Package 'oeli'

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check_correlation_matrix *Check correlation matrix*

Description

These functions check whether the input fulfills the properties of a correlation matrix.

Usage

```
check_correlation_matrix(x, dim = NULL, tolerance = sqrt(.Machine$double.eps))

assert_correlation_matrix(
  x,
  dim = NULL,
  tolerance = sqrt(.Machine$double.eps),
  .var.name = checkmate::vname(x),
  add = NULL
)

test_correlation_matrix(x, dim = NULL, tolerance = sqrt(.Machine$double.eps))
```

Arguments

x	[any] Object to check.
dim	[integer(1)] The matrix dimension.
tolerance	[numeric(1)] A non-negative tolerance value.
.var.name	[character(1)] Name of the checked object to print in assertions. Defaults to the heuristic implemented in vname.
add	[AssertCollection] Collection to store assertion messages. See AssertCollection.

Value

Same as documented in check_matrix.

See Also

Other matrix helpers: check_covariance_matrix(), check_transition_probability_matrix(), cov_to_chol(), diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(), sample_correlation_matrix(), sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
M <- matrix(c(1,  0.9,  0.9, 0.9,  1,  -0.9, 0.9,  -0.9,  1), nrow = 3)
check_correlation_matrix(M)
test_correlation_matrix(M)
## Not run:
assert_correlation_matrix(M)

## End(Not run)
```

`check_covariance_matrix`
Check covariance matrix

Description

These functions check whether the input fulfills the properties of a covariance matrix.

Usage

```
check_covariance_matrix(x, dim = NULL, tolerance = sqrt(.Machine$double.eps))

assert_covariance_matrix(
  x,
  dim = NULL,
  tolerance = sqrt(.Machine$double.eps),
  .var.name = checkmate::vname(x),
  add = NULL
)

test_covariance_matrix(x, dim = NULL, tolerance = sqrt(.Machine$double.eps))
```

Arguments

`x`	[any] Object to check.
`dim`	[`integer(1)`] The matrix dimension.
`tolerance`	[`numeric(1)`] A non-negative tolerance value.
`.var.name`	[`character(1)`] Name of the checked object to print in assertions. Defaults to the heuristic implemented in `vname`.
`add`	[`AssertCollection`] Collection to store assertion messages. See `AssertCollection`.

Value

Same as documented in check_matrix.

See Also

Other matrix helpers: check_correlation_matrix(), check_transition_probability_matrix(), cov_to_chol(), diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(), sample_correlation_matrix(), sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
M <- matrix(c(1, 2, 3, 2, 1, 2, 3, 2, 1), nrow = 3)
check_covariance_matrix(M)
test_covariance_matrix(M)
## Not run:
assert_covariance_matrix(M)

## End(Not run)
```

---

check_list_of_lists *Check list of lists*

Description

These functions check whether the input is a list that contains list elements.

Usage

```
check_list_of_lists(x, len = NULL)

assert_list_of_lists(
  x,
  len = NULL,
  .var.name = checkmate::vname(x),
  add = NULL
)

test_list_of_lists(x, len = NULL)
```

Arguments

x	[any] Object to check.
len	[integer(1)] Exact expected length of x.

.var.name	[character(1)] Name of the checked object to print in assertions. Defaults to the heuristic implemented in vname.
add	[AssertCollection] Collection to store assertion messages. See AssertCollection.

Value

Same as documented in check_list.

See Also

Other list helpers: merge_lists()

Examples

```
L <- list(list(1), list(2), 3)
check_list_of_lists(L)
test_list_of_lists(L)
## Not run:
assert_list_of_lists(L)

## End(Not run)
```

---

check_numeric_vector *Check numeric vector*

Description

These functions check whether the input is a numeric vector.

Usage

```
check_numeric_vector(
  x,
  lower = -Inf,
  upper = Inf,
  finite = FALSE,
  any.missing = TRUE,
  all.missing = TRUE,
  len = NULL,
  min.len = NULL,
  max.len = NULL,
  unique = FALSE,
  sorted = FALSE,
  names = NULL,
  typed.missing = FALSE,
  null.ok = FALSE
```

```
)

assert_numeric_vector(
  x,
  lower = -Inf,
  upper = Inf,
  finite = FALSE,
  any.missing = TRUE,
  all.missing = TRUE,
  len = NULL,
  min.len = NULL,
  max.len = NULL,
  unique = FALSE,
  sorted = FALSE,
  names = NULL,
  typed.missing = FALSE,
  null.ok = FALSE,
  .var.name = checkmate::vname(x),
  add = NULL
)

test_numeric_vector(
  x,
  lower = -Inf,
  upper = Inf,
  finite = FALSE,
  any.missing = TRUE,
  all.missing = TRUE,
  len = NULL,
  min.len = NULL,
  max.len = NULL,
  unique = FALSE,
  sorted = FALSE,
  names = NULL,
  typed.missing = FALSE,
  null.ok = FALSE
)
```

