obs_id = NULL, occur_time = NULL, MOB = NULL,
  period = "monthly", status = NULL, amount = NULL, by_out = "cnt",
  start_date = NULL, end_date = NULL, dead_status = 30)
```

```
cohort_table(dat, obs_id = NULL, occur_time = NULL, MOB = NULL,
  period = "monthly", status = NULL, amount = NULL, by_out = "cnt",
  start_date = NULL, end_date = NULL, dead_status = 30)
```

**Arguments**

dat	A data.frame contained id, occur_time, mob, status ...
obs_id	The name of ID of observations or key variable of data. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
MOB	Mobility of book
period	Period of event to analysis. Default is "monthly"
status	Status of observations
amount	The name of variable representing amount. Default is NULL.
by_out	Output: amount (amt) or count (cnt)
start_date	The earliest occurrence time of observations.
end_date	The latest occurrence time of observations.
dead_status	Status of dead observations.

---

cohort_table_plot	<i>cohort_table_plot cohort_table_plot is for plotting cohort(vintage) analysis table.</i>
-------------------	--

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
cohort_table_plot(cohort_dat)
```

```
cohort_plot(cohort_dat)
```

**Arguments**

cohort\_dat      A data.frame generated by cohort\_analysis.

---

cor_heat_plot	<i>Correlation Heat Plot</i>
---------------	------------------------------

---

**Description**

cor\_heat\_plot is for plotting correlation matrix

**Usage**

```
cor_heat_plot(cor_mat, low_color = love_color("deep_red"),  
              high_color = love_color("light_cyan"), title = "Correlation Matrix")
```

**Arguments**

cor\_mat          A correlation matrix.

low\_color        color of the lowest correlation between variables.

high\_color       color of the highest correlation between variables.

title            title of plot.

**Examples**

```
train_test <- train_test_split(UCICreditCard,  
                               split_type = "Random", prop = 0.8, save_data = FALSE)  
dat_train = train_test$train  
dat_test = train_test$test  
cor_mat = cor(dat_train[,8:12], use = "complete.obs")  
cor_heat_plot(cor_mat)
```

---

cor_plot	<i>Correlation Plot</i>
----------	-------------------------

---

**Description**

cor\_plot is for plotting correlation matrix

**Usage**

```
cor_plot(dat, dir_path = tempdir(), x_list = NULL, gtitle = NULL,
         save_data = FALSE, plot_show = FALSE)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
dir_path	The path for periodically saved graphic files. Default is <code>"/model/LR"</code>
x_list	Names of independent variables.
gtitle	The title of the graph & The name for periodically saved graphic file. Default is <code>"_correlation_of_variables"</code> .
save_data	Logical, save results in locally specified folder. Default is TRUE
plot_show	Logical, show graph in current graphic device.

**Examples**

```
train_test <- train_test_split(UCICreditCard,
                               split_type = "Random", prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
cor_plot(dat_train[,8:12], plot_show = TRUE)
```

---

cos_sim	<i>cos_sim</i>
---------	----------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
cos_sim(x, y, cos_margin = 1)
```

**Arguments**

x	A list of numbers
y	A list of numbers
cos_margin	Margin of matrix, 1 for rows and 2 for cols, Default is 1.

**Value**

A number of cosin similarity

---

cross_table	<i>cross_table</i>
-------------	--------------------

---

**Description**

cross\_table is for cross table analysis.

**Usage**

```
cross_table(dat, cross_x, cross_y = NULL, target = NULL,
            value = NULL, cross_type = "total_sum")
```

**Arguments**

dat	A data.frame with independent variables.
cross_x	Names of variables to cross.
cross_y	Names of variables to cross.
target	The name of target variable.
value	The name of the variable to sum. When this parameter is NULL, the default statistics is to sum frequency.
cross_type	Output form of the result of crosstable. Provide these four forms: "total_sum", "total_pct", "bad_sum", "bad_pct".

**Value**

A cross table.

**Examples**

```
cross_table(dat = UCICreditCard, cross_x = "SEX", cross_y = "AGE",
            target = "default.payment.next.month", cross_type = "bad_pct", value = "LIMIT_BAL")
cross_table(dat = UCICreditCard, cross_x = c("SEX", "MARRIAGE"), cross_y = "AGE",
            target = "default.payment.next.month", cross_type = "bad_pct", value = "LIMIT_BAL")
```



---

 customer\_segmentation *Customer Segmentation*


---

**Description**

customer\_segmentation is a function for clustering and find the best segment variable.

**Usage**

```
customer_segmentation(dat, x_list = NULL, ex_cols = NULL,
  cluster_control = list(meth = "Kmeans", kc = 2, nstart = 1, epsm =
  0.000001, sf = 2, max_iter = 100), tree_control = list(cv_folds = 5,
  maxdepth = kc + 1, minbucket = nrow(dat)/(kc + 1)), save_data = FALSE,
  file_name = NULL, dir_path = tempdir())
```

**Arguments**

dat	A data.frame contained only predict variables.
x_list	A list of x variables.
ex_cols	A list of excluded variables. Default is NULL.
cluster_control	<p>A list controls cluster. kc is the number of cluster center (default is 2), nstart is the number of random groups (default is 1), max_iter max iteration number(default is 100) .</p> <ul style="list-style-type: none"> <li>• meth Method of clustering. Provides two mehods,"Kmeans" and "FCM(Fuzzy Cluster Means)"(default is "Kmeans").</li> <li>• kc Number of cluster center (default is 2).</li> <li>• nstart Number of random groups (default is 1).</li> <li>• max_iter Max iteration number(default is 100).</li> </ul>
tree_control	<p>A list of controls for desision tree to find the best segment variable.</p> <ul style="list-style-type: none"> <li>• cv_folds Number of cross-validations(default is 5).</li> <li>• maxdepth Maximum depth of a tree(default is kc +1).</li> <li>• minbucket Minimum percent of observations in any terminal &lt;leaf&gt; node (default is nrow(dat) / (kc + 1)).</li> </ul>
save_data	Logical. If TRUE, save outliers analysis file to the specified folder at dir_path
file_name	The name for periodically saved segmentation file. Default is NULL.
dir_path	The path for periodically saved segmentation file.

**Value**

A "data.frame" object contains cluster results.

## References

Bezdek, James C. "FCM: The fuzzy c-means clustering algorithm". Computers & Geosciences (0098-3004), [https://doi.org/10.1016/0098-3004\(84\)90020-7](https://doi.org/10.1016/0098-3004(84)90020-7)

## Examples

```
clust <- customer_segmentation(dat = lendingclub[1:10000,20:30],
                             x_list = NULL, ex_cols = "id$|loan_status",
                             cluster_control = list(meth = "FCM", kc = 2), save_data = FALSE,
                             tree_control = list(minbucket = round(nrow(lendingclub) / 10)),
                             file_name = NULL, dir_path = tempdir())
```

---

cut\_equal

*Generating Initial Equal Size Sample Bins*

---

## Description

cut\_equal is used to generate initial breaks for equal frequency binning.

## Usage

```
cut_equal(dat_x, g = 10, sp_values = NULL, cut_bin = "equal_depth")
```

## Arguments

dat_x	A vector of an variable x.
g	numeric, number of initial bins for equal_bins.
sp_values	a list of special value. Default: list(-1, "missing")
cut_bin	A string, 'equal_depth' or 'equal_width', default is 'equal_depth'.

## See Also

[get\\_breaks](#), [get\\_breaks\\_all](#), [get\\_tree\\_breaks](#)

## Examples

```
#equal sample size breaks
equ_breaks = cut_equal(dat = UCICreditCard[, "PAY_AMT2"], g = 10)
```

---

cv_split	<i>Stratified Folds</i>
----------	-------------------------

---

**Description**

this function creates stratified folds for cross validation.

**Usage**

```
cv_split(dat, k = 5, occur_time = NULL, seed = 46)
```

**Arguments**

dat	A data.frame.
k	k is an integer specifying the number of folds.
occur_time	time variable for creating OOT folds. Default is NULL.
seed	A seed. Default is 46.

**Value**

a list of indices

**Examples**

```
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
```

---

data_cleansing	<i>Data Cleaning</i>
----------------	----------------------

---

**Description**

The data\_cleansing function is a simpler wrapper for data cleaning functions, such as delete variables that values are all NAs; checking dat and target format. delete low variance variables replace null or NULL or blank with NA; encode variables which NAs & miss value rate is more than 95 encode variables which unique value rate is more than 95 merge categories of character variables that is more than 10; transfer time variables to dateformation; remove duplicated observations; process outliers; process NAs.

**Usage**

```
data_cleansing(dat, target = NULL, obs_id = NULL, occur_time = NULL,
  pos_flag = NULL, x_list = NULL, ex_cols = NULL,
  miss_values = NULL, remove_dup = TRUE, outlier_proc = TRUE,
  missing_proc = "median", low_var = 0.999, missing_rate = 0.999,
  merge_cat = FALSE, note = TRUE, parallel = FALSE,
  save_data = FALSE, file_name = NULL, dir_path = tempdir())
```

**Arguments**

dat	A data frame with x and target.
target	The name of target variable.
obs_id	The name of ID of observations.Default is NULL.
occur_time	The name of occur time of observations.Default is NULL.
pos_flag	The value of positive class of target variable, default: "1".
x_list	A list of x variables.
ex_cols	A list of excluded variables. Default is NULL.
miss_values	Other extreme value might be used to represent missing values, e.g: -9999, -9998. These miss_values will be encoded to -1 or "missing".
remove_dup	Logical, if TRUE, remove the duplicated observations.
outlier_proc	Logical, process outliers or not. Default is TRUE.
missing_proc	If logical, process missing values or not. If "median", then Nas imputation with k neighbors median. If "avg_dist", the distance weighted average method is applied to determine the NAs imputation with k neighbors. If "default", assigning the missing values to -1 or "missing", otherwise ,processing the missing values according to the results of missing analysis.
low_var	The maximum percent of unique values (including NAs) for filtering low variance variables.
missing_rate	The maximum percent of missing values for recoding values to missing and non_missing.
merge_cat	The minimum number of categories for merging categories of character variables.
note	Logical. Outputs info. Default is TRUE.
parallel	Logical, parallel computing or not. Default is FALSE.
save_data	Logical, save the result or not. Default is FALSE.
file_name	The name for periodically saved data file. Default is NULL.
dir_path	The path for periodically saved data file. Default is tempdir().

**Value**

A preprocessed data.frame

**See Also**

[remove\\_duplicated](#), [null\\_blank\\_na](#), [entry\\_rate\\_na](#), [low\\_variance\\_filter](#), [process\\_nas](#), [process\\_outliers](#)

**Examples**

```
#data cleaning
dat_cl = data_cleansing(dat = UCICreditCard[1:2000,],
                        target = "default.payment.next.month",
                        x_list = NULL,
                        obs_id = "ID",
                        occur_time = "apply_date",
                        ex_cols = c("PAY_6|BILL_"),
                        outlier_proc = TRUE,
                        missing_proc = TRUE,
                        low_var = TRUE,
                        save_data = FALSE)
```

---

data_exploration	<i>Data Exploration</i>
------------------	-------------------------

---

**Description**

#The data\_exploration includes both univariate and bivariate analysis and ranges from univariate statistics and frequency distributions, to correlations, cross-tabulation and characteristic analysis.

**Usage**

```
data_exploration(dat, save_data = FALSE, file_name = NULL,
                 dir_path = tempdir())
```

**Arguments**

dat	A data.frame with x and target.
save_data	Logical. If TRUE, save files to the specified folder at dir_path
file_name	The file name for periodically saved outliers analysis file. Default is NULL.
dir_path	The path for periodically saved outliers analysis file. Default is tempdir().

**Value**

A list contains both category and numeric variable analysis.

**Examples**

```
data_ex = data_exploration(dat = UCICreditCard[1:1000,])
```

---

date_cut	<i>Date Time Cut Point</i>
----------	----------------------------

---

**Description**

date\_cut is a small function to get date point.

**Usage**

```
date_cut(dat_time, pct = 0.7)
```

**Arguments**

dat_time	time vectors.
pct	the percent of cutting. Default: 0.7.

**Value**

A Date.

**Examples**

```
date_cut(dat_time = lendingclub$issue_d, pct = 0.8)
#"2018-08-01"
```

---

derived_interval	<i>derived_interval</i>
------------------	-------------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
derived_interval(dat_s, interval_type = c("cnt_interval",
    "time_interval"))
```

**Arguments**

dat_s	A data.frame contained only predict variables.
interval_type	Available of c("cnt_interval", "time_interval")

---

derived_partial_acf	<i>derived_partial_acf</i>
---------------------	----------------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
derived_partial_acf(dat_s)
```

**Arguments**

dat_s	A data.frame
-------	--------------

---

derived_pct	<i>derived_pct</i>
-------------	--------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
derived_pct(dat_s, pct_type = "total_pct")
```

**Arguments**

dat_s	A data.frame contained only predict variables.
pct_type	Available of "total_pct"

---

 derived\_ts\_vars      *Derivation of Behavioral Variables*


---

**Description**

This function is used for derivating behavioral variables and is not intended to be used by end user.

**Usage**

```
derived_ts_vars(dat, grx = NULL, td = NULL, ID = NULL,
  ex_cols = NULL, x_list = NULL, der = c("cvs", "sums", "means",
  "maxs", "max_mins", "time_intervals", "cnt_intervals", "total_pcts",
  "cum_pcts", "partial_acfs"), parallel = TRUE, note = TRUE)
```

```
derived_ts(dat, grx_x = NULL, x_list = NULL, td = NULL, ID = NULL,
  ex_cols = NULL, der = c("cvs", "sums", "means", "maxs", "max_mins",
  "time_intervals", "cnt_intervals", "total_pcts", "cum_pcts",
  "partial_acfs"))
```

**Arguments**

dat	A data.frame contained only predict variables.
grx	Regular expressions used to match variable names.
td	Number of variables to derivate.
ID	The name of ID of observations or key variable of data. Default is NULL.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
x_list	Names of independent variables.
der	Variables to derivate
parallel	Logical, parallel computing. Default is FALSE.
note	Logical, outputs info. Default is TRUE.
grx_x	Regular expression used to match a group of variable names.

**Details**

The key to creating a good model is not the power of a specific modelling technique, but the breadth and depth of derived variables that represent a higher level of knowledge about the phenomena under examination.



---

de\_one\_hot\_encoding    *Recovery One-Hot Encoding*

---

### Description

de\_one\_hot\_encoding is for one-hot encoding recovery processing

### Usage

```
de_one_hot_encoding(dat_one_hot, cat_vars = NULL, na_act = TRUE,  
  note = FALSE)
```

### Arguments

dat_one_hot	A dat frame with the one hot encoding variables
cat_vars	variables to be recovery processed, default is null, if null, find these variables through regular expressions .
na_act	Logical,If true, the missing value is assigned as "missing", if FALSE missing value is omitted, the default is TRUE.
note	Logical.Outputs info.Default is TRUE.

### Value

A dat frame with the one hot encoding recovery character variables

### See Also

[one\\_hot\\_encoding](#)

### Examples

```
#one hot encoding  
dat1 = one_hot_encoding(dat = UCICreditCard,  
  cat_vars = c("SEX", "MARRIAGE"),  
  merge_cat = TRUE, na_act = TRUE)  
#de one hot encoding  
dat2 = de_one_hot_encoding(dat_one_hot = dat1,  
  cat_vars = c("SEX", "MARRIAGE"),  
  na_act = FALSE)
```

---

de_percent	<i>Recovery Percent Format</i>
------------	--------------------------------

---

**Description**

de\_percent is a small function for recovering percent format..

**Usage**

```
de_percent(x, digits = 2)
```

**Arguments**

x	Character with percent format.
digits	Number of digits.Default: 2.

**Value**

x without percent format.

**Examples**

```
de_percent("24%")
```

---

digits_num	<i>Number of digits</i>
------------	-------------------------

---

**Description**

digits\_num is for caculating optimal digits number for numeric variables.

**Usage**

```
digits_num(dat_x)
```

**Arguments**

dat_x	A numeric variable.
-------	---------------------

**Value**

A number of digits

**Examples**

```
## Not run:
digits_num(lendingclub[, "dti"])
# 7

## End(Not run)
```

---

entropy_weight	<i>Entropy Weight Method</i>
----------------	------------------------------

---

**Description**

entropy\_weight is for calculating Entropy Weight.

**Usage**

```
entropy_weight(dat, ID = NULL, pos_vars, neg_vars)
```

**Arguments**

dat	A data.frame with independent variables.
ID	The name of ID variable.
pos_vars	Names or index of positive direction variables, the bigger the better.
neg_vars	Names or index of negative direction variables, the smaller the better.

**Details**

Step1 Raw data normalization Step2 Find out the total amount of contributions of all samples to the index  $X_j$  Step3 Each element of the step generated matrix is transformed into the product of each element and the LN (element), and the information entropy is calculated. Step4 Calculate redundancy. Step5 Calculate the weight of each index.

**Value**

A data.frame with weights of each variable.

**Examples**

```
entropy_weight(dat = ewm_data, ID = "ID",
              pos_vars = -c(7,11),
              neg_vars = c(7,11))
```

---

entry_rate_na	<i>Max Percent of missing Value</i>
---------------	-------------------------------------

---

**Description**

entry\_rate\_na is the function to recode variables with missing values up to a certain percentage with missing and non\_missing.

**Usage**

```
entry_rate_na(dat, nr = 0.98, note = FALSE)
```

**Arguments**

dat	A data frame with x and target.
nr	The maximum percent of NAs.
note	Logical. Outputs info. Default is TRUE.

**Value**

A data.frame

**Examples**

```
datss = entry_rate_na(dat = lendingclub[1:1000, ], nr = 0.98)
```

---

euclid_dist	<i>euclid_dist</i>
-------------	--------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
euclid_dist(x, y, cos_margin = 1)
```

**Arguments**

x	A list
y	A list
cos_margin	rows or cols

---

`eval_auc`*Functions of xgboost feval*

---

**Description**

`eval_auc`, `eval_ks`, `eval_lift`, `eval_tnr` is for getting best params of xgboost.

**Usage**

```
eval_auc(preds, dtrain)
```

```
eval_ks(preds, dtrain)
```

```
eval_tnr(preds, dtrain)
```

```
eval_lift(preds, dtrain)
```

**Arguments**

`preds`            A list of predict probability or score.

`dtrain`           Matrix of x predictors.

**Value**

List of best value

---

`ewm_data`*Entropy Weight Method Data*

---

**Description**

This data is for Entropy Weight Method examples.

**Format**

A data frame with 10 rows and 13 variables.

---

fast\_high\_cor\_filter    *high\_cor\_filter*

---

### Description

`fast_high_cor_filter` In a highly correlated variable group, select the variable with the highest IV.  
`high_cor_filter` In a highly correlated variable group, select the variable with the highest IV.

### Usage

```
fast_high_cor_filter(dat, p = 0.95, x_list = NULL, com_list = NULL,
  ex_cols = NULL, save_data = FALSE, cor_class = TRUE,
  vars_name = TRUE, parallel = FALSE, note = FALSE,
  file_name = NULL, dir_path = tempdir(), ...)
```

```
high_cor_filter(dat, com_list = NULL, x_list = NULL, ex_cols = NULL,
  onehot = TRUE, parallel = FALSE, p = 0.7, file_name = NULL,
  dir_path = tempdir(), save_data = FALSE, note = FALSE, ...)
```

### Arguments

<code>dat</code>	A data.frame with independent variables.
<code>p</code>	Threshold of correlation between features. Default is 0.95.
<code>x_list</code>	Names of independent variables.
<code>com_list</code>	A data.frame with important values of each variable. eg : <code>IV_list</code>
<code>ex_cols</code>	A list of excluded variables. Regular expressions can also be used to match variable names. Default is <code>NULL</code> .
<code>save_data</code>	Logical, save results in locally specified folder. Default is <code>FALSE</code> .
<code>cor_class</code>	Calculate category variables's correlation matrix. Default is <code>FALSE</code> .
<code>vars_name</code>	Logical, output a list of filtered variables or table with detailed compared value of each variable. Default is <code>TRUE</code> .
<code>parallel</code>	Logical, parallel computing. Default is <code>FALSE</code> .
<code>note</code>	Logical. Outputs info. Default is <code>TRUE</code> .
<code>file_name</code>	The name for periodically saved results files. Default is "Feature_selected_COR".
<code>dir_path</code>	The path for periodically saved results files. Default is "./variable".
<code>...</code>	Additional parameters.
<code>onehot</code>	one-hot-encoding independent variables.

