

# Package: misc (via r-universe)

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**Title** Miscellaneous Useful R Functions

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**Description** Miscellaneous Useful R Functions.

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parallel, compiler

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**Repository** <https://techtonique.r-universe.dev>

**RemoteUrl** <https://github.com/thierrymoudiki/misc>

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debug_print	<i>Debug print</i>
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**Description**

Debug print

**Usage**

```
debug_print(x)
```

**Arguments**

x	An object to be printed
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**Examples**

```
misc::debug_print(1:10)
misc::debug_print("Hello, world!")
```

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fit_param_dist	<i>Fit multiple parametric distributions, compute KL divergence, simulate best fit</i>
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**Description**

Fit multiple parametric distributions, compute KL divergence, simulate best fit

**Usage**

```
fit_param_dist(vector, num_bins = 30, verbose = TRUE)
```

**Arguments**

vector	Numeric vector of data to fit
num_bins	Number of bins for the empirical histogram
verbose	Logical indicating whether to print results

**Value**

Function to simulate data from the best-fitting distribution

**Examples**

```
set.seed(123)
n <- 1000
vector <- rnorm(n)

start <- proc.time()[3]
simulate_function <- fit_param_dist(vector)
end <- proc.time()[3]
print(paste("Time taken:", end - start))
simulated_data <- simulate_function(n) # Generate 100 samples from the best-fit distribution
par(mfrow = c(1, 2))
hist(vector, main = "Original Data", xlab = "Value", ylab = "Frequency")
hist(simulated_data, main = "Simulated Data", xlab = "Value", ylab = "Frequency")
```

---

is\_package\_available *Check if a package is available*

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**Description**

Check if a package is available

**Usage**

```
is_package_available(pkg_name)
```

**Arguments**

pkg\_name      A package name

**Value**

A logical value

**Examples**

```
misc::is_package_available("dplyr")
```

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is_wholenumber	<i>Check if a number is a whole number</i>
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**Description**

Check if a number is a whole number

**Usage**

```
is_wholenumber(x, tol = .Machine$double.eps^0.5)
```

**Arguments**

x	A number
tol	A tolerance level

**Value**

A logical value

**Examples**

```
is_wholenumber(1)
is_wholenumber(1.1)
is_wholenumber(1L)
```

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KL_divergence_hist	<i>Function to calculate KL divergence for continuous distributions using histograms</i>
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**Description**

Function to calculate KL divergence for continuous distributions using histograms

**Usage**

```
KL_divergence_hist(P, Q)
```

**Arguments**

P	Numeric vector representing the empirical distribution
Q	Numeric vector representing the theoretical distribution

**Value**

KL divergence between P and Q

**Examples**

```
P <- c(0.2, 0.3, 0.5)
Q <- c(0.1, 0.4, 0.5)
misc::KL_divergence_hist(P, Q)
```

---

one_hot_encode	<i>One-hot encoding</i>
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**Description**

One-hot encoding

**Usage**

```
one_hot_encode(y)
```

**Arguments**

y	A vector of class labels
n_classes	The number of classes

**Value**

A matrix of one-hot encoded labels

**Examples**

```
y <- as.factor(c(1, 2, 1, 1, 2))
misc::one_hot_encode(y)
```

---

parfor	<i>Sequential or parallel for loop.</i>
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**Description**

Sequential or parallel for loop.

## Usage

```
parfor(  
  what,  
  args,  
  cl = NULL,  
  combine = c,  
  errorhandling = c("stop", "remove", "pass"),  
  verbose = FALSE,  
  show_progress = TRUE,  
  export = NULL,  
  ...  
)
```

## Arguments

what	A function.
args	A list of arguments.
cl	Number of cores to use. If NULL, the loop will be sequential. If -1, the number of cores will be detected automatically.
combine	A function to combine the results.
errorhandling	A character string specifying how to handle errors. Possible values are "stop", "remove", and "pass".
verbose	A logical indicating whether to print progress.
show_progress	A logical indicating whether to show a progress bar.
export	A list of objects to export to the workers.
...	Additional arguments to pass to what for <code>foreach::foreach</code> (excluding <code>.combine</code> , <code>.errorhandling</code> , <code>.options.snow</code> , <code>.verbose</code> , and <code>.export</code> ).

## Value

A list of results.