### Value

A list of selected variables.

**See Also**

[get\\_correlation\\_group](#), [high\\_cor\\_selector](#), [char\\_cor\\_vars](#)

**Examples**

```
# calculate iv for each variable.
iv_list = feature_selector(dat_train = UCICreditCard[1:1000,], dat_test = NULL,
target = "default.payment.next.month",
occur_time = "apply_date",
filter = c("IV"), cv_folds = 1, iv_cp = 0.01,
ex_cols = "ID$|date$|default.payment.next.month$",
save_data = FALSE, vars_name = FALSE)
fast_high_cor_filter(dat = UCICreditCard[1:1000,],
com_list = iv_list, save_data = FALSE,
ex_cols = "ID$|date$|default.payment.next.month$",
p = 0.9, cor_class = FALSE ,var_name = FALSE)
```

---

feature_selector	<i>Feature Selection Wrapper</i>
------------------	----------------------------------

---

**Description**

`feature_selector` This function uses four different methods (IV, PSI, correlation, xgboost) in order to select important features. The correlation algorithm must be used with IV.

**Usage**

```
feature_selector(dat_train, dat_test = NULL, x_list = NULL,
target = NULL, pos_flag = NULL, occur_time = NULL,
ex_cols = NULL, filter = c("IV", "PSI", "XGB", "COR"),
cv_folds = 1, iv_cp = 0.01, psi_cp = 0.1, xgb_cp = 0,
cor_cp = 0.98, breaks_list = NULL, hopper = FALSE,
vars_name = TRUE, parallel = FALSE, note = TRUE, seed = 46,
save_data = FALSE, file_name = NULL, dir_path = tempdir(), ...)
```

**Arguments**

<code>dat_train</code>	A data.frame with independent variables and target variable.
<code>dat_test</code>	A data.frame of test data. Default is NULL.
<code>x_list</code>	Names of independent variables.
<code>target</code>	The name of target variable.
<code>pos_flag</code>	The value of positive class of target variable, default: "1".
<code>occur_time</code>	The name of the variable that represents the time at which each observation takes place.
<code>ex_cols</code>	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.

filter	The methods for selecting important and stable variables.
cv_folds	Number of cross-validations. Default: 5.
iv_cp	The minimum threshold of IV. $0 < iv\_i ; 0.01$ to $0.1$ usually work. Default: 0.02
psi_cp	The maximum threshold of PSI. $0 \leq psi\_i \leq 1$ ; $0.05$ to $0.2$ usually work. Default: 0.1
xgb_cp	Threshold of XGB feature's Gain. $0 \leq xgb\_cp \leq 1$ . Default is $1/\text{number of independent variables}$ .
cor_cp	Threshold of correlation between features. $0 \leq cor\_cp \leq 1$ ; $0.7$ to $0.98$ usually work. Default is $0.98$ .
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
hopper	Logical.Filtering screening. Default is FALSE.
vars_name	Logical, output a list of filtered variables or table with detailed IV and PSI value of each variable. Default is FALSE.
parallel	Logical, parallel computing. Default is FALSE.
note	Logical.Outputs info. Default is TRUE.
seed	Random number seed. Default is 46.
save_data	Logical, save results in locally specified folder. Default is FALSE.
file_name	The name for periodically saved results files. Default is "select_vars".
dir_path	The path for periodically saved results files. Default is "./variable"
...	Other parameters.

**Value**

A list of selected features

**See Also**

[psi\\_iv\\_filter](#), [xgb\\_filter](#), [gbm\\_filter](#)

**Examples**

```
feature_selector(dat_train = UCICreditCard[1:1000,c(2,8:12,26)],
               dat_test = NULL, target = "default.payment.next.month",
               occur_time = "apply_date", filter = c("IV", "PSI"),
               cv_folds = 1, iv_cp = 0.01, psi_cp = 0.1, xgb_cp = 0, cor_cp = 0.98,
               vars_name = FALSE, note = FALSE)
```



---

fuzzy\_cluster\_means     *Fuzzy Cluster means.*

---

### Description

This function is used for Fuzzy Clustering.

### Usage

```
fuzzy_cluster_means(dat, kc = 2, sf = 2, nstart = 1,
  max_iter = 100, epsm = 0.000001)
```

```
fuzzy_cluster(dat, kc = 2, init_centers, sf = 3, max_iter = 100,
  epsm = 0.000001)
```

### Arguments

dat	A data.frame contained only predict variables.
kc	The number of cluster center (default is 2),
sf	Default is 2.
nstart	The number of random groups (default is 1),
max_iter	Max iteration number(default is 100) .
epsm	Default is 1e-06.
init_centers	Initial centers of obs.

### References

Bezdek, James C. "FCM: The fuzzy c-means clustering algorithm". Computers & Geosciences (0098-3004),[https://doi.org/10.1016/0098-3004\(84\)90020-7](https://doi.org/10.1016/0098-3004(84)90020-7)

---

gather\_data     *gather or aggregate data*

---

### Description

This function is used for gathering or aggregating data.

### Usage

```
gather_data(dat, x_list = NULL, ID = NULL, FUN = sum_x)
```

**Arguments**

dat	A data.frame contained only predict variables.
x_list	The names of variables to gather.
ID	The name of ID of observations or key variable of data. Default is NULL.
FUN	The function of gathering method.

**Details**

The key to creating a good model is not the power of a specific modelling technique, but the breadth and depth of derived variables that represent a higher level of knowledge about the phenomena under examination.

**Examples**

```
dat = data.frame(id = c(1,1,1,2,2,3,3,3,4,4,4,4,4,5,5,6,7,7,
                      8,8,8,9,9,9,10,10,11,11,11,11,11,11),
                terms = c('a', 'b', 'c', 'a', 'c', 'd', 'd', 'a',
                          'b', 'c', 'a', 'c', 'd', 'a', 'c',
                          'd', 'a', 'e', 'f', 'b', 'c', 'f', 'b',
                          'c', 'h', 'h', 'i', 'c', 'd', 'g', 'k', 'k'),
                time = c(8,3,1,9,6,1,4,9,1,3,4,8,2,7,1,
                        3,4,1,8,7,2,5,7,8,8,2,1,5,7,2,7,3))

gather_data(dat = dat, x_list = "time", ID = 'id', FUN = sum_x)
```

---

gbm\_filter

*Select Features using GBM*


---

**Description**

gbm\_filter is for selecting important features using GBM.

**Usage**

```
gbm_filter(dat, target = NULL, x_list = NULL, ex_cols = NULL,
           pos_flag = NULL, GBM.params = gbm_params(), cores_num = 2,
           vars_name = TRUE, note = TRUE, save_data = FALSE,
           file_name = NULL, dir_path = tempdir(), seed = 46, ...)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
target	The name of target variable.
x_list	Names of independent variables.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.

pos_flag	The value of positive class of target variable, default: "1".
GBM.params	Parameters of GBM.
cores_num	The number of CPU cores to use.
vars_name	Logical, output a list of filtered variables or table with detailed IV and PSI value of each variable. Default is TRUE.
note	Logical, outputs info. Default is TRUE.
save_data	Logical, save results results in locally specified folder. Default is FALSE.
file_name	The name for periodically saved results files. Default is "Feature_importance_GBDT".
dir_path	The path for periodically saved results files. Default is "./variable".
seed	Random number seed. Default is 46.
...	Other parameters to pass to gbdm_params.

**Value**

Selected variables.

**See Also**

[psi\\_iv\\_filter](#), [xgb\\_filter](#), [feature\\_selector](#)

**Examples**

```
GBM.params = gbm_params(n.trees = 2, interaction.depth = 2, shrinkage = 0.1,
                        bag.fraction = 1, train.fraction = 1,
                        n.minobsinnode = 30,
                        cv.folds = 2)

## Not run:
features <- gbm_filter(dat = UCICreditCard[1:1000, c(8:12, 26)],
                      target = "default.payment.next.month",
                      occur_time = "apply_date",
                      GBM.params = GBM.params
                      , vars_name = FALSE)

## End(Not run)
```

---

gbm\_params

*GBM Parameters*


---

**Description**

gbm\_params is the list of parameters to train a GBM using in [training\\_model](#).

**Usage**

```
gbm_params(n.trees = 1000, interaction.depth = 6, shrinkage = 0.01,
           bag.fraction = 0.5, train.fraction = 0.7, n.minobsinnode = 30,
           cv.folds = 5, ...)
```

**Arguments**

n.trees	Integer specifying the total number of trees to fit. This is equivalent to the number of iterations and the number of basis functions in the additive expansion. Default is 100.
interaction.depth	Integer specifying the maximum depth of each tree(i.e., the highest level of variable interactions allowed) . A value of 1 implies an additive model, a value of 2 implies a model with up to 2 - way interactions, etc. Default is 1.
shrinkage	a shrinkage parameter applied to each tree in the expansion. Also known as the learning rate or step - size reduction; 0.001 to 0.1 usually work, but a smaller learning rate typically requires more trees. Default is 0.1 .
bag.fraction	the fraction of the training set observations randomly selected to propose the next tree in the expansion. This introduces randomness into the model fit. If bag.fraction < 1 then running the same model twice will result in similar but different fits. gbm uses the R random number generator so set.seed can ensure that the model can be reconstructed. Preferably, the user can save the returned gbm.object using save. Default is 0.5 .
train.fraction	The first train.fraction * nrows(data) observations are used to fit the gbm and the remainder are used for computing out-of-sample estimates of the loss function.
n.minobsinnode	Integer specifying the minimum number of observations in the terminal nodes of the trees. Note that this is the actual number of observations, not the total weight.
cv.folds	Number of cross - validation folds to perform. If cv.folds > 1 then gbm, in addition to the usual fit, will perform a cross - validation, calculate an estimate of generalization error returned in cv.error.
...	Other parameters

**Details**

See details at: [gbm](#)

**Value**

A list of parameters.

**See Also**

[training\\_model](#), [lr\\_params](#), [xgb\\_params](#), [rf\\_params](#)

---

get_auc_ks_lambda	<i>get_auc_ks_lambda</i> get_auc_ks_lambda is for get best lambda required in lasso_filter. This function required in lasso_filter
-------------------	--

---

### Description

get\_auc\_ks\_lambda get\_auc\_ks\_lambda is for get best lambda required in lasso\_filter. This function required in lasso\_filter

### Usage

```
get_auc_ks_lambda(lasso_model, x_test, y_test, save_data = FALSE,
  plot_show = TRUE, file_name = NULL, dir_path = tempdir())
```

### Arguments

lasso_model	A lasso model generated by glmnet.
x_test	A matrix of test dataset with x.
y_test	A matrix of y test dataset with y.
save_data	Logical, save results in locally specified folder. Default is FALSE
plot_show	Logical, if TRUE plot the results. Default is TRUE.
file_name	The name for periodically saved results files. Default is NULL.
dir_path	The path for periodically saved results files.

### Value

Lambda values with max K-S and AUC.

### See Also

[lasso\\_filter](#), [get\\_sim\\_sign\\_lambda](#)

---

get_bins_table_all	<i>Table of Binning</i>
--------------------	-------------------------

---

### Description

get\_bins\_table is used to generates summary information of variables. get\_bins\_table\_all can generates bins table for all specified independent variables.

**Usage**

```
get_bins_table_all(dat, x_list = NULL, target = NULL,
  pos_flag = NULL, dat_test = NULL, ex_cols = NULL,
  breaks_list = NULL, parallel = FALSE, note = FALSE,
  bins_total = TRUE, save_data = FALSE, file_name = NULL,
  dir_path = tempdir())
```

```
get_bins_table(dat, x, target = NULL, pos_flag = NULL,
  dat_test = NULL, breaks = NULL, breaks_list = NULL,
  bins_total = TRUE, note = FALSE)
```

**Arguments**

<code>dat</code>	A data.frame with independent variables and target variable.
<code>x_list</code>	Names of independent variables.
<code>target</code>	The name of target variable.
<code>pos_flag</code>	Value of positive class, Default is "1".
<code>dat_test</code>	A data.frame of test data. Default is NULL.
<code>ex_cols</code>	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
<code>breaks_list</code>	A table containing a list of splitting points for each independent variable. Default is NULL.
<code>parallel</code>	Logical, parallel computing. Default is FALSE.
<code>note</code>	Logical, outputs info. Default is TRUE.
<code>bins_total</code>	Logical, total sum for each columns.
<code>save_data</code>	Logical, save results in locally specified folder. Default is FALSE.
<code>file_name</code>	The name for periodically saved bins table file. Default is "bins_table".
<code>dir_path</code>	The path for periodically saved bins table file. Default is "./variable".
<code>x</code>	The name of an independent variable.
<code>breaks</code>	Splitting points for an independent variable. Default is NULL.

**See Also**

[get\\_iv](#), [get\\_iv\\_all](#), [get\\_psi](#), [get\\_psi\\_all](#)

**Examples**

```
breaks_list = get_breaks_all(dat = UCICreditCard, x_list = names(UCICreditCard)[3:4],
  target = "default.payment.next.month", equal_bins = TRUE, best = FALSE, g=5,
  ex_cols = "ID|apply_date", save_data = FALSE)
get_bins_table_all(dat = UCICreditCard, breaks_list = breaks_list,
  target = "default.payment.next.month")
```

---

get_breaks_all	<i>Generates Best Breaks for Binning</i>
----------------	--

---

## Description

get\_breaks is for generating optimal binning for numerical and nominal variables. The get\_breaks\_all is a simpler wrapper for get\_breaks.

## Usage

```
get_breaks_all(dat, target = NULL, x_list = NULL, ex_cols = NULL,
  pos_flag = NULL, occur_time = NULL, oot_pct = 0.7, best = TRUE,
  equal_bins = FALSE, cut_bin = "equal_depth", g = 10,
  sp_values = NULL, tree_control = list(p = 0.05, cp = 0.000001, xval =
  5, maxdepth = 10), bins_control = list(bins_num = 10, bins_pct = 0.05,
  b_chi = 0.05, b_odds = 0.1, b_psi = 0.05, b_or = 0.15, mono = 0.3,
  odds_psi = 0.2, kc = 1), parallel = FALSE, note = FALSE,
  save_data = FALSE, file_name = NULL, dir_path = tempdir(), ...)
```

```
get_breaks(dat, x, target = NULL, pos_flag = NULL, best = TRUE,
  equal_bins = FALSE, cut_bin = "equal_depth", g = 10,
  sp_values = NULL, occur_time = NULL, oot_pct = 0.7,
  tree_control = NULL, bins_control = NULL, note = FALSE, ...)
```

## Arguments

dat	A data frame with x and target.
target	The name of target variable.
x_list	A list of x variables.
ex_cols	A list of excluded variables. Default is NULL.
pos_flag	The value of positive class of target variable, default: "1".
occur_time	The name of the variable that represents the time at which each observation takes place.
oot_pct	Percentage of observations retained for overtime test (especially to calculate PSI). Default is 0.7
best	Logical, if TRUE, merge initial breaks to get optimal breaks for binning.
equal_bins	Logical, if TRUE, equal sample size initial breaks generates.If FALSE , tree breaks generates using desision tree.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
g	Integer, number of initial bins for equal_bins.
sp_values	A list of missing values.
tree_control	the list of tree parameters.

- p the minimum percent of observations in any terminal <leaf> node.  $0 < p < 1$ ; 0.01 to 0.1 usually work.
- cp complexity parameter. the larger, the more conservative the algorithm will be.  $0 < cp < 1$ ; 0.0001 to 0.0000001 usually work.
- xval number of cross-validations. Default: 5
- max\_depth maximum depth of a tree. Default: 10

bins\_control the list of parameters.

- bins\_num The maximum number of bins. 5 to 10 usually work. Default: 10
- bins\_pct The minimum percent of observations in any bins.  $0 < bins\_pct < 1$ , 0.01 to 0.1 usually work. Default: 0.02
- b\_chi The minimum threshold of chi-square merge.  $0 < b\_chi < 1$ ; 0.01 to 0.1 usually work. Default: 0.02
- b\_odds The minimum threshold of odds merge.  $0 < b\_odds < 1$ ; 0.05 to 0.2 usually work. Default: 0.1
- b\_psi The maximum threshold of PSI in any bins.  $0 < b\_psi < 1$ ; 0 to 0.1 usually work. Default: 0.05
- b\_or The maximum threshold of G/B index in any bins.  $0 < b\_or < 1$ ; 0.05 to 0.3 usually work. Default: 0.15
- odds\_psi The maximum threshold of Training and Testing G/B index PSI in any bins.  $0 < odds\_psi < 1$ ; 0.01 to 0.3 usually work. Default: 0.1
- mono Monotonicity of all bins, the larger, the more nonmonotonic the bins will be.  $0 < mono < 0.5$ ; 0.2 to 0.4 usually work. Default: 0.2
- kc number of cross-validations. 1 to 5 usually work. Default: 1

parallel Logical, parallel computing or not. Default is FALSE.

note Logical. Outputs info. Default is TRUE.

save\_data Logical, save results in locally specified folder. Default is TRUE

file\_name File name that save results in locally specified folder. Default is "breaks\_list".

dir\_path Path to save results. Default is "./variable"

... Additional parameters.

x The Name of an independent variable.

**Value**

A table containing a list of splitting points for each independent variable.

**See Also**

[get\\_tree\\_breaks](#), [cut\\_equal](#), [select\\_best\\_class](#), [select\\_best\\_breaks](#)

**Examples**

```
#controls
tree_control = list(p = 0.02, cp = 0.000001, xval = 5, maxdepth = 10)
bins_control = list(bins_num = 10, bins_pct = 0.02, b_chi = 0.02, b_odds = 0.1,
```



```

        b_psi = 0.05, b_or = 15, mono = 0.2, odds_psi = 0.1, kc = 5)
# get category variable breaks
b <- get_breaks(dat = UCICreditCard[1:1000,], x = "MARRIAGE",
               target = "default.payment.next.month",
               occur_time = "apply_date",
               sp_values = list(-1, "missing"),
               tree_control = tree_control, bins_control = bins_control)
# get numeric variable breaks
b2 <- get_breaks(dat = UCICreditCard[1:1000,], x = "PAY_2",
                 target = "default.payment.next.month",
                 occur_time = "apply_date",
                 sp_values = list(-1, "missing"),
                 tree_control = tree_control, bins_control = bins_control)
# get breaks of all predictive variables
b3 <- get_breaks_all(dat = UCICreditCard[1:1000,], target = "default.payment.next.month",
                    x_list = c("MARRIAGE", "PAY_2"),
                    occur_time = "apply_date", ex_cols = "ID",
                    sp_values = list(-1, "missing"),
                    tree_control = tree_control, bins_control = bins_control,
                    save_data = FALSE)

```

---

get\_correlation\_group *get\_correlation\_group*

---

## Description

get\_correlation\_group is function for obtaining highly correlated variable groups. select\_cor\_group is function for selecting highly correlated variable group. select\_cor\_list is function for selecting highly correlated variable list.

## Usage

```
get_correlation_group(cor_mat, p = 0.8)
```

```
select_cor_group(cor_vars)
```

```
select_cor_list(cor_vars_list)
```

## Arguments

cor_mat	A correlation matrix of independent variables.
p	Threshold of correlation between features. Default is 0.7.
cor_vars	Correlated variables.
cor_vars_list	List of correlated variable

## Value

A list of selected variables.

**Examples**

```
## Not run:
cor_mat = cor(UCICreditCard[8:20],
use = "complete.obs", method = "spearman")
get_correlation_group(cor_mat, p = 0.6 )

## End(Not run)
```

---

get_ctree_rules	<i>Parse desision tree rules</i>
-----------------	----------------------------------

---

**Description**

get\_ctree\_rules This function is used to desision tree rules and percentage of 1 under each rule.

**Usage**

```
get_ctree_rules(tree_fit = NULL, train_dat = NULL, target = NULL,
test_dat = NULL, tree_control = list(p = 0.05, cp = 0.0001, xval = 1,
maxdepth = 10), seed = 46)
```

**Arguments**

tree_fit	A tree model object.
train_dat	A data.frame of train.
target	The name of target variable.
test_dat	A data.frame of test.
tree_control	the list of parameters to control cutting initial breaks by decision tree.
seed	Random number seed. Default is 46.

**Value**

A data frame with tree rules and 1 percent under each rule.

**See Also**

[rules\\_filter](#), [check\\_rules](#)

**Examples**

```
train_test <- train_test_split(UCICreditCard, split_type = "Random", prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
dat_train$default.payment.next.month = as.numeric(dat_train$default.payment.next.month)
get_ctree_rules(tree_fit = NULL, train_dat = dat_train[, 8:26],
target = "default.payment.next.month", test_dat = dat_test)
```

---

get_iv_all	<i>Calculate Information Value (IV) get_iv is used to calculate Information Value (IV) of an independent variable. get_iv_all can loop through IV for all specified independent variables.</i>
------------	--

---

### Description

Calculate Information Value (IV) get\_iv is used to calculate Information Value (IV) of an independent variable. get\_iv\_all can loop through IV for all specified independent variables.

### Usage

```
get_iv_all(dat, x_list = NULL, ex_cols = NULL, breaks_list = NULL,
  target = NULL, pos_flag = NULL, best = TRUE, equal_bins = FALSE,
  tree_control = NULL, bins_control = NULL, g = 10,
  parallel = FALSE, note = FALSE)
```

```
get_iv(dat, x, target = NULL, pos_flag = NULL, breaks = NULL,
  breaks_list = NULL, best = TRUE, equal_bins = FALSE,
  tree_control = NULL, bins_control = NULL, g = 10, note = FALSE)
```

### Arguments

dat	A data.frame with independent variables and target variable.
x_list	Names of independent variables.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
target	The name of target variable.
pos_flag	Value of positive class, Default is "1".
best	Logical, merge initial breaks to get optimal breaks for binning.
equal_bins	Logical, generates initial breaks for equal frequency binning.
tree_control	Parameters of using Decision Tree to segment initial breaks. See details: <a href="#">get_tree_breaks</a>
bins_control	Parameters used to control binning. See details: <a href="#">select_best_class</a> , <a href="#">select_best_breaks</a>
g	Number of initial breakpoints for equal frequency binning.
parallel	Logical, parallel computing. Default is FALSE.
note	Logical, outputs info. Default is TRUE.
x	The name of an independent variable.
breaks	Splitting points for an independent variable. Default is NULL.

**Details**

IV Rules of Thumb for evaluating the strength a predictor  
 Less than 0.02:unpredictive 0.02 to 0.1:weak 0.1 to 0.3:medium 0.3 + :strong

**References**

Information Value Statistic:Bruce Lund, Magnify Analytics Solutions, a Division of Marketing Associates, Detroit, MI(Paper AA - 14 - 2013)

**See Also**

[get\\_iv](#),[get\\_iv\\_all](#),[get\\_psi](#),[get\\_psi\\_all](#)

**Examples**

```
get_iv_all(dat = UCICreditCard,
  x_list = names(UCICreditCard)[3:10],
  equal_bins = TRUE, best = FALSE,
  target = "default.payment.next.month",
  ex_cols = "ID|apply_date")
get_iv(UCICreditCard, x = "PAY_3",
  equal_bins = TRUE, best = FALSE,
  target = "default.payment.next.month")
```

---

get\_logistic\_coef      *get logistic coef*

---

**Description**

get\_logistic\_coef is for getting logistic coefficients.

**Usage**

```
get_logistic_coef(lg_model, file_name = NULL, dir_path = tempdir(),
  save_data = FALSE)
```

**Arguments**

lg_model	An object of logistic model.
file_name	The name for periodically saved coefficient file. Default is "LR_coef".
dir_path	The Path for periodically saved coefficient file. Default is "./model".
save_data	Logical, save the result or not. Default is FALSE.

**Value**

A data.frame with logistic coefficients.

**Examples**

```

# dataset splitting
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
#rename the target variable
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
  occur_time = "apply_date", miss_values = list("", -1))
#train_ test pliting
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
  x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
  save_data = FALSE, note = FALSE)
#woe transforming
train_woe = woe_trans_all(dat = dat_train,
  target = "target",
  breaks_list = breaks_list,
  woe_name = FALSE)
test_woe = woe_trans_all(dat = dat_test,
  target = "target",
  breaks_list = breaks_list,
  note = FALSE)
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = train_woe[, c("target", x_list)], family = binomial(logit))
#get LR coefficient
dt_imp_LR = get_logistic_coef(lg_model = lr_model, save_data = FALSE)
bins_table = get_bins_table_all(dat = dat_train, target = "target",
  x_list = x_list, dat_test = dat_test,
  breaks_list = breaks_list, note = FALSE)

#score card
LR_score_card <- get_score_card(lg_model = lr_model, bins_table, target = "target")
#scoring
train_pred = dat_train[, c("ID", "apply_date", "target")]
test_pred = dat_test[, c("ID", "apply_date", "target")]
train_pred$pred_LR = score_transfer(model = lr_model,
  tbl_woe = train_woe,
  save_data = TRUE)[, "score"]

test_pred$pred_LR = score_transfer(model = lr_model,
  tbl_woe = test_woe, save_data = FALSE)[, "score"]

```

**Description**

This function is not intended to be used by end user.

**Usage**

```
get_median(x, weight_avg = NULL)
```

**Arguments**

x	A vector or list.
weight_avg	avg weight to calculate means.

---

get\_names

*Get Variable Names*

---

**Description**

get\_names is for getting names of particular classes of variables

**Usage**

```
get_names(dat, types = c("logical", "factor", "character", "numeric",
  "integer64", "integer", "double", "Date", "POSIXlt", "POSIXct",
  "POSIXt"), ex_cols = NULL, get_ex = FALSE)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
types	The class or types of variables which names to get. Default: c('numeric', 'integer', 'double')
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
get_ex	Logical ,if TRUE, return a list contains names of excluded variables.

**Value**

A list contains names of variables

**See Also**

[get\\_x\\_list](#)

**Examples**

```
x_list = get_names(dat = UCICreditCard, types = c('factor', 'character'),
  ex_cols = c("default.payment.next.month", "ID$_date$"), get_ex = FALSE)
x_list = get_names(dat = UCICreditCard, types = c('numeric', 'character', 'integer'),
  ex_cols = c("default.payment.next.month", "ID$SEX "), get_ex = FALSE)
```

---

get_nas_random	<i>get_nas_random</i>
----------------	-----------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
get_nas_random(dat)
```

**Arguments**

dat	A data.frame contained only predict variables.
-----	--

---

get_plots	<i>Plot Independent Variables Distribution</i>
-----------	--

---

**Description**

You can use the plot\_vars to produce plots that characterize the frequency or the distribution of your data. get\_plots can loop through plots for all specified independent variables.

**Usage**

```
get_plots(dat_train, dat_test = NULL, x_list = NULL, target = NULL,
  ex_cols = NULL, breaks_list = NULL, pos_flag = NULL,
  equal_bins = FALSE, cut_bin = "equal_depth", best = TRUE, g = 20,
  tree_control = NULL, bins_control = NULL, plot_show = TRUE,
  save_data = FALSE, file_name = NULL, parallel = FALSE,
  g_width = 8, dir_path = tempdir())
```

```
plot_vars(dat_train, x, target, dat_test = NULL, g_width = 8,
  breaks_list = NULL, breaks = NULL, pos_flag = list("1", 1, "bad",
  "positive"), equal_bins = TRUE, cut_bin = "equal_depth",
  best = FALSE, g = 10, tree_control = NULL, bins_control = NULL,
  plot_show = TRUE, save_data = FALSE, dir_path = tempdir())
```

**Arguments**

dat_train	A data.frame with independent variables and target variable.
dat_test	A data.frame of test data. Default is NULL.
x_list	Names of independent variables.
target	The name of target variable.

ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
pos_flag	Value of positive class, Default is "1".
equal_bins	Logical, generates initial breaks for equal frequency or width binning.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
best	Logical, merge initial breaks to get optimal breaks for binning.
g	Number of initial breakpoints for equal frequency binning.
tree_control	Parameters of using Decision Tree to segment initial breaks. See details: <a href="#">get_tree_breaks</a>
bins_control	Parameters used to control binning. See details: <a href="#">select_best_class</a> , <a href="#">select_best_breaks</a>
plot_show	Logical, show model performance in current graphic device. Default is FALSE.
save_data	Logical, save results in locally specified folder. Default is FALSE.
file_name	The name for periodically saved data file. Default is NULL.
parallel	Logical, parallel computing. Default is FALSE.
g_width	The width of graphs.
dir_path	The path for periodically saved graphic files.
x	The name of an independent variable.
breaks	Splitting points for an independent variable. Default is NULL.

### Examples

```
train_test <- train_test_split(UCICreditCard[1:1000,], split_type = "Random",
  prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
get_plots(dat_train[, c(8, 26)], dat_test = dat_test[, c(8, 26)],
  target = "default.payment.next.month")
```

---

get_psi_all	<i>Calculate Population Stability Index (PSI) get_psi is used to calculate Population Stability Index (PSI) of an independent variable. get_psi_all can loop through PSI for all specified independent variables.</i>
-------------	---

---

### Description

Calculate Population Stability Index (PSI) get\_psi is used to calculate Population Stability Index (PSI) of an independent variable. get\_psi\_all can loop through PSI for all specified independent variables.



**Usage**

```
get_psi_all(dat, x_list = NULL, target = NULL, dat_test = NULL,
  breaks_list = NULL, occur_time = NULL, start_date = NULL,
  cut_date = NULL, oot_pct = 0.7, pos_flag = NULL,
  parallel = FALSE, ex_cols = NULL, as_table = FALSE, g = 10,
  bins_no = TRUE, note = FALSE)
```

```
get_psi(dat, x, target = NULL, dat_test = NULL, occur_time = NULL,
  start_date = NULL, cut_date = NULL, pos_flag = NULL,
  breaks = NULL, breaks_list = NULL, oot_pct = 0.7, g = 10,
  as_table = TRUE, note = FALSE, bins_no = TRUE)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
x_list	Names of independent variables.
target	The name of target variable.
dat_test	A data.frame of test data. Default is NULL.
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
start_date	The earliest occurrence time of observations.
cut_date	Time points for splitting data sets, e.g. : splitting Actual and Expected data sets.
oot_pct	Percentage of observations retained for overtime test (especially to calculate PSI). Default is 0.7
pos_flag	Value of positive class, Default is "1".
parallel	Logical, parallel computing. Default is FALSE.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
as_table	Logical, output results in a table. Default is TRUE.
g	Number of initial breakpoints for equal frequency binning.
bins_no	Logical, add serial numbers to bins. Default is TRUE.
note	Logical, outputs info. Default is TRUE.
x	The name of an independent variable.
breaks	Splitting points for an independent variable. Default is NULL.

**Details**

PSI Rules for evaluating the stability of a predictor Less than 0.02: Very stable 0.02 to 0.1: Stable 0.1 to 0.2: Unstable 0.2 to 0.5] : Change more than 0.5: Great change

**See Also**

[get\\_iv](#), [get\\_iv\\_all](#), [get\\_psi](#), [get\\_psi\\_all](#)

**Examples**

```
# dat_test is null
get_psi(dat = UCICreditCard, x = "PAY_3", occur_time = "apply_date")
# dat_test is not all
# train_test split
train_test = train_test_split(dat = UCICreditCard, prop = 0.7, split_type = "OOT",
                              occur_time = "apply_date", start_date = NULL, cut_date = NULL,
                              save_data = FALSE, note = FALSE)

dat_ex = train_test$train
dat_ac = train_test$test
# generate psi table
get_psi(dat = dat_ex, dat_test = dat_ac, x = "PAY_3",
        occur_time = "apply_date", bins_no = TRUE)
```

---

get\_psi\_iv\_all

*Calculate IV & PSI*

---

**Description**

get\_iv\_psi is used to calculate Information Value (IV) and Population Stability Index (PSI) of an independent variable. get\_iv\_psi\_all can loop through IV & PSI for all specified independent variables.

**Usage**

```
get_psi_iv_all(dat, dat_test = NULL, x_list = NULL, target,
              ex_cols = NULL, pos_flag = NULL, breaks_list = NULL,
              occur_time = NULL, oot_pct = 0.7, equal_bins = FALSE,
              cut_bin = "equal_depth", tree_control = NULL, bins_control = NULL,
              bins_total = FALSE, best = TRUE, g = 10, as_table = TRUE,
              note = FALSE, parallel = FALSE, bins_no = TRUE)
```

```
get_psi_iv(dat, dat_test = NULL, x, target, pos_flag = NULL,
           breaks = NULL, breaks_list = NULL, occur_time = NULL,
           oot_pct = 0.7, equal_bins = FALSE, cut_bin = "equal_depth",
           tree_control = NULL, bins_control = NULL, bins_total = FALSE,
           best = TRUE, g = 10, as_table = TRUE, note = FALSE,
           bins_no = TRUE)
```

**Arguments**

dat                    A data.frame with independent variables and target variable.  
dat\_test                A data.frame of test data. Default is NULL.

x_list	Names of independent variables.
target	The name of target variable.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
pos_flag	The value of positive class of target variable, default: "1".
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
oot_pct	Percentage of observations retained for overtime test (especially to calculate PSI). Default is 0.7
equal_bins	Logical, generates initial breaks for equal frequency or width binning.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
tree_control	Parameters of using Decision Tree to segment initial breaks. See details: <a href="#">get_tree_breaks</a>
bins_control	Parameters used to control binning. See details: <a href="#">select_best_class</a> , <a href="#">select_best_breaks</a>
bins_total	Logical, total sum for each variable.
best	Logical, merge initial breaks to get optimal breaks for binning.
g	Number of initial breakpoints for equal frequency binning.
as_table	Logical, output results in a table. Default is TRUE.
note	Logical, outputs info. Default is TRUE.
parallel	Logical, parallel computing. Default is FALSE.
bins_no	Logical, add serial numbers to bins. Default is FALSE.
x	The name of an independent variable.
breaks	Splitting points for an independent variable. Default is NULL.

### See Also

[get\\_iv](#), [get\\_iv\\_all](#), [get\\_psi](#), [get\\_psi\\_all](#)

### Examples

```
iv_list = get_psi_iv_all(dat = UCICreditCard[1:1000, ],
x_list = names(UCICreditCard)[3:5], equal_bins = TRUE,
target = "default.payment.next.month", ex_cols = "ID|apply_date")
get_psi_iv(UCICreditCard, x = "PAY_3",
target = "default.payment.next.month", bins_total = TRUE)
```

---

<code>get_psi_plots</code>	<i>Plot PSI(Population Stability Index)</i>
----------------------------	---

---

### Description

You can use the `psi_plot` to plot PSI of your data. `get_psi_plots` can loop through plots for all specified independent variables.

### Usage

```
get_psi_plots(dat_train, dat_test = NULL, x_list = NULL,
             ex_cols = NULL, breaks_list = NULL, occur_time = NULL, g = 10,
             plot_show = TRUE, save_data = FALSE, file_name = NULL,
             parallel = FALSE, g_width = 8, dir_path = tempdir())
```

```
psi_plot(dat_train, x, dat_test = NULL, occur_time = NULL,
         g_width = 8, breaks_list = NULL, breaks = NULL, g = 10,
         plot_show = TRUE, save_data = FALSE, dir_path = tempdir())
```

### Arguments

<code>dat_train</code>	A data.frame with independent variables.
<code>dat_test</code>	A data.frame of test data. Default is NULL.
<code>x_list</code>	Names of independent variables.
<code>ex_cols</code>	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
<code>breaks_list</code>	A table containing a list of splitting points for each independent variable. Default is NULL.
<code>occur_time</code>	The name of occur time.
<code>g</code>	Number of initial breakpoints for equal frequency binning.
<code>plot_show</code>	Logical, show model performance in current graphic device. Default is FALSE.
<code>save_data</code>	Logical, save results in locally specified folder. Default is FALSE.
<code>file_name</code>	The name for periodically saved data file. Default is NULL.
<code>parallel</code>	Logical, parallel computing. Default is FALSE.
<code>g_width</code>	The width of graphs.
<code>dir_path</code>	The path for periodically saved graphic files.
<code>x</code>	The name of an independent variable.
<code>breaks</code>	Splitting points for a continues variable.

**Examples**

```

train_test <- train_test_split(UCICreditCard[1:1000,], split_type = "Random",
  prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
get_psi_plots(dat_train[, c(8, 9)], dat_test = dat_test[, c(8, 9)])

```

---

get_score_card	<i>Score Card</i>
----------------	-------------------

---

**Description**

get\_score\_card is for generating a standard scorecard

**Usage**

```

get_score_card(lg_model, target, bins_table, a = 600, b = 50,
  file_name = NULL, dir_path = tempdir(), save_data = FALSE)

```

**Arguments**

lg_model	An object of glm model.
target	The name of target variable.
bins_table	a data.frame generated by <a href="#">get_bins_table</a>
a	Base line of score.
b	Numeric.Increased scores from doubling Odds.
file_name	The name for periodically saved scorecard file. Default is "LR_Score_Card".
dir_path	The path for periodically saved scorecard file. Default is "./model"
save_data	Logical, save results in locally specified folder. Default is FALSE.

**Value**

scorecard

**Examples**

```

# dataset splitting
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
#rename the target variable
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
  occur_time = "apply_date", miss_values = list("", -1))
#train_ test pliting
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

```

```

dat_train = train_test$train
dat_test = train_test$test
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
                             x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
save_data = FALSE, note = FALSE)
#woe transforming
train_woe = woe_trans_all(dat = dat_train,
                          target = "target",
                          breaks_list = breaks_list,
                          woe_name = FALSE)
test_woe = woe_trans_all(dat = dat_test,
                          target = "target",
                          breaks_list = breaks_list,
                          note = FALSE)
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = train_woe[, c("target", x_list)], family = binomial(logit))
#get LR coefficient
dt_imp_LR = get_logistic_coef(lg_model = lr_model, save_data = FALSE)
bins_table = get_bins_table_all(dat = dat_train, target = "target",
                                dat_test = dat_test,
                                x_list = x_list,
                                breaks_list = breaks_list, note = FALSE)

#score card
LR_score_card <- get_score_card(lg_model = lr_model, bins_table, target = "target")
#scoring
train_pred = dat_train[, c("ID", "apply_date", "target")]
test_pred = dat_test[, c("ID", "apply_date", "target")]
train_pred$pred_LR = score_transfer(model = lr_model,
                                   tbl_woe = train_woe,
                                   save_data = FALSE)[, "score"]

test_pred$pred_LR = score_transfer(model = lr_model,
                                   tbl_woe = test_woe, save_data = FALSE)[, "score"]

```

---

get\_shadow\_nas

get\_shadow\_nas

---

## Description

This function is not intended to be used by end user.

## Usage

```
get_shadow_nas(dat)
```

## Arguments

**dat**                    A data.frame contained only predict variables.

---

get\_sim\_sign\_lambda     *get\_sim\_sign\_lambda* get\_sim\_sign\_lambda is for get Best lambda required in lasso\_filter. This function required in lasso\_filter

---

### Description

get\_sim\_sign\_lambda get\_sim\_sign\_lambda is for get Best lambda required in lasso\_filter. This function required in lasso\_filter

### Usage

```
get_sim_sign_lambda(lasso_model, sim_sign = "negative")
```

### Arguments

lasso\_model     A lasso model generated by glmnet.  
sim\_sign        Default is "negative". This is related to pos\_flag. If pos\_flag equals 1 or 1, the value must be set to negative. If pos\_flag equals 0 or 0, the value must be set to positive.

### Details

lambda.sim\_sign give the model with the same positive or negative coefficients of all variables.

### Value

Lambda value

---

get\_tree\_breaks     *Getting the breaks for terminal nodes from decision tree*

---

### Description

get\_tree\_breaks is for generating initial breaks by decision tree for a numerical or nominal variable. The get\_breaks function is a simpler wrapper for get\_tree\_breaks.