## Examples

```
# Sequential  
print(misc::parfor(function(x) x^2, 1:10))  
  
# Parallel  
print(misc::parfor(function(x) x^2, 1:10, cl = 2))
```

---

rm_zero_cols	<i>Removing columns containing only zeros</i>
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**Description**

Removing columns containing only zeros

**Usage**

```
rm_zero_cols(X)
```

**Arguments**

X                    A matrix or data frame

**Value**

A matrix or data frame

**Examples**

```
X <- matrix(c(1, 0, 3, 0, 5, 0, 0, 0), nrow = 2)
print(misc::rm_zero_cols(X))
```

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scale_matrix	<i>Scale matrix</i>
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**Description**

Scale matrix

**Usage**

```
scale_matrix(X, X_mean = NULL, X_sd = NULL)
```

**Arguments**

X                    A matrix  
X\_mean              Mean of each column  
X\_sd                 Standard deviation of each column

**Value**

A list containing the scaled matrix, mean of each column, and standard deviation of each column

**Examples**

```
X <- matrix(c(1, 2, 3, 4, 5, 6), nrow = 2)
(X_scaled <- misc::scale_matrix(X))
(X_scaled <- misc::scale_matrix(X, X_mean = colMeans(X), X_sd = apply(X, 2, stats::sd)))
print(colMeans(X_scaled$X))
print(apply(X_scaled$X, 2, stats::sd))
```

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sort\_df

*Sort data frame*

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**Description**

Sort data frame

**Usage**

```
sort_df(df, by, decreasing = FALSE)
```

**Arguments**

df	data frame
by	column to sort by
decreasing	logical. Should sorting be decreasing?

**Value**

A sorted data frame

**Examples**

```
df <- data.frame(a = c(2, 4, 3), b = c(3, 5, 1))
misc::sort_df(df, "a")
misc::sort_df(df, "b", decreasing = TRUE)
```



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splitts	<i>Partition a time series object</i>
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**Description**

Partition a time series object

**Usage**

```
splitts(y, split_prob = 0.5, return_indices = FALSE)
```

**Arguments**

`y` A time series object  
`split_prob` Splitting ratio  
`return_indices` if TRUE, returns series' indices, otherwise, time series objects

**Examples**

```
misc::splitts(ts(1:10))
```

---

split_data	<i>Split a dataset</i>
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**Description**

Split a dataset

**Usage**

```
split_data(y, p = 0.5, seed = 123, type_split = c("stratify", "sequential"))
```

**Arguments**

`y` A vector of labels  
`p` A proportion of the dataset to split  
`seed` An integer to set the seed  
`type_split` A character string specifying the type of split

**Value**

A vector of indices

**Examples**

```
set.seed(123)
(y <- rnorm(10))
misc::split_data(y, 0.5)
misc::split_data(y, 0.5, type_split = "sequential")
```

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timeit	<i>Timing an expression</i>
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**Description**

Timing an expression

**Usage**

```
timeit(expr, times = 1, ...)
```

**Arguments**

expr	an R expression
times	number of repetitions
...	additional arguments passed to <a href="#">base::eval</a>

**Value**

the elapsed time in seconds

**Examples**

```
timeit(1 + 1)
timeit(1 + 1, times = 10)
```

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vlookup	<i>VLOOKUP</i>
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**Description**

A simple implementation similar to the VLOOKUP function in Excel.

**Usage**

```
vlookup(this, df, key, value)
```

**Arguments**

this	The value to look up
df	A data frame
key	The column to look up
value	The column to return

**Value**

The value in the value column corresponding to the key column

**Examples**

```
df <- data.frame(key = c("a", "b", "c"), value = c(1, 2, 3))
print(misc::vlookup("b", df, "key", "value"))
```

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winkler\_score

*Winkler score for probabilistic forecasts*


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**Description**

Winkler score for probabilistic forecasts

**Usage**

```
winkler_score(actual, lower, upper, level = 95, scale = FALSE)
```

**Arguments**

actual	numeric vector of actual values
lower	numeric vector of lower bounds
upper	numeric vector of upper bounds
level	numeric level of confidence
scale	logical, if TRUE, the score is scaled by the range of the bounds

**Value**

numeric score

**Examples**

```
actual <- c(1, 2, 3, 4, 5)
lower <- c(0, 1, 2, 3, 4)
upper <- c(2, 3, 4, 5, 6)
winkler_score(actual, lower, upper)
winkler_score(actual, lower, upper, scale = TRUE)
winkler_score(actual, lower, upper, level = 99)
winkler_score(actual, lower, upper, level = 99, scale = TRUE)
```

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