### Usage

```
get_tree_breaks(dat, x, target, pos_flag = NULL, tree_control = list(p
= 0.02, cp = 0.000001, xval = 5, maxdepth = 10), sp_values = NULL)
```

**Arguments**

dat	A data frame with x and target.
x	name of variable to cut breaks by tree.
target	The name of target variable.
pos_flag	The value of positive class of target variable, default: "1".
tree_control	the list of parameters to control cutting initial breaks by decision tree. <ul style="list-style-type: none"> <li>• p the minimum percent of observations in any terminal &lt;leaf&gt; node. <math>0 &lt; p &lt; 1</math>; 0.01 to 0.1 usually work.</li> <li>• cp complexity parameter. the larger, the more conservative the algorithm will be. <math>0 &lt; cp &lt; 1</math> ; 0.0001 to 0.0000001 usually work.</li> <li>• xval number of cross-validations. Default: 5</li> <li>• max_depth maximum depth of a tree. Default: 10</li> </ul>
sp_values	A list of special value. Default: NULL.

**See Also**

[get\\_breaks](#), [get\\_breaks\\_all](#)

**Examples**

```
#tree breaks
tree_control = list(p = 0.02, cp = 0.000001, xval = 5, maxdepth = 10)
tree_breaks = get_tree_breaks(dat = UCICreditCard, x = "MARRIAGE",
target = "default.payment.next.month", tree_control = tree_control)
```

---

get\_x\_list

*Get X List.*

---

**Description**

get\_x\_list is for getting intersect names of x\_list, train and test.

**Usage**

```
get_x_list(dat_train = NULL, dat_test = NULL, x_list = NULL,
ex_cols = NULL, note = FALSE)
```

**Arguments**

dat_train	A data.frame with independent variables.
dat_test	Another data.frame.
x_list	Names of independent variables.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
note	Logical. Outputs info. Default is TRUE.



**Value**

A list contains names of variables

**See Also**

[get\\_names](#)

**Examples**

```
x_list = get_x_list(x_list = NULL, dat_train = UCICreditCard,  
ex_cols = c("default.payment.next.month", "ID$_date$"))
```

---

high_cor_selector	<i>Compare the two highly correlated variables</i>
-------------------	--

---

**Description**

high\_cor\_selector is function for comparing the two highly correlated variables, select a variable with the largest IV value.

**Usage**

```
high_cor_selector(cor_mat, p = 0.95, x_list = NULL, com_list = NULL,  
retain = TRUE)
```

**Arguments**

cor_mat	A correlation matrix.
p	The threshold of high correlation.
x_list	Names of independent variables.
com_list	A data.frame with important values of each variable. eg : IV_list.
retain	Logical, output selected variables, if FALSE, output filtered variables.

**Value**

A list of selected variables.

---

is_date	<i>is_date</i>
---------	----------------

---

**Description**

is\_date is a small function for distinguishing time formats

**Usage**

```
is_date(x)
```

**Arguments**

x	list or vectors
---	-----------------

**Value**

A Date.

**Examples**

```
is_date(lendingclub$issue_d)
```

---

knn_nas_imp	<i>Impute nas using KNN</i>
-------------	-----------------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
knn_nas_imp(dat, x, nas_rate = NULL, mat_nas_shadow = NULL,
  dt_nas_random = NULL, k = 10, scale = FALSE, method = "median",
  miss_value_num = -1)
```

**Arguments**

dat	A data.frame with independent variables.
x	The name of variable to process.
nas_rate	A list contains nas rate of each variable.
mat_nas_shadow	A shadow matrix of variables which contain nas.
dt_nas_random	A data.frame with random nas imputation.
k	Number of neighbors of each obs which x is missing.

scale	Logical. Standardization of variable.
method	The methods of imputation by knn. "median" is knn imputation with k neighbors median, "avg_dist" is knn imputation with k neighbors of distance weighted mean.
miss_value_num	Default value of missing data imputation for numeric variables, Default is -1.

---

ks_table	<i>ks_table &amp; plot</i>
----------	----------------------------

---

### Description

ks\_table is for generating a model performance table. ks\_table\_plot is for plotting the table generated by ks\_table ks\_psi\_plot is for K-S & PSI distribution plotting.

### Usage

```
ks_table(train_pred, test_pred = NULL, target = NULL, score = NULL,
         g = 10, breaks = NULL, pos_flag = list("1", "1", "Bad", 1))
```

```
ks_table_plot(train_pred, test_pred, target = "target",
              score = "score", g = 10, plot_show = TRUE, g_width = 12,
              file_name = NULL, save_data = FALSE, dir_path = tempdir(),
              gtitle = NULL)
```

```
ks_psi_plot(train_pred, test_pred, target = "target", score = "score",
            gtitle = NULL, plot_show = TRUE, g_width = 12, save_data = FALSE,
            breaks = NULL, g = 10, dir_path = tempdir())
```

```
model_key_index(tb_pred)
```

### Arguments

train_pred	A data frame of training with predicted prob or score.
test_pred	A data frame of validation with predict prob or score.
target	The name of target variable.
score	The name of prob or score variable.
g	Number of breaks for prob or score.
breaks	Splitting points of prob or score.
pos_flag	The value of positive class of target variable, default: "1".
plot_show	Logical, show model performance in current graphic device. Default is FALSE.
g_width	Width of graphs.
file_name	The name for periodically saved data file. Default is NULL.
save_data	Logical, save results in locally specified folder. Default is FALSE.

dir_path	The path for periodically saved graphic files.
gtitle	The title of the graph & The name for periodically saved graphic file. Default is "_ks_psi_table".
tb_pred	A table generated by code <a href="#">ks_table</a>

### Examples

```

sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
  occur_time = "apply_date", miss_values = list("", -1))

train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))

dat_train$pred_LR = round(predict(lr_model, dat_train[, x_list], type = "response"), 5)
dat_test$pred_LR = round(predict(lr_model, dat_test[, x_list], type = "response"), 5)
# model evaluation
ks_psi_plot(train_pred = dat_train, test_pred = dat_test,
  score = "pred_LR", target = "target",
  plot_show = TRUE)
tb_pred <- ks_table_plot(train_pred = dat_train, test_pred = dat_test,
  score = "pred_LR", target = "target",
  g = 10, g_width = 13, plot_show = FALSE)

key_index = model_key_index(tb_pred)

```

---

ks_value	<i>ks_value</i>
----------	-----------------

---

### Description

ks\_value is for get K-S value for a prob or score.

### Usage

```
ks_value(target, prob)
```

### Arguments

target	Vector of target.
prob	A list of predict probability or score.

**Value**

KS value

---

lasso_filter	<i>Variable selection by LASSO</i>
--------------	------------------------------------

---

**Description**

lasso\_filter filter variables by lasso.

**Usage**

```
lasso_filter(dat_train, dat_test = NULL, target = NULL,
            x_list = NULL, pos_flag = NULL, ex_cols = NULL,
            sim_sign = "negative", best_lambda = "lambda.auc",
            save_data = FALSE, plot.it = TRUE, seed = 46, file_name = NULL,
            dir_path = tempdir(), note = FALSE)
```

**Arguments**

dat_train	A data.frame with independent variables and target variable.
dat_test	A data.frame of test data. Default is NULL.
target	The name of target variable.
x_list	Names of independent variables.
pos_flag	The value of positive class of target variable, default: "1".
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
sim_sign	The coefficients of all variables should be all negative or positive, after turning to woe. Default is "negative" for pos_flag is "1".
best_lambda	Methods of best lambda standards using to filter variables by LASSO. There are 3 methods: ("lambda.auc", "lambda.ks", "lambda.sim_sign"). Default is "lambda.auc".
save_data	Logical, save results in locally specified folder. Default is FALSE
plot.it	Logical, shrinkage plot. Default is TRUE.
seed	Random number seed. Default is 46.
file_name	The name for periodically saved results files. Default is "Feature_selected_LASSO".
dir_path	The path for periodically saved results files. Default is "./variable".
note	Logical, outputs info. Default is FALSE.

**Value**

A list of filtered x variables by lasso.

**Examples**

```

sub = cv_split(UCICreditCard, k = 40)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat_train = data_cleansing(dat, target = "target", obs_id = "ID", occur_time = "apply_date",
  miss_values = list("", -1))
dat_train = process_nas(dat_train)
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
  x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
  save_data = FALSE, note = FALSE)
#woe transform
train_woe = woe_trans_all(dat = dat_train,
  target = "target",
  breaks_list = breaks_list,
  woe_name = FALSE)
lasso_filter(dat_train = train_woe,
  target = "target", x_list = x_list,
  save_data = FALSE, plot.it = FALSE)

```

---

lendingclub

*Lending Club data*


---

**Description**

This data contains complete loan data for all loans issued through the time period stated, including the current loan status (Current, Late, Fully Paid, etc.) and latest payment information. The data containing loan data through the "present" contains complete loan data for all loans issued through the previous completed calendar quarter(time period: 2018Q1:2018Q4).

**Format**

A data frame with 63532 rows and 145 variables.

**Details**

- id: A unique LC assigned ID for the loan listing.
- issue\_d: The month which the loan was funded.
- loan\_status: Current status of the loan.
- addr\_state: The state provided by the borrower in the loan application.
- acc\_open\_past\_24mths: Number of trades opened in past 24 months.
- all\_util: Balance to credit limit on all trades.
- annual\_inc: The self:reported annual income provided by the borrower during registration.
- avg\_cur\_bal: Average current balance of all accounts.
- bc\_open\_to\_buy: Total open to buy on revolving bankcards.

- `bc_util`: Ratio of total current balance to high credit/credit limit for all bankcard accounts.
- `dti`: A ratio calculated using the borrower's total monthly debt payments on the total debt obligations, excluding mortgage and the requested LC loan, divided by the borrower's self-reported monthly income.
- `dti_joint`: A ratio calculated using the co:borrowers' total monthly payments on the total debt obligations, excluding mortgages and the requested LC loan, divided by the co:borrowers' combined self-reported monthly income
- `emp_length`: Employment length in years. Possible values are between 0 and 10 where 0 means less than one year and 10 means ten or more years.
- `emp_title`: The job title supplied by the Borrower when applying for the loan.
- `funded_amnt_inv`: The total amount committed by investors for that loan at that point in time.
- `grade`: LC assigned loan grade
- `inq_last_12m`: Number of credit inquiries in past 12 months
- `installment`: The monthly payment owed by the borrower if the loan originates.
- `max_bal_bc`: Maximum current balance owed on all revolving accounts
- `mo_sin_old_il_acct`: Months since oldest bank installment account opened
- `mo_sin_old_rev_tl_op`: Months since oldest revolving account opened
- `mo_sin_rcnt_rev_tl_op`: Months since most recent revolving account opened
- `mo_sin_rcnt_tl`: Months since most recent account opened
- `mort_acc`: Number of mortgage accounts.
- `pct_tl_nvr_dlq`: Percent of trades never delinquent
- `percent_bc_gt_75`: Percentage of all bankcard accounts > 75
- `purpose`: A category provided by the borrower for the loan request.
- `sub_grade`: LC assigned loan subgrade
- `term`: The number of payments on the loan. Values are in months and can be either 36 or 60.
- `tot_cur_bal`: Total current balance of all accounts
- `tot_hi_cred_lim`: Total high credit/credit limit
- `total_acc`: The total number of credit lines currently in the borrower's credit file
- `total_bal_ex_mort`: Total credit balance excluding mortgage
- `total_bc_limit`: Total bankcard high credit/credit limit
- `total_cu_tl`: Number of finance trades
- `total_il_high_credit_limit`: Total installment high credit/credit limit
- `verification_status_joint`: Indicates if the co:borrowers' joint income was verified by LC, not verified, or if the income source was verified
- `zip_code`: The first 3 numbers of the zip code provided by the borrower in the loan application.

**See Also**

[UCICreditCard](#)

---

lift_value	<i>lift_value</i>
------------	-------------------

---

**Description**

lift\_value is for getting max lift value for a prob or score.

**Usage**

```
lift_value(target, prob)
```

**Arguments**

target	Vector of target.
prob	A list of predict probability or score.

**Value**

Max lift value

---

local_outlier_factor	<i>local_outlier_factor</i> local_outlier_factor is function for calculating the lof factor for a data set using knn This function is not intended to be used by end user.
----------------------	--

---

**Description**

local\_outlier\_factor local\_outlier\_factor is function for calculating the lof factor for a data set using knn This function is not intended to be used by end user.

**Usage**

```
local_outlier_factor(dat, k = 10)
```

**Arguments**

dat	A data.frame contained only predict variables.
k	Number of neighbors for LOF.Default is 10.



---

log_trans	<i>Logarithmic transformation</i>
-----------	-----------------------------------

---

**Description**

log\_trans is for logarithmic transformation

**Usage**

```
log_trans(dat, target, x_list = NULL, cor_dif = 0.01, ex_cols = NULL,
          note = TRUE)
```

```
log_vars(dat, x_list = NULL, target = NULL, cor_dif = 0.01,
          ex_cols = NULL)
```

**Arguments**

dat	A data.frame.
target	The name of target variable.
x_list	A list of x variables.
cor_dif	The correlation coefficient difference with the target of logarithm transformed variable and original variable.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
note	Logical, outputs info. Default is TRUE.

**Value**

Log transformed data.frame.

**Examples**

```
dat = log_trans(dat = UCICreditCard, target = "default.payment.next.month",
                x_list = NULL, cor_dif = 0.01, ex_cols = "ID", note = TRUE)
```

---

loop_function	<i>Loop Function. #' loop_function is an iterator to loop through</i>
---------------	---

---

**Description**

Loop Function. #' loop\_function is an iterator to loop through

**Usage**

```
loop_function(func = NULL, args = list(data = NULL), x_list = NULL,
  bind = "rbind", parallel = TRUE, as_list = FALSE)
```

**Arguments**

func	A function.
args	A list of arguments required by function.
x_list	Names of objects to loop through.
bind	Complie results, "rbind" & "cbind" are available.
parallel	Logical, parallel computing.
as_list	Logical, whether outputs to be a list.

**Value**

A data.frame or list

**Examples**

```
dat = UCICreditCard[24:26]
num_x_list = get_names(dat = dat, types = c('numeric', 'integer', 'double'),
  ex_cols = NULL, get_ex = FALSE)
dat[, num_x_list] = loop_function(func = outliers_kmeans_lof, x_list = num_x_list,
  args = list(dat = dat),
  bind = "cbind", as_list = FALSE,
  parallel = FALSE)
```

---

love\_color

*love\_color*

---

**Description**

love\_color is for get plots for a variable.

**Usage**

```
love_color(color = NULL, type = NULL, all = FALSE)
```

**Arguments**

color	The name of colors.
type	The type of colors, "deep".
all	all of colors.

**Examples**

```
love_color(color="dark_cyan")
```

---

**low\_variance\_filter**     *Filtering Low Variance Variables*

---

**Description**

low\_variance\_filter is for removing variables with repeated values up to a certain percentage.

**Usage**

```
low_variance_filter(dat, lvp = 0.97, only_NA = FALSE, note = FALSE,  
  ex_cols = NULL)
```

**Arguments**

dat	A data frame with x and target.
lvp	The maximum percent of unique values (including NAs).
only_NA	Logical, only process variables which NA's rate are more than lvp.
note	Logical.Outputs info.Default is TRUE.
ex_cols	A list of excluded variables. Default is NULL.

**Value**

A data.frame

**Examples**

```
dat = low_variance_filter(lendingclub[1:1000, ], lvp = 0.9)
```

---

**lr\_params**     *Logistic Regression & Scorecard Parameters*

---

**Description**

lr\_params is the list of parameters to train a LR model or Scorecard using in [training\\_model](#).  
lr\_params\_search is for searching the optimal parameters of logistic regression,if any parameters of params in [lr\\_params](#) is more than one.

**Usage**

```
lr_params(tree_control = list(p = 0.02, cp = 0.00000001, xval = 5,
  maxdepth = 10), bins_control = list(bins_num = 10, bins_pct = 0.05,
  b_chi = 0.02, b_odds = 0.1, b_psi = 0.03, b_or = 0.15, mono = 0.2,
  odds_psi = 0.15, kc = 1), f_eval = "ks", best_lambda = "lambda.ks",
  method = "random_search", iters = 10, lasso = TRUE,
  step_wise = TRUE, score_card = TRUE, sp_values = NULL,
  forced_in = NULL, obsweight = c(1, 1), thresholds = list(cor_p =
  0.8, iv_i = 0.02, psi_i = 0.1, cos_i = 0.5), ...)
```

```
lr_params_search(method = "random_search", dat_train, target,
  dat_test = NULL, occur_time = NULL, x_list = NULL, prop = 0.7,
  iters = 10, tree_control = list(p = 0.02, cp = 0, xval = 1, maxdepth
  = 10), bins_control = list(bins_num = 10, bins_pct = 0.02, b_chi =
  0.02, b_odds = 0.1, b_psi = 0.05, b_or = 0.1, mono = 0.1, odds_psi =
  0.03, kc = 1), thresholds = list(cor_p = 0.8, iv_i = 0.02, psi_i = 0.1,
  cos_i = 0.6), step_wise = FALSE, lasso = FALSE, f_eval = "ks")
```

**Arguments**

tree_control	the list of parameters to control cutting initial breaks by decision tree. See details at: <a href="#">get_tree_breaks</a>
bins_control	the list of parameters to control merging initial breaks. See details at: <a href="#">select_best_breaks</a> , <a href="#">select_best</a>
f_eval	Customized evaluation function, "ks" & "auc" are available.
best_lambda	Methods of best lambda standards using to filter variables by LASSO. There are 3 methods: ("lambda.auc", "lambda.ks", "lambda.sim_sign") . Default is "lambda.auc".
method	Method of searching optimal parameters. "random_search", "grid_search", "local_search" are available.
iters	Number of iterations of "random_search" optimal parameters.
lasso	Logical, if TRUE, variables filtering by LASSO. Default is TRUE.
step_wise	Logical, stepwise method. Default is TRUE.
score_card	Logical, transfer woe to a standard scorecard. If TRUE, Output scorecard, and score prediction, otherwise output probability. Default is TRUE.
sp_values	Values will be in separate bins.e.g. list(-1, "missing") means that -1 & missing as special values.Default is NULL.
forced_in	Names of forced input variables. Default is NULL.
obsweight	An optional vector of 'prior weights' to be used in the fitting process. Should be NULL or a numeric vector. If you oversample or cluster different datasets to training the LR model, you need to set this parameter to ensure that the probability of logistic regression output is the same as that before oversampling or segmentation. e.g.:There are 10,000 0 obs and 500 1 obs before oversampling or under-sampling, 5,000 0 obs and 3,000 1 obs after oversampling. Then this parameter should be set to c(10000/5000, 500/3000). Default is NULL..
thresholds	Thresholds for selecting variables.

- cor\_p The maximum threshold of correlation. Default: 0.8.
- iv\_i The minimum threshold of IV. 0.01 to 0.1 usually work. Default: 0.02
- psi\_i The maximum threshold of PSI. 0.1 to 0.3 usually work. Default: 0.1.
- cos\_i cos\_similarity of positive rate of train and test. 0.7 to 0.9 usually work. Default: 0.5.

...	Other parameters
dat_train	data.frame of train data. Default is NULL.
target	name of target variable.
dat_test	data.frame of test data. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place. Default is NULL.
x_list	names of independent variables. Default is NULL.
prop	Percentage of train-data after the partition. Default: 0.7.

**Value**

A list of parameters.

**See Also**

[training\\_model](#), [xgb\\_params](#), [gbm\\_params](#), [rf\\_params](#)

---

lr_vif	<i>Variance-Inflation Factors</i>
--------	-----------------------------------

---

**Description**

lr\_vif is for calculating Variance-Inflation Factors.

**Usage**

```
lr_vif(lr_model)
```

**Arguments**

lr\_model      An object of logistic model.

## Examples

```
sub = cv_split(UCICreditCard, k = 30)[[1]]
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
dat = re_name(UCICreditCard[sub,], "default.payment.next.month", "target")
dat = dat[,c("target",x_list)]

dat = data_cleansing(dat, miss_values = list("", -1))

train_test <- train_test_split(dat, prop = 0.7)
dat_train = train_test$train
dat_test = train_test$test

Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))
lr_vif(lr_model)
get_logistic_coef(lr_model)
class(dat)
mod = lr_model
lr_vif(lr_model)
```

---

max\_min\_norm

*Max Min Normalization*

---

## Description

max\_min\_norm is for normalizing each column vector of matrix 'x' using max\_min normalization

## Usage

```
max_min_norm(x)
```

## Arguments

x                    Vector

## Value

Normalized vector

## Examples

```
dat_s = apply(UCICreditCard[,12:14], 2, max_min_norm)
```

---

merge_category	<i>Merge Category</i>
----------------	-----------------------

---

**Description**

merge\_category is for merging category of nominal variables which number of categories is more than m or percent of samples in any categories is less than p.

**Usage**

```
merge_category(dat, char_list = NULL, ex_cols = NULL, m = 30,
              note = TRUE)
```

**Arguments**

dat	A data frame with x and target.
char_list	The list of characteristic variables that need to merge categories, Default is NULL. In case of NULL,merge categories for all variables of string type.
ex_cols	A list of excluded variables. Default is NULL.
m	The minimum number of categories.
note	Logical, outputs info. Default is TRUE.

**Value**

A data.frame with merged category variables.

**Examples**

```
#merge_catagory
dat = merge_category(lendingclub,ex_cols = "id$_d$")
char_list = get_names(dat = dat,types = c('factor', 'character'),
ex_cols = "id$_d$", get_ex = FALSE)
str(dat[,char_list])
```

---

min_max_norm	<i>Min Max Normalization</i>
--------------	------------------------------

---

**Description**

min\_max\_norm is for normalizing each column vector of matrix 'x' using min\_max normalization

**Usage**

```
min_max_norm(x)
```

**Arguments**

x                    Vector

**Value**

Normalized vector

**Examples**

```
dat_s = apply(UCICreditCard[,12:14], 2, min_max_norm)
```

---

model_result_plot	<i>model result plots model_result_plot is a wrapper of following: perf_table is for generating a model performance table. ks_plot is for K-S. roc_plot is for ROC. lift_plot is for Lift Chart. score_distribution_plot is for plotting the score distribution.</i>
-------------------	--

---

**Description**

model result plots model\_result\_plot is a wrapper of following: perf\_table is for generating a model performance table. ks\_plot is for K-S. roc\_plot is for ROC. lift\_plot is for Lift Chart. score\_distribution\_plot is for plotting the score distribution.

performance table

ks\_plot

lift\_plot

roc\_plot

score\_distribution\_plot

**Usage**

```
model_result_plot(train_pred, score, target, test_pred = NULL,
  gtitle = NULL, perf_dir_path = NULL, save_data = FALSE,
  plot_show = TRUE, total = TRUE, g = 10, cut_bin = "equal_depth")
```

```
perf_table(train_pred, test_pred = NULL, target = NULL, score = NULL,
  g = 10, cut_bin = "equal_depth", breaks = NULL,
  pos_flag = list("1", "1", "Bad", 1), total = FALSE)
```

```
ks_plot(train_pred, test_pred = NULL, target = NULL, score = NULL,
  gtitle = NULL, breaks = NULL, g = 10, cut_bin = "equal_depth",
  perf_tb = NULL)
```

```
lift_plot(train_pred, test_pred = NULL, target = NULL, score = NULL,
  gtitle = NULL, breaks = NULL, g = 10, cut_bin = "equal_depth",
  perf_tb = NULL)
```



```
roc_plot(train_pred, test_pred = NULL, target = NULL, score = NULL,
         gtitle = NULL)
```

```
score_distribution_plot(train_pred, test_pred, target, score,
                       gtitle = NULL, breaks = NULL, g = 10, cut_bin = "equal_depth",
                       perf_tb = NULL)
```

### Arguments

train_pred	A data frame of training with predicted prob or score.
score	The name of prob or score variable.
target	The name of target variable.
test_pred	A data frame of validation with predict prob or score.
gtitle	The title of the graph & The name for periodically saved graphic file.
perf_dir_path	The path for periodically saved graphic files.
save_data	Logical, save results in locally specified folder. Default is FALSE.
plot_show	Logical, show model performance in current graphic device. Default is TRUE.
total	Whether to summarize the table. default: TRUE.
g	Number of breaks for prob or score.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
breaks	Splitting points of prob or score.
pos_flag	The value of positive class of target variable, default: "1".
perf_tb	Performance table.

### Examples

```
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
dat = data_cleansing(dat, target = "target", obs_id = "ID", x_list = x_list,
                    occur_time = "apply_date", miss_values = list("", -1))
dat = process_nas(dat, default_miss = TRUE)
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
                              occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))

dat_train$pred_LR = round(predict(lr_model, dat_train[, x_list], type = "response"), 5)
dat_test$pred_LR = round(predict(lr_model, dat_test[, x_list], type = "response"), 5)
# model evaluation
```

```

perf_table(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
ks_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
roc_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
#lift_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
#score_distribution_plot(train_pred = dat_train, test_pred = dat_test,
#target = "target", score = "pred_LR")
#model_result_plot(train_pred = dat_train, test_pred = dat_test,
#target = "target", score = "pred_LR")

```

---

multi\_grid

*Arrange list of plots into a grid*


---

### Description

Plot multiple ggplot-objects as a grid-arranged single plot.

### Usage

```
multi_grid(..., grobs = list(...), nrow = NULL, ncol = NULL)
```

### Arguments

...	Other parameters.
grobs	A list of ggplot-objects to be arranged into the grid.
nrow	Number of rows in the plot grid.
ncol	Number of columns in the plot grid.

### Details

This function takes a list of ggplot-objects as argument. Plotting functions of this package that produce multiple plot objects (e.g., when there is an argument `facet.grid`) usually return multiple plots as list.

### Value

An object of class `gtable`.

### Examples

```

library(ggplot2)
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
occur_time = "apply_date", miss_values = list("", -1))
dat = process_nas(dat)
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
occur_time = "apply_date")

```

```

dat_train = train_test$train
dat_test = train_test$test
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))

dat_train$pred_LR = round(predict(lr_model, dat_train[, x_list], type = "response"), 5)
dat_test$pred_LR = round(predict(lr_model, dat_test[, x_list], type = "response"), 5)
# model evaluation
p1 = ks_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
p2 = roc_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
p3 = lift_plot(train_pred = dat_train, test_pred = dat_test, target = "target", score = "pred_LR")
p4 = score_distribution_plot(train_pred = dat_train, test_pred = dat_test,
target = "target", score = "pred_LR")
p_plots= multi_grid(p1,p2,p3,p4)
plot(p_plots)

```

---

multi\_left\_join

*multi\_left\_join*


---

## Description

multi\_left\_join is for left join a list of datasets fast.

## Usage

```
multi_left_join(..., df_list = list(...), key_dt = NULL, by = NULL)
```

## Arguments

...	Datasets need join
df_list	A list of datasets.
key_dt	Name or index of Key table to left join.
by	Name of Key columns to join.

## Examples

```
multi_left_join(UCICreditCard[1:10, 1:10], UCICreditCard[1:10, c(1,8:14)],
UCICreditCard[1:10, c(1,20:25)], by = "ID")
```

---

null_blank_na	<i>Encode NAs</i>
---------------	-------------------

---

**Description**

null\_blank\_na is the function to replace null ,NULL, blank or other missing vaules with NA.

**Usage**

```
null_blank_na(dat, miss_values = NULL, note = FALSE)
```

**Arguments**

dat	A data frame with x and target.
miss_values	Other extreme value might be used to represent missing values, e.g: -9999, -9998. These miss_values will be encoded to -1 or "missing".
note	Logical.Outputs info.Default is TRUE.

**Value**

A data.frame

**Examples**

```
datss = null_blank_na(dat = UCICreditCard[1:1000, ], miss_values =list(-1,-2))
```

---

one_hot_encoding	<i>One-Hot Encoding</i>
------------------	-------------------------

---

**Description**

one\_hot\_encoding is for converting the factor or character variables into multiple columns

**Usage**

```
one_hot_encoding(dat, cat_vars = NULL, ex_cols = NULL,
  merge_cat = TRUE, na_act = TRUE, note = FALSE)
```

**Arguments**

dat	A dat frame.
cat_vars	The name or Column index list to be one_hot encoded.
ex_cols	Variables to be excluded, use regular expression matching
merge_cat	Logical. If TRUE, to merge categories greater than 8, default is TRUE.
na_act	Logical,If true, the missing value is processed, if FALSE missing value is omitted .
note	Logical.Outputs info.Default is TRUE.

**Value**

A dat frame with the one hot encoding applied to all the variables with type as factor or character.

**See Also**

[de\\_one\\_hot\\_encoding](#)

**Examples**

```
dat1 = one_hot_encoding(dat = UCICreditCard,
  cat_vars = c("SEX", "MARRIAGE"),
  merge_cat = TRUE, na_act = TRUE)
dat2 = de_one_hot_encoding(dat_one_hot = dat1,
  cat_vars = c("SEX", "MARRIAGE"), na_act = FALSE)
```

---

outliers_detection	<i>Outliers Detection outliers_detection is for outliers detecting using Kmeans and Local Outlier Factor (lof)</i>
--------------------	--

---

**Description**

Outliers Detection outliers\_detection is for outliers detecting using Kmeans and Local Outlier Factor (lof)

**Usage**

```
outliers_detection(dat, x, kc = 3, kn = 5)
```

**Arguments**

dat	A data.frame with independent variables.
x	The name of variable to process.
kc	Number of clustering centers for Kmeans
kn	Number of neighbors for LOF.

**Value**

Outliers of each variable.

---

 partial\_dependence\_plot

*partial\_dependence\_plot*


---

## Description

partial\_dependence\_plot is for generating a partial dependence plot. get\_partial\_dependence\_plots is for plotting partial dependence of all variables in x\_list.

## Usage

```
partial_dependence_plot(model, x, x_train, n.trees = NULL)
```

```
get_partial_dependence_plots(model, x_train, x_list, n.trees = NULL,
  dir_path = getwd(), save_data = TRUE, plot_show = FALSE,
  parallel = FALSE)
```

## Arguments

model	A data frame of training with predicted prob or score.
x	The name of an independent variable.
x_train	A data.frame with independent variables.
n.trees	Number of trees for best.iter of gbm.
x_list	Names of independent variables.
dir_path	The path for periodically saved graphic files.
save_data	Logical, save results in locally specified folder. Default is FALSE.
plot_show	Logical, show model performance in current graphic device. Default is FALSE.
parallel	Logical, parallel computing. Default is FALSE.

## Examples

```
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
  occur_time = "apply_date", miss_values = list("", -1))

train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))
#plot partial dependency of one variable
```

```
partial_dependence_plot(model = lr_model, x = "LIMIT_BAL", x_train = dat_train)
#plot partial dependency of all variables
pd_list = get_partial_dependence_plots(model = lr_model, x_list = x_list[1:2],
  x_train = dat_train, save_data = FALSE, plot_show = TRUE)
```

---

PCA\_reduce

*PCA Dimension Reduction*


---

### Description

PCA\_reduce is used for PCA reduction of high demension data .

### Usage

```
PCA_reduce(train = train, test = NULL, mc = 0.9)
```

### Arguments

train	A data.frame with independent variables and target variable.
test	A data.frame of test data.
mc	Threshold of cumulative imp.

### Examples

```
## Not run:
num_x_list = get_names(dat = UCICreditCard, types = c('numeric'),
  ex_cols = "ID$|date$|default.payment.next.month$", get_ex = FALSE)
  PCA_dat = PCA_reduce(train = UCICreditCard[num_x_list])

## End(Not run)
```

---

plot\_bar

*Plot Bar*


---

### Description

You can use the plot\_bar to produce the barplot.

### Usage

```
plot_bar(dat, x, breaks = NULL, g = 10, cut_bin = "equal_width")
```

**Arguments**

dat	A data.frame with independent variables and target variable.
x	The name of an independent variable.
breaks	Breaks points of x.
g	Number of initial breakpoints for equal frequency binning.
cut_bin	'equal_width' or 'equal_depth'

**Examples**

```
plot_bar(dat = lendingclub, x = "grade")
```

---

plot\_box

*Plot Box*


---

**Description**

You can use the plot\_box to produce boxplot.

**Usage**

```
plot_box(dat, y, x = NULL, g = 5, colors_x = c(love_color(type =
  "deep"), love_color(type = "light"), love_color(type = "shallow"),
  love_color(type = "dark")))
```

**Arguments**

dat	A data.frame with independent variables and target variable.
y	The name of target variable.
x	The name of an independent variable.
g	Number of initial breakpoints for equal frequency binning.
colors_x	colors of x.

**Examples**

```
plot_box(lendingclub, x = "grade", y = "installment", g = 7)
```



---

plot_density	<i>Plot Density</i>
--------------	---------------------

---

**Description**

You can use the `plot_density` to produce plots that characterize the density.

**Usage**

```
plot_density(dat, x, y = NULL, m = 3, g = 5, y_breaks = NULL,
             binwidth = NULL, hist = TRUE, colors_y = c(love_color(type =
"deep"), love_color(type = "light"), love_color(type = "shallow")))
```

**Arguments**

<code>dat</code>	A data.frame with independent variables and target variable.
<code>x</code>	The name of an independent variable.
<code>y</code>	The name of target variable.
<code>m</code>	The outlier cutoff.
<code>g</code>	Number of initial breakpoints for equal frequency binning.
<code>y_breaks</code>	Breaks points of y.
<code>binwidth</code>	Windth of bins for histogram.
<code>hist</code>	If plot the histogram.
<code>colors_y</code>	colors of y.

**Examples**

```
plot_density(dat = lendingclub, x = "annual_inc", y = "emp_length", m = 0, hist = FALSE)
plot_density(dat = lendingclub, x = "annual_inc", m = 2,
             colors_y = love_color(type = "line")[c(1,3)])
```

---

plot_distribution	<i>Plot Distribution</i>
-------------------	--------------------------

---

**Description**

You can use the `plot_distribution_x` to produce the distribution plot of a variable. You can use the `plot_distribution` to produce the plots of distributions of all variables.

**Usage**

```
plot_distribution(dat, x_list = NULL, dir_path = tempdir(),
                 breaks_list = NULL, g = 10, m = 3, cut_bin = "equal_width")

plot_distribution_x(dat, x, breaks = NULL, g = 10, m = 3,
                  cut_bin = "equal_width", binwidth = NULL)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
x_list	Names of independent variables.
dir_path	The path for periodically saved graphic files.
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
g	Number of initial breakpoints for equal frequency binning.
m	The outlier cutoff.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
x	The name of an independent variable.
breaks	Splitting points for an independent variable. Default is NULL.
binwidth	Windth of bins for histogram.

**Examples**

```
plot_distribution_x(dat = lendingclub, x = "max_bal_bc", g = 10,
  cut_bin = 'equal_width')
plot_distribution(dat = lendingclub, x_list = c("max_bal_bc", "installment"),
  g = 10, dir_path = tempdir(),
  cut_bin = 'equal_width')
```

---

plot_oot_perf	<i>plot_oot_perf</i> plot_oot_perf is for plotting performance of cross time samples in the future
---------------	--

---

**Description**

plot\_oot\_perf plot\_oot\_perf is for plotting performance of cross time samples in the future

**Usage**

```
plot_oot_perf(dat_test, x, occur_time, target, k = 3, g = 10,
  period = "month", best = FALSE, equal_bins = TRUE, pl = "rate",
  breaks = NULL, cut_bin = "equal_depth", gtitle = NULL,
  perf_dir_path = NULL, save_data = FALSE, plot_show = TRUE)
```

**Arguments**

dat_test	A data frame of testing dataset with predicted prob or score.
x	The name of prob or score variable.
occur_time	The name of the variable that represents the time at which each observation takes place.
target	The name of target variable.

k	If period is NULL, number of equal frequency samples.
g	Number of breaks for prob or score.
period	OOT period, 'weekly' and 'month' are available.if NULL, use k equal frequency samples.
best	Logical, merge initial breaks to get optimal breaks for binning.
equal_bins	Logical, generates initial breaks for equal frequency or width binning.
pl	'lift' is for lift chart plot,'rate' is for positive rate plot.
breaks	Splitting points of prob or score.
cut_bin	A string, if equal_bins is TRUE, 'equal_depth' or 'equal_width', default is 'equal_depth'.
gtitle	The title of the graph & The name for periodically saved graphic file.
perf_dir_path	The path for periodically saved graphic files.
save_data	Logical, save results in locally specified folder. Default is FALSE.
plot_show	Logical, show model performance in current graphic device. Default is TRUE.

### Examples

```

sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "PAY_3", "PAY_2")
dat = data_cleansing(dat, target = "target", obs_id = "ID", x_list = x_list,
  occur_time = "apply_date", miss_values = list("", -1))
dat = process_nas(dat)
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = dat_train[, c("target", x_list)], family = binomial(logit))

dat_train$pred_LR = round(predict(lr_model, dat_train[, x_list], type = "response"), 5)
dat_test$pred_LR = round(predict(lr_model, dat_test[, x_list], type = "response"), 5)
plot_oot_perf(dat_test = dat_test, occur_time = "apply_date", target = "target", x = "pred_LR")

```

---

plot\_relative\_freq\_histogram

*Plot Relative Frequency Histogram*

---

### Description

You can use the plot\_relative\_freq\_histogram to produce the relative frequency histogram plots.

**Usage**

```
plot_relative_freq_histogram(dat, x, y = NULL, x_breaks = NULL,
  y_breaks = NULL, g = 10, cut_bin = "equal_width")
```

**Arguments**

dat	A data.frame with independent variables and target variable.
x	The name of an independent variable.
y	The name of target variable. Default is NULL.
x_breaks	Breaks points of x.
y_breaks	Breaks points of y.
g	Number of initial breakpoints for equal frequency binning.
cut_bin	'equal_width' or 'equal_depth' to produce the breaks points.

**Examples**

```
plot_relative_freq_histogram(dat = lendingclub, x = "grade", y = "dti_joint", g = 7,
  cut_bin = 'equal_width')
```

---

plot\_table

*plot\_table*


---

**Description**

plot\_table is for table visualizaiton.

**Usage**

```
plot_table(grid_table, theme = c("cyan", "grey", "green", "red", "blue",
  "purple"), title = NULL, title.size = 12, title.color = "black",
  title.face = "bold", title.position = "middle", subtitle = NULL,
  subtitle.size = 8, subtitle.color = "black",
  subtitle.face = "plain", subtitle.position = "middle",
  tile.color = "white", tile.size = 1, colname.size = 3,
  colname.color = "white", colname.face = "bold",
  colname.fill.color = love_color("dark_cyan"), text.size = 3,
  text.color = love_color("dark_grey"), text.face = "plain",
  text.fill.color = c("white", love_color("pale_grey"))))
```

**Arguments**

grid_table	A data.frame or table
theme	The theme of color, "cyan","grey","green","red","blue","purple" are available.
title	The title of table
title.size	The title size of plot.
title.color	The title color.
title.face	The title face, such as "plain", "bold".
title.position	The title position,such as "left","middle","right".
subtitle	The subtitle of table
subtitle.size	The subtitle size.
subtitle.color	The subtitle color.
subtitle.face	The subtitle face, such as "plain", "bold",default is "bold".
subtitle.position	The subtitle position,such as "left","middle","right", default is "middle".
tile.color	The color of table lines, default is 'white'.
tile.size	The size of table lines , default is 1.
colname.size	The size of colnames, default is 3.
colname.color	The color of colnames, default is 'white'.
colname.face	The face of colnames,default is 'bold'.
colname.fill.color	The fill color of colnames, default is love_color("dark_cyan").
text.size	The size of text, default is 3.
text.color	The color of text, default is love_color("dark_grey").
text.face	The face of text, default is 'plain'.
text.fill.color	The fill color of text, default is c('white',love_color("pale_grey")).

**Examples**

```

iv_list = get_psi_iv_all(dat = UCICreditCard[1:1000, ],
                        x_list = names(UCICreditCard)[3:5], equal_bins = TRUE,
                        target = "default.payment.next.month", ex_cols = "ID|apply_date")
iv_dt =get_psi_iv(UCICreditCard, x = "PAY_3",
                  target = "default.payment.next.month", bins_total = TRUE)

plot_table(iv_dt)

```

---

plot_theme	<i>plot_theme</i>
------------	-------------------

---

## Description

plot\_theme is a simpler wrapper of theme for ggplot2.

## Usage

```
plot_theme(legend.position = "top", angle = 30, legend_size = 7,  
  axis_size_y = 8, axis_size_x = 8, axis_title_size = 10,  
  title_size = 11, title_vjust = 0, title_hjust = 0,  
  linetype = "dotted", face = "bold")
```

## Arguments

legend.position	see details at: codelegend.position
angle	see details at: codeaxis.text.x
legend_size	see details at: codelegend.text
axis_size_y	see details at: codeaxis.text.y
axis_size_x	see details at: codeaxis.text.x
axis_title_size	see details at: codeaxis.title.x
title_size	see details at: codeplot.title
title_vjust	see details at: codeplot.title
title_hjust	see details at: codeplot.title
linetype	see details at: codepanel.grid.major
face	see details at: codeaxis.title.x

## Details

see details at: codetheme

---

pred_score	<i>pred_score</i>
------------	-------------------

---

**Description**

pred\_score is for using logistic regression model model to predict new data.

**Usage**

```
pred_score(model, dat, x_list = NULL, bins_table = NULL,
  obs_id = NULL, miss_values = list(-1, "-1", "NULL", "-1", "-9999",
  "-9996", "-9997", "-9995", "-9998", -9999, -9998, -9997, -9996, -9995),
  woe_name = TRUE)
```

**Arguments**

model	Logistic Regression Model generated by <a href="#">training_model</a> .
dat	Dataframe of new data.
x_list	Into the model variables.
bins_table	a data.frame generated by <a href="#">get_bins_table</a>
obs_id	The name of ID of observations or key variable of data. Default is NULL.
miss_values	Special values.
woe_name	Logical. Whether woe variable's name contains 'woe'.Default is TRUE.

**Value**

new scores.

**See Also**

[training\\_model](#), [lr\\_params](#), [xgb\\_params](#), [rf\\_params](#)

---

pred_xgb	<i>pred_xgb</i>
----------	-----------------

---

**Description**

pred\_xgb is for using xgboost model to predict new data.

**Usage**

```
pred_xgb(xgb_model = NULL, dat, x_list = NULL, miss_values = NULL,
  model_name = NULL, model_path = getwd())
```

**Arguments**

xgb_model	XGboost Model generated by <a href="#">training_model</a> .
dat	Dataframe of new data.
x_list	Into the model variables.
miss_values	missing values.
model_name	Name of model
model_path	dir_path of model.

**Value**

new prob.

**See Also**

[training\\_model](#), [pred\\_score](#)

---

process_nas	<i>missing Treatment</i>
-------------	--------------------------

---

**Description**

process\_nas\_var is for missing value analysis and treatment using knn imputation, central imputation and random imputation. process\_nas is a simpler wrapper for process\_nas\_var.

**Usage**

```
process_nas(dat, x_list = NULL, class_var = FALSE,
  miss_values = list(-1, "missing"), default_miss = list(-1,
  "missing"), parallel = FALSE, ex_cols = NULL, method = "median",
  note = FALSE, save_data = FALSE, file_name = NULL,
  dir_path = tempdir(), ...)
```

```
process_nas_var(dat = dat, x, nas_rate = NULL,
  default_miss = list("missing", -1), mat_nas_shadow = NULL,
  dt_nas_random = NULL, missing_type = NULL, method = "median",
  note = FALSE, save_data = FALSE, file_name = NULL,
  dir_path = tempdir(), ...)
```

**Arguments**

dat	A data.frame with independent variables.
x_list	Names of independent variables.
class_var	Logical, nas analysis of the nominal variables. Default is TRUE.
miss_values	Other extreme value might be used to represent missing values, e.g:-1, -9999, -9998. These miss_values will be encoded to NA.



default_miss	Default value of missing data imputation, Default is list(-1,'missing').
parallel	Logical, parallel computing. Default is FALSE.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
method	The methods of imputation by knn. If "median", then Nas imputation with k neighbors median. If "avg_dist", the distance weighted average method is applied to determine the NAs imputation with k neighbors. If "default", assigning the missing values to -1 or "missing", otherwise ,processing the missing values according to the results of missing analysis.
note	Logical, outputs info. Default is TRUE.
save_data	Logical. If TRUE, save missing analysis to dir_path
file_name	The file name for periodically saved missing analysis file. Default is NULL.
dir_path	The path for periodically saved missing analysis file. Default is "./variable".
...	Other parameters.
x	The name of variable to process.
nas_rate	A list contains nas rate of each variable.
mat_nas_shadow	A shadow matrix of variables which contain nas.
dt_nas_random	A data.frame with random nas imputation.
missing_type	Type of missing, generated by code <a href="#">analysis_nas</a>

**Value**

A dat frame with no NAs.

**Examples**

```
dat_na = process_nas(dat = UCICreditCard[1:1000,],
parallel = FALSE,ex_cols = "ID$", method = "median")
```

---

process\_outliers      *Outliers Treatment*

---

**Description**

outliers\_kmeans\_lof is for outliers detection and treatment using Kmeans and Local Outlier Factor (lof) process\_outliers is a simpler wrapper for outliers\_kmeans\_lof.

**Usage**

```
process_outliers(dat, target, ex_cols = NULL, kc = 3, kn = 5,
x_list = NULL, parallel = FALSE, note = FALSE, process = TRUE,
save_data = FALSE, file_name = NULL, dir_path = tempdir())
```

```
outliers_kmeans_lof(dat, x, target = NULL, kc = 3, kn = 5,
note = FALSE, process = TRUE, save_data = FALSE,
file_name = NULL, dir_path = tempdir())
```

**Arguments**

dat	Dataset with independent variables and target variable.
target	The name of target variable.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
kc	Number of clustering centers for Kmeans
kn	Number of neighbors for LOF.
x_list	Names of independent variables.
parallel	Logical, parallel computing.
note	Logical, outputs info. Default is TRUE.
process	Logical, process outliers, not just analysis.
save_data	Logical. If TRUE, save outliers analysis file to the specified folder at dir_path
file_name	The file name for periodically saved outliers analysis file. Default is NULL.
dir_path	The path for periodically saved outliers analysis file. Default is "./variable".
x	The name of variable to process.

**Value**

A data frame with outliers process to all the variables.

**Examples**

```
dat_out = process_outliers(UCICreditCard[1:10000,c(18:21,26)],
  target = "default.payment.next.month",
  ex_cols = "date$", kc = 3, kn = 10,
  parallel = FALSE,note = TRUE)
```

---

psi_iv_filter	<i>Variable reduction based on Information Value &amp; Population Stability Index filter</i>
---------------	--

---

**Description**

psi\_iv\_filter is for selecting important and stable features using IV & PSI.

**Usage**

```
psi_iv_filter(dat, dat_test = NULL, target, x_list = NULL,
  breaks_list = NULL, pos_flag = NULL, ex_cols = NULL,
  occur_time = NULL, best = FALSE, equal_bins = TRUE, g = 10,
  sp_values = NULL, tree_control = list(p = 0.05, cp = 0.000001, xval =
  5, maxdepth = 10), bins_control = list(bins_num = 10, bins_pct = 0.05,
  b_chi = 0.05, b_odds = 0.1, b_psi = 0.05, b_or = 0.15, mono = 0.3,
  odds_psi = 0.2, kc = 1), oot_pct = 0.7, psi_i = 0.1, iv_i = 0.01,
  cos_i = 0.7, vars_name = FALSE, note = TRUE, parallel = FALSE,
  save_data = FALSE, file_name = NULL, dir_path = tempdir(), ...)
```

**Arguments**

dat	A data.frame with independent variables and target variable.
dat_test	A data.frame of test data. Default is NULL.
target	The name of target variable.
x_list	Names of independent variables.
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
pos_flag	The value of positive class of target variable, default: "1".
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
best	Logical, if TRUE, merge initial breaks to get optimal breaks for binning.
equal_bins	Logical, if TRUE, equal sample size initial breaks generates.If FALSE , tree breaks generates using desision tree.
g	Integer, number of initial bins for equal_bins.
sp_values	A list of missing values.
tree_control	the list of tree parameters.
bins_control	the list of parameters.
oot_pct	Percentage of observations retained for overtime test (especially to calculate PSI). Default is 0.7
psi_i	The maximum threshold of PSI. $0 \leq \text{psi}_i \leq 1$ ; 0.05 to 0.2 usually work. Default: 0.1
iv_i	The minimum threshold of IV. $0 < \text{iv}_i$ ; 0.01 to 0.1 usually work. Default: 0.01
cos_i	cos_similarity of posive rate of train and test. 0.7 to 0.9 usually work.Default: 0.5.
vars_name	Logical, output a list of filtered variables or table with detailed IV and PSI value of each variable. Default is FALSE.
note	Logical, outputs info. Default is TRUE.
parallel	Logical, parallel computing. Default is FALSE.
save_data	Logical, save results in locally specified folder. Default is FALSE.
file_name	The name for periodically saved results files. Default is "Feature_importance_IV_PSI".
dir_path	The path for periodically saved results files. Default is tempdir().
...	Other parameters.

**Value**

A list with the following elements:

- Feature Selected variables.
- IV IV of variables.
- PSI PSI of variables.
- COS cos\_similarity of posive rate of train and test.

**See Also**

[xgb\\_filter](#), [gbm\\_filter](#), [feature\\_selector](#)

**Examples**

```
psi_iv_filter(dat= UCICreditCard[1:1000,c(2,4,8:9,26)],
              target = "default.payment.next.month",
              occur_time = "apply_date",
              parallel = FALSE)
```

---

p_ij	<i>Entropy</i>
------	----------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

p\_ij(x)

e\_ij(x)

**Arguments**

x                    A numeric vector.

**Value**

A numeric vector of entropy.

---

p_to_score	<i>prob to score</i>
------------	----------------------

---

**Description**

p\_to\_score is for transforming probability to score.

**Usage**

```
p_to_score(p, PDO = 20, base = 600, ratio = 1)
```

**Arguments**

p	Probability.
PDO	Point-to-Double Odds.
base	Base Point.
ratio	The corresponding odds when the score is base.

**Value**

new prob.

**See Also**

[training\\_model](#), [pred\\_score](#)

---

quick\_as\_df                      *List as data.frame quickly*

---

**Description**

quick\_as\_df is function for fast dat frame transfromation.

**Usage**

```
quick_as_df(df_list)
```

**Arguments**

df\_list                      A list of data.

**Value**

packages installed and library,

**Examples**

```
UCICreditCard = quick_as_df(UCICreditCard)
```

---

ranking\_percent\_proc *Ranking Percent Process*

---

### Description

ranking\_percent\_proc is for processing ranking percent variables. ranking\_percent\_dict is for generating ranking percent dictionary.

### Usage

```
ranking_percent_proc(dat, ex_cols = NULL, x_list = NULL,
  rank_dict = NULL, pct = 0.01, parallel = FALSE, note = FALSE,
  save_data = FALSE, file_name = NULL, dir_path = tempdir(), ...)
```

```
ranking_percent_proc_x(dat, x, rank_dict = NULL, pct = 0.01)
```

```
ranking_percent_dict(dat, x_list = NULL, ex_cols = NULL, pct = 0.01,
  parallel = FALSE, save_data = FALSE, file_name = NULL,
  dir_path = tempdir(), ...)
```

```
ranking_percent_dict_x(dat, x = NULL, pct = 0.01)
```

### Arguments

dat	A data.frame.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
x_list	A list of x variables.
rank_dict	The dictionary of rank_percent generated by ranking_percent_dict .
pct	Percent of rank. Default is 0.01.
parallel	Logical, parallel computing. Default is FALSE.
note	Logical, outputs info. Default is TRUE.
save_data	Logical, save results in locally specified folder. Default is FALSE
file_name	The name for periodically saved rank_percent data file. Default is "dat_rank_percent".
dir_path	The path for periodically saved rank_percent data file Default is "tempdir()"
...	Additional parameters.
x	The name of an independent variable.

### Value

Data.frame with new processed variables.

**Examples**

```
rank_dict = ranking_percent_dict(dat = UCICreditCard[1:1000,],
x_list = c("LIMIT_BAL", "BILL_AMT2", "PAY_AMT3"), ex_cols = NULL )
UCICreditCard_new = ranking_percent_proc(dat = UCICreditCard[1:1000,],
x_list = c("LIMIT_BAL", "BILL_AMT2", "PAY_AMT3"), rank_dict = rank_dict, parallel = FALSE)
```

---

read\_data

*Read data*


---

**Description**

read\_data is for loading data, formats like csv, txt, data and so on.

**Usage**

```
read_data(path, pattern = NULL, encoding = "unknown", header = TRUE,
sep = "auto", stringsAsFactors = FALSE, select = NULL,
drop = NULL, nrows = Inf)
```

```
check_data_format(path)
```

**Arguments**

path	Path to file or file name in working directory & path to file.
pattern	An optional regular expression. Only file names which match the regular expression will be returned.
encoding	Default is "unknown". Other possible options are "UTF-8" and "Latin-1".
header	Does the first data line contain column names?
sep	The separator between columns.
stringsAsFactors	Logical. Convert all character columns to factors?
select	A vector of column names or numbers to keep, drop the rest.
drop	A vector of column names or numbers to drop, keep the rest.
nrows	The maximum number of rows to read.

---

 reduce\_high\_cor\_filter

*Filtering highly correlated variables with reduce method*


---

### Description

reduce\_high\_cor\_filter is function for filtering highly correlated variables with reduce method.

### Usage

```
reduce_high_cor_filter(dat, x_list = NULL, size = ncol(dat)/10,
  p = 0.95, com_list = NULL, ex_cols = NULL, cor_class = TRUE,
  parallel = FALSE)
```

### Arguments

dat	A data.frame with independent variables.
x_list	Names of independent variables.
size	Size of vairable group.
p	Threshold of correlation between features. Default is 0.7.
com_list	A data.frame with important values of each variable. eg : IV_list
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
cor_class	Culcate catagery variables's correlation matrix. Default is FALSE.
parallel	Logical, parallel computing. Default is FALSE.

---

remove\_duplicated

*Remove Duplicated Observations*


---

### Description

remove\_duplicated is the function to remove duplicated observations

### Usage

```
remove_duplicated(dat = dat, obs_id = NULL, occur_time = NULL,
  target = NULL, note = FALSE)
```

### Arguments

dat	A data frame with x and target.
obs_id	The name of ID of observations. Default is NULL.
occur_time	The name of occur time of observations.Default is NULL.
target	The name of target variable.
note	Logical.Outputs info.Default is TRUE.



**Value**

A data.frame

**Examples**

```
datss = remove_duplicated(dat = UCICreditCard,
  target = "default.payment.next.month",
  obs_id = "ID", occur_time = "apply_date")
```

---

replace_value	<i>Replace Value</i>
---------------	----------------------

---

**Description**

replace\_value is for replacing values of some variables . replace\_value\_x is for replacing values of a variable.

**Usage**

```
replace_value(dat = dat, x_list = NULL, x_pattern = NULL,
  replace_dat, MARGIN = 2, VALUE = if (MARGIN == 2)
  colnames(replace_dat) else rownames(replace_dat), RE_NAME = TRUE,
  parallel = FALSE)
```

```
replace_value_x(dat, x, replace_dat, MARGIN = 2, VALUE = if (MARGIN ==
  2) colnames(replace_dat) else rownames(replace_dat), RE_NAME = TRUE)
```

**Arguments**

dat	A data.frame.
x_list	Names of variables to replace value.
x_pattern	Regular expressions, used to match variable names.
replace_dat	A data.frame contains value to replace.
MARGIN	A vector giving the subscripts which the function will be applied over. E.g., for a matrix 1 indicates rows, 2 indicates columns, c(1, 2) indicates rows and columns. Where X has named dimnames, it can be a character vector selecting dimension names.
VALUE	Values to replace.
RE_NAME	Logical, rename the replaced variable.
parallel	Logical, parallel computing. Default is TRUE.
x	Name of variable to replace value.

---

require_packages	<i>Packages required and intallment</i>
------------------	---

---

**Description**

require\_packages is function for librarying required packages and installing missing packages if needed.

**Usage**

```
require_packages(..., pkg = as.character(substitute(list(...))))
```

**Arguments**

...	Packages need loaded
pkg	A list or vector of names of required packages.

**Value**

packages installed and library.

**Examples**

```
## Not run:
require_packages(data.table, ggplot2, dplyr)

## End(Not run)
```

---

re_code	<i>re_code re_code search for matches to argument pattern within each element of a character vector:</i>
---------	--

---

**Description**

re\_code re\_code search for matches to argument pattern within each element of a character vector:

**Usage**

```
re_code(x, codes)
```

**Arguments**

x	Variable to recode.
codes	A data.frame of original value & recode value

**Examples**

```
SEX = sample(c("F","M"),1000,replace = TRUE)
codes= data.frame(ori_value = c('F','M'), code = c(0,1) )
SEX_re = re_code(SEX,codes)
```

---

re_name	<i>Rename</i>
---------	---------------

---

**Description**

re\_name is for renaming variables.

**Usage**

```
re_name(dat, oldname = c(), newname = c())
```

**Arguments**

dat	A data frame with vairables to rename.
oldname	Old names of vairables.
newname	New names of vairables.

**Value**

data with new variable names.

**Examples**

```
dt = re_name(dat = UCICreditCard, "default.payment.next.month" , "target")
names(dt['target'])
```

---

rf_params	<i>Random Forest Parameters</i>
-----------	---------------------------------

---

**Description**

rf\_params is the list of parameters to train a Random Forest using in [training\\_model](#).

**Usage**

```
rf_params(ntree = 100, nodesize = 30, samp_rate = 0.5,
  tune_rf = FALSE, ...)
```

**Arguments**

n <sub>tree</sub>	Number of trees to grow. This should not be set to too small a number, to ensure that every input row gets predicted at least a few times.
n <sub>odesize</sub>	Minimum size of terminal nodes. Setting this number larger causes smaller trees to be grown (and thus take less time). Note that the default values are different for classification (1) and regression (5).
samp <sub>rate</sub>	Percentage of sample to draw. Default is 0.2.
tune <sub>rf</sub>	A logical.If TRUE, then tune Random Forest model.Default is FALSE.
...	Other parameters

**Details**

See details at : [https://www.stat.berkeley.edu/~breiman/Using\\_random\\_forests\\_V3.1.pdf](https://www.stat.berkeley.edu/~breiman/Using_random_forests_V3.1.pdf)

**Value**

A list of parameters.

**See Also**

[training\\_model](#), [lr\\_params](#), [gbm\\_params](#), [xgb\\_params](#)

---

rowAny

*Functions for vector operation.*

---

**Description**

Functions for vector operation.

**Usage**

rowAny(x)

rowAllnas(x)

colAllnas(x)

colAllzeros(x)

rowAll(x)

rowCVs(x, na.rm = FALSE)

rowSds(x, na.rm = FALSE)

```
colSds(x, na.rm = TRUE)
rowMaxs(x, na.rm = FALSE)
rowMins(x, na.rm = FALSE)
rowMaxMins(x, na.rm = FALSE)
colMaxMins(x, na.rm = FALSE)
cnt_x(x)
sum_x(x)
max_x(x)
min_x(x)
avg_x(x)
```

### Arguments

x	A data.frame or Matrix.
na.rm	Logical, remove NAs.

### Value

A data.frame or Matrix.

### Examples

```
#any row has missing values
row_amy = rowAny(UCICreditCard[8:10])
#rows which is all missing values
row_na = rowAllnas(UCICreditCard[8:10])
#cols which is all missing values
col_na = colAllnas(UCICreditCard[8:10])
#cols which is all zeros
row_zero = colAllzeros(UCICreditCard[8:10])
#sum all numbers of a row
row_all = rowAll(UCICreditCard[8:10])
#caculate cv of a row
row_cv = rowCVs(UCICreditCard[8:10])
#caculate sd of a row
row_sd = rowSds(UCICreditCard[8:10])
#caculate sd of a column
col_sd = colSds(UCICreditCard[8:10])
```

---

rules_filter	<i>rules_filter</i>
--------------	---------------------

---

### Description

rules\_filter This function is used to filter or select samples by rules.

### Usage

```
rules_filter(dat, rules_list, drop = FALSE, logic = "or")
```

### Arguments

dat	A data.frame.
rules_list	A list of rules.
drop	Logical, if TRUE, dropping samples, if FALSE, selecting samples. Default is FALSE.
logic	The logic between rules in the rules_list: 'and','or'. Default is 'or'.

### Value

A data frame.

### See Also

[get\\_ctree\\_rules](#), [check\\_rules](#)

### Examples

```
train_test <- train_test_split(UCICreditCard, split_type = "Random", prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
dat_train$default.payment.next.month = as.numeric(dat_train$default.payment.next.month)
rules_list = get_ctree_rules(tree_fit = NULL, train_dat = dat_train[, 8:26],
                             target = "default.payment.next.month", test_dat = dat_test)[1:3,2]
new_dat = rules_filter(rules_list = rules_list[3], dat = dat_test)
```

---

rules_result	<i>rules_result</i>
--------------	---------------------

---

## Description

`rules_result` This function is used to get rules results.

## Usage

```
rules_result(dat, rules_list, yes = "reject", no = "pass")
```

## Arguments

<code>dat</code>	A data.frame
<code>rules_list</code>	A list of rules.
<code>yes</code>	Default is 'reject'.
<code>no</code>	Default is 'pass'.

## Value

A vector with 'pass' and 'reject'.

## See Also

[get\\_ctree\\_rules](#), [check\\_rules](#), [rules\\_filter](#)

## Examples

```
train_test <- train_test_split(UCICreditCard, split_type = "Random", prop = 0.8, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
dat_train$default.payment.next.month = as.numeric(dat_train$default.payment.next.month)
rules_list = get_ctree_rules(tree_fit = NULL, train_dat = dat_train[, 8:26],
                             target = "default.payment.next.month", test_dat = dat_test)[1:3,2]
dat_test$rules_result = rules_result(rules_list = rules_list[3], dat = dat_test)
```

---

rule_value_replace	<i>rule_value_replace</i>
--------------------	---------------------------

---

**Description**

rule\_value\_replace is for generating new variables by rules.

**Usage**

```
rule_value_replace(dat, rules_list, VALUE = 1:length(rules_list),
  x_name = "x_level")
```

**Arguments**

dat	A data.frame.
rules_list	Names of variables to replace value.
VALUE	values to replace.
x_name	name of the new variable.

---

save_data	<i>Save data</i>
-----------	------------------

---

**Description**

save\_data is for saving a data.frame or a list fast.

**Usage**

```
save_data(..., files = list(...),
  file_name = as.character(substitute(list(...))), dir_path = getwd(),
  note = FALSE, as_list = FALSE, row_names = FALSE, append = FALSE)
```

**Arguments**

...	datasets
files	A dataset or a list of datasets.
file_name	The file name of data.
dir_path	A string. The dir path to save breaks_list.
note	Logical. Outputs info.Default is TRUE.
as_list	Logical. List format or data.frame format to save. Default is FALSE.
row_names	Logical,retain rownames.
append	Logical, append newdata to old.

**Examples**

```
save_data(UCICreditCard,"UCICreditCard", tempdir())
```



---

score_transfer	<i>Score Transformation</i>
----------------	-----------------------------

---

**Description**

score\_transfer is for transfer woe to score.

**Usage**

```
score_transfer(model, tbl_woe, a = 600, b = 50, file_name = NULL,
  dir_path = tempdir(), save_data = FALSE)
```

**Arguments**

model	A data frame with x and target.
tbl_woe	a data.frame with woe variables.
a	Base line of score.
b	Numeric.Increased scores from doubling Odds.
file_name	The name for periodically saved score file. Default is "dat_score".
dir_path	The path for periodically saved score file. Default is "./data"
save_data	Logical, save results in locally specified folder. Default is FALSE.

**Value**

A data.frame with variables which values transferred to score.

**Examples**

```
# dataset splitting
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
#rename the target variable
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID",
  occur_time = "apply_date", miss_values = list("", -1))
#train_ test pliting
train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
  occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
  x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
  save_data = FALSE, note = FALSE)
#woe transforming
train_woe = woe_trans_all(dat = dat_train,
```

```

        target = "target",
        breaks_list = breaks_list,
        woe_name = FALSE)
test_woe = woe_trans_all(dat = dat_test,
                        target = "target",
                        breaks_list = breaks_list,
                        note = FALSE)
Formula = as.formula(paste("target", paste(x_list, collapse = ' + '), sep = ' ~ '))
set.seed(46)
lr_model = glm(Formula, data = train_woe[, c("target", x_list)], family = binomial(logit))
#get LR coefficient
dt_imp_LR = get_logistic_coef(lg_model = lr_model, save_data = FALSE)
bins_table = get_bins_table_all(dat = dat_train, target = "target",
                               x_list = x_list, dat_test = dat_test,
                               breaks_list = breaks_list, note = FALSE)

#score card
LR_score_card <- get_score_card(lg_model = lr_model, bins_table, target = "target")
#scoring
train_pred = dat_train[, c("ID", "apply_date", "target")]
test_pred = dat_test[, c("ID", "apply_date", "target")]
train_pred$pred_LR = score_transfer(model = lr_model,
                                   tbl_woe = train_woe,
                                   save_data = FALSE)[, "score"]

test_pred$pred_LR = score_transfer(model = lr_model,
                                   tbl_woe = test_woe, save_data = FALSE)[, "score"]

```

---

select\_best\_class      *Generates Best Binning Breaks*

---

## Description

select\_best\_class & select\_best\_breaks are for merging initial breaks of variables using chi-square, odds-ratio, PSI, G/B index and so on. The get\_breaks is a simpler wrapper for select\_best\_class & select\_best\_class.

## Usage

```
select_best_class(dat, x, target, breaks = NULL, occur_time = NULL,
                 oot_pct = 0.7, pos_flag = NULL, bins_control = NULL,
                 sp_values = NULL, ...)
```

```
select_best_breaks(dat, x, target, breaks = NULL, pos_flag = NULL,
                  sp_values = NULL, occur_time = NULL, oot_pct = 0.7,
                  bins_control = NULL, ...)
```

## Arguments

dat                    A data frame with x and target.

x	The name of variable to process.
target	The name of target variable.
breaks	Splitting points for an independent variable. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place.
oot_pct	The percentage of Actual and Expected set for PSI calculating.
pos_flag	The value of positive class of target variable, default: "1".
bins_control	the list of parameters. <ul style="list-style-type: none"> <li>• bins_num The maximum number of bins. 5 to 10 usually work. Default: 10</li> <li>• bins_pct The minimum percent of observations in any bins. <math>0 &lt; \text{bins\_pct} &lt; 1</math>, 0.01 to 0.1 usually work. Default: 0.02.</li> <li>• b_chi The minimum threshold of chi-square merge. <math>0 &lt; \text{b\_chi} &lt; 1</math>; 0.01 to 0.1 usually work. Default: 0.02.</li> <li>• b_odds The minimum threshold of odds merge. <math>0 &lt; \text{b\_odds} &lt; 1</math>; 0.05 to 0.2 usually work. Default: 0.1.</li> <li>• b_psi The maximum threshold of PSI in any bins. <math>0 &lt; \text{b\_psi} &lt; 1</math>; 0 to 0.1 usually work. Default: 0.05.</li> <li>• b_or The maximum threshold of G/B index in any bins. <math>0 &lt; \text{b\_or} &lt; 1</math>; 0.05 to 0.3 usually work. Default: 0.15.</li> <li>• odds_psi The maximum threshold of Training and Testing G/B index PSI in any bins. <math>0 &lt; \text{odds\_psi} &lt; 1</math>; 0.01 to 0.3 usually work. Default: 0.1.</li> <li>• mono Monotonicity of all bins, the larger, the more nonmonotonic the bins will be. <math>0 &lt; \text{mono} &lt; 0.5</math>; 0.2 to 0.4 usually work. Default: 0.2.</li> <li>• kc number of cross-validations. 1 to 5 usually work. Default: 1.</li> </ul>
sp_values	A list of special value.
...	Other parameters.

## Details

The following is the list of Reference Principles

- 1.The increasing or decreasing trend of variables is consistent with the actual business experience.(The percent of Non-monotonic intervals of which are not head or tail is less than 0.35)
- 2.Maximum 10 intervals for a single variable.
- 3.Each interval should cover more than 2
- 4.Each interval needs at least 30 or 1
- 5.Combining the values of blank, missing or other special value into the same interval called missing.
- 6.The difference of Chi effect size between intervals should be at least 0.02 or more.
- 7.The difference of absolute odds ratio between intervals should be at least 0.1 or more.

- 8.The difference of positive rate between intervals should be at least 1/10 of the total positive rate.
- 9.The difference of G/B index between intervals should be at least 15 or more.
- 10.The PSI of each interval should be less than 0.1.

### Value

A list of breaks for x.

### See Also

[get\\_tree\\_breaks](#), [cut\\_equal](#), [get\\_breaks](#)

### Examples

```
#equal sample size breaks
equ_breaks = cut_equal(dat = UCICreditCard[, "PAY_AMT2"], g = 10)

# select best bins
bins_control = list(bins_num = 10, bins_pct = 0.02, b_chi = 0.02,
b_odds = 0.1, b_psi = 0.05, b_or = 0.15, mono = 0.3, odds_psi = 0.1, kc = 1)
select_best_breaks(dat = UCICreditCard, x = "PAY_AMT2", breaks = equ_breaks,
target = "default.payment.next.month", occur_time = "apply_date",
sp_values = NULL, bins_control = bins_control)
```

---

sim\_str

*sim\_str*

---

### Description

This function is not intended to be used by end user.

### Usage

```
sim_str(a, b, sep = "_|[.]|[A-Z]")
```

### Arguments

a	A string
b	A string
sep	Seprater of strings. Default is "_ [.] [A-Z]".

---

split_bins	<i>split_bins</i>
------------	-------------------

---

**Description**

split\_bins is for binning using breaks.

**Usage**

```
split_bins(dat, x, breaks = NULL, bins_no = TRUE)
```

**Arguments**

dat	A data.frame with independent variables.
x	The name of an independent variable.
breaks	Breaks for binning.
bins_no	Number the generated bins. Default is TRUE.

**Value**

A data.frame with Bined x.

**Examples**

```
bins = split_bins(dat = UCICreditCard,  
x = "PAY_AMT1", breaks = NULL, bins_no = TRUE)
```

---

split_bins_all	<i>Split bins all</i>
----------------	-----------------------

---

**Description**

split\_bins is for transforming data to bins. The split\_bins\_all function is a simpler wrapper for split\_bins.

**Usage**

```
split_bins_all(dat, x_list = NULL, ex_cols = NULL,  
breaks_list = NULL, bins_no = TRUE, note = FALSE,  
save_data = FALSE, file_name = NULL, dir_path = tempdir(), ...)
```

**Arguments**

dat	A data.frame with independent variables.
x_list	A list of x variables.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
breaks_list	A list contains breaks of variables. it is generated by <a href="#">codeget_breaks_all</a> , <a href="#">codeget_breaks</a>
bins_no	Number the generated bins. Default is TRUE.
note	Logical, outputs info. Default is TRUE.
save_data	Logical, save results in locally specified folder. Default is TRUE
file_name	The name for periodically saved woe file. Default is "dat_woe".
dir_path	The path for periodically saved woe file Default is "./data"
...	Additional parameters.

**Value**

A data.frame with splitted bins.

**See Also**

[get\\_tree\\_breaks](#), [cut\\_equal](#), [select\\_best\\_class](#), [select\\_best\\_breaks](#)

**Examples**

```
sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID", occur_time = "apply_date",
miss_values = list("", -1))

train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
                             occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
                             x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
save_data = FALSE, note = FALSE)
#woe transform
train_bins = split_bins_all(dat = dat_train,
                             breaks_list = breaks_list,
                             woe_name = FALSE)
test_bins = split_bins_all(dat = dat_test,
                             breaks_list = breaks_list,
                             note = FALSE)
```

---

`start_parallel_computing`*Parallel computing and export variables to global Env.*

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
start_parallel_computing(parallel = TRUE)
```

**Arguments**

`parallel`      A logical, default is TRUE.

**Value**

parallel works.

---

`stop_parallel_computing`*Stop parallel computing*

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
stop_parallel_computing(cluster)
```

**Arguments**

`cluster`      Parallel works.

**Value**

stop clusters.

---

str_match	<i>string match #' str_match search for matches to argument pattern within each element of a character vector:</i>
-----------	--

---

### Description

string match #' str\_match search for matches to argument pattern within each element of a character vector:

### Usage

```
str_match(pattern, str_r)
```

### 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 regexpr and gregexpr.
str_r	a character vector where matches are sought, or an object which can be coerced by as.character to a character vector. Long vectors are supported.

### Examples

```
original_nam = c("12mdd","11mdd","10mdd")
str_match(str_r = original_nam,pattern= "\\d+")
```

---

swap_analysis	<i>Swap Out/Swap In Analysis</i>
---------------	----------------------------------

---

### Description

swap\_analysis is for swap out/swap in analysis.

### Usage

```
swap_analysis(dat, new_rules, old_rules, target = NULL,
  cross_type = "total_pct", value = NULL)
```



**Arguments**

dat	A data.frame with independent variables.
new_rules	A list of new rules.
old_rules	A list of old rules.
target	The name of target variable.
cross_type	Output form of the result of crosstable. Provide these four forms: "total_sum", "total_pct", "bad_sum", "bad_pct".
value	The name of the variable to sum. When this parameter is NULL, the default statistics is to sum frequency.

**Value**

A cross table.

**Examples**

```
swap_analysis(dat = UCICreditCard, new_rules = list("SEX == 'male' & AGE < 25"),
             old_rules = list("SEX == 'male' & AGE < 30"),
             target = "default.payment.next.month", cross_type = "bad_pct", value = "LIMIT_BAL")
```

---

term_tfidf	<i>TF-IDF</i>
------------	---------------

---

**Description**

The `term_filter` is for filtering `stop_words` and low frequency words. The `term_idf` is for computing `idf`(inverse documents frequency) of terms. The `term_tfidf` is for computing `tf-idf` of documents.

**Usage**

```
term_tfidf(term_df, idf = NULL)

term_idf(term_df, n_total = NULL)

term_filter(term_df, low_freq = 0.01, stop_words = NULL)
```

**Arguments**

term_df	A data.frame with id and term.
idf	A data.frame with idf.
n_total	Number of documents.
low_freq	Use rate of terms or use numbers of terms.
stop_words	Stop words.

**Value**

A data.frame

**Examples**

```
term_df = data.frame(id = c(1,1,1,2,2,3,3,3,4,4,4,4,4,5,5,6,7,7,
                           8,8,8,9,9,9,10,10,11,11,11,11,11,11),
                    terms = c('a','b','c','a','c','d','d','a','b','c','a','c','d','a','c',
                               'd','a','e','f','b','c','f','b','c','h','h','i','c','d','g','k','k'))
term_df = term_filter(term_df = term_df, low_freq = 1)
idf = term_idf(term_df)
tf_idf = term_tfidf(term_df, idf = idf)
```

---

time\_series\_proc

*Process time series data*


---

**Description**

This function is used for time series data processing.

**Usage**

```
time_series_proc(dat, ID = NULL, group = NULL, time = NULL)
```

**Arguments**

dat	A data.frame contained only predict variables.
ID	The name of ID of observations or key variable of data. Default is NULL.
group	The group of behavioral or status variables.
time	The name of variable which is time when behavior was happened.

**Details**

The key to creating a good model is not the power of a specific modelling technique, but the breadth and depth of derived variables that represent a higher level of knowledge about the phenomena under examination.

**Examples**

```
dat = data.frame(id = c(1,1,1,2,2,3,3,3,4,4,4,4,4,5,5,6,7,7,
                       8,8,8,9,9,9,10,10,11,11,11,11,11,11),
                terms = c('a','b','c','a','c','d','d','a',
                           'b','c','a','c','d','a','c',
                           'd','a','e','f','b','c','f','b',
                           'c','h','h','i','c','d','g','k','k'),
                time = c(8,3,1,9,6,1,4,9,1,3,4,8,2,7,1,
                          3,4,1,8,7,2,5,7,8,8,2,1,5,7,2,7,3))

time_series_proc(dat = dat, ID = 'id', group = 'terms', time = 'time')
```

---

time_transfer	<i>Time Format Transferring</i>
---------------	---------------------------------

---

**Description**

time\_transfer is for transferring time variables to time format.

**Usage**

```
time_transfer(dat, date_cols = NULL, ex_cols = NULL, note = FALSE)
```

**Arguments**

dat	A data frame
date_cols	Names of time variable or regular expressions for finding time variables. Default is "DATE\$time\$date\$timestamp\$stamp\$".
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
note	Logical, outputs info. Default is TRUE.

**Value**

A data.frame with transferred time variables.

**Examples**

```
#transfer a variable.
dat = time_transfer(dat = lendingclub,date_cols = "issue_d")
class(dat["issue_d"])
#transfer a group of variables with similar name.
#transfer all time variables.
dat = time_transfer(dat = lendingclub[1:3],date_cols = "_d$")
class(dat["issue_d"])
```

---

time_variable	<i>time_variable</i>
---------------	----------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
time_variable(dat, date_cols = NULL, enddate = NULL)
```

**Arguments**

dat	A data.frame.
date_cols	Time variables.
enddate	End time.

---

time_vars_process	<i>Processing of Time or Date Variables</i>
-------------------	---

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
time_vars_process(df_tm = df_tm, x, enddate = NULL)
```

**Arguments**

df_tm	A data.frame
x	Time variable.
enddate	End time.

---

tnr_value	<i>tnr_value</i>
-----------	------------------

---

**Description**

tnr\_value is for get true negative rate for a prob or score.

**Usage**

```
tnr_value(prob, target)
```

**Arguments**

prob	A list of predict probability or score.
target	Vector of target.

**Value**

True Positive Rate

---

training_model	<i>Training model</i>
----------------	-----------------------

---

## Description

training\_model Model builder

## Usage

```
training_model(model_name = "mymodel", dat, dat_test = NULL,
  target = NULL, occur_time = NULL, obs_id = NULL, x_list = NULL,
  ex_cols = NULL, pos_flag = NULL, prop = 0.7, split_type = if
  (!is.null(occur_time)) "OOT" else "Random", preproc = TRUE,
  low_var = 0.99, missing_rate = 0.98, merge_cat = 30,
  remove_dup = TRUE, outlier_proc = TRUE, missing_proc = "median",
  default_miss = list(-1, "missing"), miss_values = NULL,
  one_hot = FALSE, trans_log = FALSE, feature_filter = list(filter =
  c("IV", "PSI", "COR", "XGB"), iv_cp = 0.02, psi_cp = 0.1, xgb_cp = 0,
  cv_folds = 1, hopper = FALSE), algorithm = list("LR", "XGB", "GBM",
  "RF"), LR.params = lr_params(), XGB.params = xgb_params(),
  GBM.params = gbm_params(), RF.params = rf_params(),
  breaks_list = NULL, parallel = FALSE, cores_num = NULL,
  save_pmml = FALSE, plot_show = FALSE, vars_plot = TRUE,
  model_path = tempdir(), seed = 46, ...)
```

## Arguments

model_name	A string, name of the project. Default is "mymodel"
dat	A data.frame with independent variables and target variable.
dat_test	A data.frame of test data. Default is NULL.
target	The name of target variable.
occur_time	The name of the variable that represents the time at which each observation takes place. Default is NULL.
obs_id	The name of ID of observations or key variable of data. Default is NULL.
x_list	Names of independent variables. Default is NULL.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
pos_flag	The value of positive class of target variable, default: "1".
prop	Percentage of train-data after the partition. Default: 0.7.
split_type	Methods for partition. See details at : <a href="#">train_test_split</a> .
preproc	Logical. Preprocess data. Default is TRUE.
low_var	Logical, delete low variance variables or not. Default is TRUE.

missing_rate	The maximum percent of missing values for recoding values to missing and non_missing.
merge_cat	merge categories of character variables that is more than m.
remove_dup	Logical, if TRUE, remove the duplicated observations.
outlier_proc	Logical, process outliers or not. Default is TRUE.
missing_proc	If logical, process missing values or not. If "median", then NAs imputation with k neighbors median. If "avg_dist", the distance weighted average method is applied to determine the NAs imputation with k neighbors. If "default", assigning the missing values to -1 or "missing", otherwise ,processing the missing values according to the results of missing analysis.
default_miss	Default value of missing data imputation, Default is list(-1,'missing').
miss_values	Other extreme value might be used to represent missing values, e.g: -9999, -9998. These miss_values will be encoded to -1 or "missing".
one_hot	Logical. If TRUE, one-hot_encoding of category variables. Default is FASLE.
trans_log	Logical, Logarithmic transformation. Default is FALSE.
feature_filter	Parameters for selecting important and stable features. See details at: <a href="#">feature_selector</a>
algorithm	Algorithms for training a model. list("LR", "XGB", "GBDT", "RF") are available.
LR.params	Parameters of logistic regression & scorecard. See details at : <a href="#">lr_params</a> .
XGB.params	Parameters of xgboost. See details at : <a href="#">xgb_params</a> .
GBM.params	Parameters of GBM. See details at : <a href="#">gbm_params</a> .
RF.params	Parameters of Random Forest. See details at : <a href="#">rf_params</a> .
breaks_list	A table containing a list of splitting points for each independent variable. Default is NULL.
parallel	Default is FALSE.
cores_num	The number of CPU cores to use.
save_pmml	Logical, save model in PMML format. Default is TRUE.
plot_show	Logical, show model performance in current graphic device. Default is FALSE.
vars_plot	Logical, if TRUE, plot distribution ,correlation or partial dependence of model input variables . Default is TRUE.
model_path	The path for periodically saved data file. Default is tempdir().
seed	Random number seed. Default is 46.
...	Other parameters.

**Value**

A list containing Model Objects.

**See Also**

[train\\_test\\_split](#), [data\\_cleansing](#), [feature\\_selector](#), [lr\\_params](#), [xgb\\_params](#), [gbm\\_params](#), [rf\\_params](#), [fast\\_high\\_cor\\_filter](#), [get\\_breaks\\_all](#), [lasso\\_filter](#), [woe\\_trans\\_all](#), [get\\_logistic\\_coef](#), [score\\_transfer](#), [get\\_score\\_card](#), [model\\_key\\_index](#), [ks\\_psi\\_plot](#), [get\\_plots](#), [ks\\_table\\_plot](#)

**Examples**

```

sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
x_list = c("LIMIT_BAL")
B_model = training_model(dat = dat,
                          model_name = "UCICreditCard",
                          target = "default.payment.next.month",
                          occur_time = NULL,
                          obs_id = NULL,
                          dat_test = NULL,
                          preproc = FALSE,
                          outlier_proc = FALSE,
                          missing_proc = FALSE,
                          feature_filter = NULL,
                          algorithm = list("LR"),
                          LR.params = lr_params(lasso = FALSE,
                                                  step_wise = FALSE,
                                                  score_card = FALSE),
                          breaks_list = NULL,
                          parallel = FALSE,
                          cores_num = NULL,
                          save_pmm1 = FALSE,
                          plot_show = FALSE,
                          vars_plot = FALSE,
                          model_path = tempdir(),
                          seed = 46)

```

---

**train\_lr***Trainig LR model*

---

**Description**

train\_lr is for training the logistic regression model using in [training\\_model](#).

**Usage**

```

train_lr(dat_train, dat_test = NULL, target, x_list = NULL,
         occur_time = NULL, prop = 0.7, tree_control = list(p = 0.02, cp =
         0.00000001, xval = 5, maxdepth = 10), bins_control = list(bins_num =
         10, bins_pct = 0.05, b_chi = 0.02, b_odds = 0.1, b_psi = 0.03, b_or =
         0.15, mono = 0.2, odds_psi = 0.15, kc = 1), thresholds = list(cor_p =
         0.8, iv_i = 0.02, psi_i = 0.1, cos_i = 0.6), lasso = TRUE,
         step_wise = TRUE, best_lambda = "lambda.auc", seed = 1234, ...)

```

**Arguments**

dat_train	data.frame of train data. Default is NULL.
dat_test	data.frame of test data. Default is NULL.
target	name of target variable.
x_list	names of independent variables. Default is NULL.
occur_time	The name of the variable that represents the time at which each observation takes place. Default is NULL.
prop	Percentage of train-data after the partition. Default: 0.7.
tree_control	the list of parameters to control cutting initial breaks by decision tree. See details at: <a href="#">get_tree_breaks</a>
bins_control	the list of parameters to control merging initial breaks. See details at: <a href="#">select_best_breaks</a> , <a href="#">select_best</a>
thresholds	Thresholds for selecting variables. <ul style="list-style-type: none"> <li>• cor_p The maximum threshold of correlation. Default: 0.8.</li> <li>• iv_i The minimum threshold of IV. 0.01 to 0.1 usually work. Default: 0.02</li> <li>• psi_i The maximum threshold of PSI. 0.1 to 0.3 usually work. Default: 0.1.</li> <li>• cos_i cos_similarity of posive rate of train and test. 0.7 to 0.9 usually work. Default: 0.5.</li> </ul>
lasso	Logical, if TRUE, variables filtering by LASSO. Default is TRUE.
step_wise	Logical, stepwise method. Default is TRUE.
best_lambda	Methods of best lambda standards using to filter variables by LASSO. There are 3 methods: ("lambda.auc", "lambda.ks", "lambda.sim_sign") . Default is "lambda.auc".
seed	Random number seed. Default is 1234.
...	Other parameters

---

train_test_split	<i>Train-Test-Split</i>
------------------	-------------------------

---

**Description**

train\_test\_split Functions for partition of data.

**Usage**

```
train_test_split(dat, prop = 0.7, split_type = "Random",
  occur_time = NULL, cut_date = NULL, start_date = NULL,
  save_data = FALSE, dir_path = tempdir(), file_name = NULL,
  note = FALSE, seed = 43)
```



**Arguments**

dat	A data.frame with independent variables and target variable.
prop	The percentage of train data samples after the partition.
split_type	Methods for partition. <ul style="list-style-type: none"> <li>• "Random" is to split train &amp; test set randomly.</li> <li>• "OOT" is to split by time for observation over time test.</li> <li>• "byRow" is to split by rownumbers.</li> </ul>
occur_time	The name of the variable that represents the time at which each observation takes place. It is used for "OOT" split.
cut_date	Time points for splitting data sets, e.g. : splitting Actual and Expected data sets.
start_date	The earliest occurrence time of observations.
save_data	Logical, save results in locally specified folder. Default is FALSE.
dir_path	The path for periodically saved data file. Default is "./data".
file_name	The name for periodically saved data file. Default is "dat".
note	Logical. Outputs info. Default is TRUE.
seed	Random number seed. Default is 46.

**Value**

A list of indices (train-test)

**Examples**

```
train_test <- train_test_split(lendingclub,
  split_type = "OOT", prop = 0.7,
  occur_time = "issue_d", seed = 12, save_data = FALSE)
dat_train = train_test$train
dat_test = train_test$test
```

---

train\_xgb

*Training XGboost*


---

**Description**

train\_xgb is for training a xgb model using in [training\\_model](#).

**Usage**

```
train_xgb(seed_number = 1234, dtrain, nthread = 2, nfold = 1,
  watchlist = NULL, nrounds = 100, f_eval = "ks",
  early_stopping_rounds = 10, verbose = 0, params = NULL, ...)
```

**Arguments**

seed_number	Random number seed. Default is 1234.
dtrain	train-data of xgb.DMatrix datasets.
nthread	Number of threads
nfold	Number of the cross validation of xgboost
watchlist	named list of xgb.DMatrix datasets to use for evaluating model performance.generating by <a href="#">xgb_data</a>
nrounds	Max number of boosting iterations.
f_eval	Custimized evaluation function,"ks" & "auc" are available.
early_stopping_rounds	If NULL, the early stopping function is not triggered. If set to an integer k, training with a validation set will stop if the performance doesn't improve for k rounds.
verbose	If 0, xgboost will stay silent. If 1, it will print information about performance.
params	List of contains parameters of xgboost. The complete list of parameters is available at: <a href="http://xgboost.readthedocs.io/en/latest/parameter.html">http://xgboost.readthedocs.io/en/latest/parameter.html</a>
...	Other parameters

UCICreditCard

*UCI Credit Card data***Description**

This research aimed at the case of customers's default payments in Taiwan and compares the predictive accuracy of probability of default among six data mining methods. This research employed a binary variable, default payment (Yes = 1, No = 0), as the response variable. This study reviewed the literature and used the following 24 variables as explanatory variables

**Format**

A data frame with 30000 rows and 26 variables.

**Details**

- ID: Customer id
- apply\_date: This is a fake occur time.
- LIMIT\_BAL: Amount of the given credit (NT dollar): it includes both the individual consumer credit and his/her family (supplementary) credit.
- SEX: Gender (male; female).
- EDUCATION: Education (1 = graduate school; 2 = university; 3 = high school; 4 = others).
- MARRIAGE: Marital status (1 = married; 2 = single; 3 = others).

- AGE: Age (year) History of past payment. We tracked the past monthly payment records (from April to September, 2005) as follows:
- PAY\_0: the repayment status in September
- PAY\_2: the repayment status in August
- PAY\_3: ...
- PAY\_4: ...
- PAY\_5: ...
- PAY\_6: the repayment status in April The measurement scale for the repayment status is: -1 = pay duly; 1 = payment delay for one month; 2 = payment delay for two months;...;8 = payment delay for eight months; 9 = payment delay for nine months and above. Amount of bill statement (NT dollar)
- BILL\_AMT1: amount of bill statement in September
- BILL\_AMT2: mount of bill statement in August
- BILL\_AMT3: ...
- BILL\_AMT4: ...
- BILL\_AMT5: ...
- BILL\_AMT6: amount of bill statement in April Amount of previous payment (NT dollar)
- PAY\_AMT1: amount paid in September
- PAY\_AMT2: amount paid in August
- PAY\_AMT3: ....
- PAY\_AMT4: ...
- PAY\_AMT5: ...
- PAY\_AMT6: amount paid in April
- default.payment.next.month: default payment (Yes = 1, No = 0), as the response variable

**Source**

<http://archive.ics.uci.edu/ml/datasets/default+of+credit+card+clients>

**See Also**

[lendingclub](#)

---

variable_process	<i>variable_process</i>
------------------	-------------------------

---

**Description**

This function is not intended to be used by end user.

**Usage**

```
variable_process(add)
```

**Arguments**

add	A data.frame
-----	--------------

---

var_group_proc	<i>Process group numeric variables</i>
----------------	--

---

**Description**

This function is used for grouped numeric data processing.

**Usage**

```
var_group_proc(dat, ID = NULL, group = NULL, num_var = NULL)
```

**Arguments**

dat	A data.frame contained only predict variables.
ID	The name of ID of observations or key variable of data. Default is NULL.
group	The group of behavioral or status variables.
num_var	The name of numeric variable to process.

**Examples**

```
dat = data.frame(id = c(1,1,1,2,2,3,3,3,4,4,4,4,4,5,5,6,7,7,
                      8,8,8,9,9,9,10,10,11,11,11,11,11,11),
                terms = c('a','b','c','a','c','d','d','a',
                          'b','c','a','c','d','a','c',
                          'd','a','e','f','b','c','f','b',
                          'c','h','h','i','c','d','g','k','k'),
                time = c(8,3,1,9,6,1,4,9,1,3,4,8,2,7,1,
                        3,4,1,8,7,2,5,7,8,8,2,1,5,7,2,7,3))

time_series_proc(dat = dat, ID = 'id', group = 'terms', time = 'time')
```

---

woe_trans_all	<i>WOE Transformation</i>
---------------	---------------------------

---

### Description

woe\_trans is for transforming data to woe. The woe\_trans\_all function is a simpler wrapper for woe\_trans.

### Usage

```
woe_trans_all(dat, x_list = NULL, ex_cols = NULL, bins_table = NULL,
  target = NULL, breaks_list = NULL, note = FALSE,
  save_data = FALSE, parallel = FALSE, woe_name = FALSE,
  file_name = NULL, dir_path = tempdir(), ...)
```

```
woe_trans(dat, x, bins_table = NULL, target = NULL,
  breaks_list = NULL, woe_name = FALSE)
```

### Arguments

dat	A data.frame with independent variables.
x_list	A list of x variables.
ex_cols	Names of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
bins_table	A table contains woe of each bin of variables, it is generated by <a href="#">code</a> <code>get_bins_table_all</code> , <a href="#">code</a> <code>get_bins_table</code>
target	The name of target variable. Default is NULL.
breaks_list	A list contains breaks of variables. it is generated by <a href="#">code</a> <code>get_breaks_all</code> , <a href="#">code</a> <code>get_breaks</code>
note	Logical, outputs info. Default is TRUE.
save_data	Logical, save results in locally specified folder. Default is TRUE
parallel	Logical, parallel computing. Default is FALSE.
woe_name	Logical. Add "_woe" at the end of the variable name.
file_name	The name for periodically saved woe file. Default is "dat_woe".
dir_path	The path for periodically saved woe file Default is "./data"
...	Additional parameters.
x	The name of an independent variable.

### Value

A list of breaks for each variables.

### See Also

[get\\_tree\\_breaks](#), [cut\\_equal](#), [select\\_best\\_class](#), [select\\_best\\_breaks](#)

**Examples**

```

sub = cv_split(UCICreditCard, k = 30)[[1]]
dat = UCICreditCard[sub,]
dat = re_name(dat, "default.payment.next.month", "target")
dat = data_cleansing(dat, target = "target", obs_id = "ID", occur_time = "apply_date",
miss_values = list("", -1))

train_test <- train_test_split(dat, split_type = "OOT", prop = 0.7,
                             occur_time = "apply_date")

dat_train = train_test$train
dat_test = train_test$test
#get breaks of all predictive variables
x_list = c("PAY_0", "LIMIT_BAL", "PAY_AMT5", "EDUCATION", "PAY_3", "PAY_2")
breaks_list <- get_breaks_all(dat = dat_train, target = "target",
                             x_list = x_list, occur_time = "apply_date", ex_cols = "ID",
save_data = FALSE, note = FALSE)
#woe transform
train_woe = woe_trans_all(dat = dat_train,
                          target = "target",
                          breaks_list = breaks_list,
                          woe_name = FALSE)
test_woe = woe_trans_all(dat = dat_test,
                          target = "target",
                          breaks_list = breaks_list,
                          note = FALSE)

```

---

xgb\_data

*XGboost data*


---

**Description**

xgb\_data is for prepare data using in [training\\_model](#).

**Usage**

```
xgb_data(dat_train, target, dat_test = NULL, x_list = NULL,
prop = 0.7, occur_time = NULL)
```

**Arguments**

dat_train	data.frame of train data. Default is NULL.
target	name of target variable.
dat_test	data.frame of test data. Default is NULL.
x_list	names of independent variables of raw data. Default is NULL.
prop	Percentage of train-data after the partition. Default: 0.7.
occur_time	The name of the variable that represents the time at which each observation takes place. Default is NULL.

---

xgb_filter	<i>Select Features using XGB</i>
------------	----------------------------------

---

### Description

xgb\_filter is for selecting important features using xgboost.

### Usage

```
xgb_filter(dat_train, dat_test = NULL, target = NULL,
           pos_flag = NULL, x_list = NULL, occur_time = NULL,
           ex_cols = NULL, xgb_params = list(nrounds = 100, max_depth = 6, eta =
           0.1, min_child_weight = 1, subsample = 1, colsample_bytree = 1, gamma =
           0, scale_pos_weight = 1, early_stopping_rounds = 10, objective =
           "binary:logistic"), f_eval = "auc", cv_folds = 1, cp = NULL,
           seed = 46, vars_name = TRUE, note = TRUE, save_data = FALSE,
           file_name = NULL, dir_path = tempdir(), ...)
```

### Arguments

dat_train	A data.frame with independent variables and target variable.
dat_test	A data.frame of test data. Default is NULL.
target	The name of target variable.
pos_flag	The value of positive class of target variable, default: "1".
x_list	Names of independent variables.
occur_time	The name of the variable that represents the time at which each observation takes place.
ex_cols	A list of excluded variables. Regular expressions can also be used to match variable names. Default is NULL.
xgb_params	Parameters of xgboost. The complete list of parameters is available at: <a href="http://xgboost.readthedocs.io/en/latest/parameter.html">http://xgboost.readthedocs.io/en/latest/parameter.html</a> .
f_eval	Customized evaluation function, "ks" & "auc" are available.
cv_folds	Number of cross-validations. Default: 5.
cp	Threshold of XGB feature's Gain. Default is 1/number of independent variables.
seed	Random number seed. Default is 46.
vars_name	Logical, output a list of filtered variables or table with detailed IV and PSI value of each variable. Default is FALSE.
note	Logical, outputs info. Default is TRUE.
save_data	Logical, save results results in locally specified folder. Default is FALSE.
file_name	The name for periodically saved results files. Default is "Feature_importance_XGB".
dir_path	The path for periodically saved results files. Default is "./variable".
...	Other parameters to pass to xgb_params.

**Value**

Selected variables.

**See Also**

[psi\\_iv\\_filter](#), [gbm\\_filter](#), [feature\\_selector](#)

**Examples**

```
dat = UCICreditCard[1:1000,c(2,4,8:9,26)]
xgb_params = list(nrounds = 100, max_depth = 6, eta = 0.1,
                 min_child_weight = 1, subsample = 1,
                 colsample_bytree = 1, gamma = 0, scale_pos_weight = 1,
                 early_stopping_rounds = 10,
                 objective = "binary:logistic")

## Not run:
xgb_features <- xgb_filter(dat_train = dat, dat_test = NULL,
                          target = "default.payment.next.month", occur_time = "apply_date", f_eval = 'ks',
                          xgb_params = xgb_params,
                          cv_folds = 1, ex_cols = "ID$|date$|default.payment.next.month$", vars_name = FALSE)

## End(Not run)
```

---

xgb\_params

*XGboost Parameters*

---

**Description**

xgb\_params is the list of parameters to train a XGB model using in [training\\_model](#). xgb\_params\_search is for searching the optimal parameters of xgboost, if any parameters of params in [xgb\\_params](#) is more than one.

**Usage**

```
xgb_params(nrounds = 1000, params = list(max_depth = 6, eta = 0.01,
                                         gamma = 0, min_child_weight = 1, subsample = 1, colsample_bytree = 1,
                                         scale_pos_weight = 1), early_stopping_rounds = 100,
           method = "random_search", iters = 10, f_eval = "auc", nfold = 1,
           nthread = 2, ...)
```

```
xgb_params_search(dat_train, target, dat_test = NULL, x_list = NULL,
                  prop = 0.7, occur_time = NULL, method = "random_search",
                  iters = 10, nrounds = 100, early_stopping_rounds = 10,
                  params = list(max_depth = 6, eta = 0.01, gamma = 0, min_child_weight = 1,
                                subsample = 1, colsample_bytree = 1, scale_pos_weight = 1),
                  f_eval = "auc", nfold = 1, nthread = 2, ...)
```



**Arguments**

nrounds	Max number of boosting iterations.
params	List of contains parameters of xgboost. The complete list of parameters is available at: <a href="http://xgboost.readthedocs.io/en/latest/parameter.html">http://xgboost.readthedocs.io/en/latest/parameter.html</a>
early_stopping_rounds	If NULL, the early stopping function is not triggered. If set to an integer k, training with a validation set will stop if the performance doesn't improve for k rounds.
method	Method of searching optimal parameters. "random_search", "grid_search", "local_search" are available.
iters	Number of iterations of "random_search" optimal parameters.
f_eval	Custimized evaluation function, "ks" & "auc" are available.
nfold	Number of the cross validation of xgboost
nthread	Number of threads
...	Other parameters
dat_train	A data.frame of train data. Default is NULL.
target	Name of target variable.
dat_test	A data.frame of test data. Default is NULL.
x_list	Names of independent variables. Default is NULL.
prop	Percentage of train-data after the partition. Default: 0.7.
occur_time	The name of the variable that represents the time at which each observation takes place. Default is NULL.

**Value**

A list of parameters.

**See Also**

[training\\_model](#), [lr\\_params](#), [gbm\\_params](#), [rf\\_params](#)

%alike%

*Fuzzy String matching*

**Description**

Fuzzy String matching

**Usage**

x %alike% y

**Arguments**

x            A string.  
y            A string.

**Value**

Logical.

**Examples**

```
"xyz" %alike% "xy"
```

---

%islike%

*Fuzzy String matching*

---

**Description**

Fuzzy String matching

**Usage**

```
x %islike% y
```

**Arguments**

x            A string.  
y            A string.

**Value**

Logical.

**Examples**

```
"xyz" %islike% "yz$"
```

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