Arguments

x	[any] Object to check.
lower	[numeric(1)] Lower value all elements of x must be greater than or equal to.
upper	[numeric(1)] Upper value all elements of x must be lower than or equal to.
finite	[logical(1)] Check for only finite values? Default is FALSE.

any.missing [logical(1)]
Are vectors with missing values allowed? Default is TRUE.

all.missing [logical(1)]
Are vectors with no non-missing values allowed? Default is TRUE. Note that empty vectors do not have non-missing values.

len [integer(1)]
Exact expected length of x.

min.len [integer(1)]
Minimal length of x.

max.len [integer(1)]
Maximal length of x.

unique [logical(1)]
Must all values be unique? Default is FALSE.

sorted [logical(1)]
Elements must be sorted in ascending order. Missing values are ignored.

names [character(1)]
Check for names. See checkNamed for possible values. Default is "any" which performs no check at all. Note that you can use checkSubset to check for a specific set of names.

typed.missing [logical(1)]
If set to FALSE (default), all types of missing values (NA, NA_integer_, NA_real_, NA_character_ or NA_character_) as well as empty vectors are allowed while type-checking atomic input. Set to TRUE to enable strict type checking.

null.ok [logical(1)]
If set to TRUE, x may also be NULL. In this case only a type check of x is performed, all additional checks are disabled.

.var.name [character(1)]
Name of the checked object to print in assertions. Defaults to the heuristic implemented in vname.

add [AssertCollection]
Collection to store assertion messages. See AssertCollection.

Value

Same as documented in check_numeric.

See Also

Other vector helpers: check_probability_vector(), chunk_vector(), insert_vector_entry(), match_numerics(), permutations(), subsets(), vector_occurrence()

Examples

```
x <- c(1, 2, "3")
check_numeric_vector(x)
test_numeric_vector(x)
```

```
## Not run:
assert_numeric_vector(x)

## End(Not run)
```

`check_probability_vector` *Check probability vector*

Description

These functions check whether the input fulfills the properties of a probability matrix.

Usage

```
check_probability_vector(x, len = NULL, tolerance = sqrt(.Machine$double.eps))

assert_probability_vector(
  x,
  len = NULL,
  tolerance = sqrt(.Machine$double.eps),
  .var.name = checkmate::vname(x),
  add = NULL
)

test_probability_vector(x, len = NULL, tolerance = sqrt(.Machine$double.eps))
```

Arguments

`x`	[any] Object to check.
`len`	[`integer(1)`] Exact expected length of `x`.
`tolerance`	[`numeric(1)`] A non-negative tolerance value.
`.var.name`	[`character(1)`] Name of the checked object to print in assertions. Defaults to the heuristic implemented in `vname`.
`add`	[`AssertCollection`] Collection to store assertion messages. See `AssertCollection`.

Value

Same as documented in `check_numeric`.

See Also

Other vector helpers: check_numeric_vector(), chunk_vector(), insert_vector_entry(), match_numerics(), permutations(), subsets(), vector_occurrence()

Examples

```
p <- c(0.2, 0.3, 0.6)
check_probability_vector(p)
test_probability_vector(p)
## Not run:
assert_probability_vector(p)

## End(Not run)
```

---

check_transition_probability_matrix

Check transition probability matrix

Description

These functions check whether the input is a transition probability matrix.

Usage

```
check_transition_probability_matrix(
  x,
  dim = NULL,
  tolerance = sqrt(.Machine$double.eps)
)

assert_transition_probability_matrix(
  x,
  dim = NULL,
  tolerance = sqrt(.Machine$double.eps),
  .var.name = checkmate::vname(x),
  add = NULL
)

test_transition_probability_matrix(
  x,
  dim = NULL,
  tolerance = sqrt(.Machine$double.eps)
)
```

Arguments

x	[any] Object to check.
dim	[integer(1)] The matrix dimension.
tolerance	[numeric(1)] A non-negative tolerance value.
.var.name	[character(1)] Name of the checked object to print in assertions. Defaults to the heuristic implemented in vname.
add	[AssertCollection] Collection to store assertion messages. See AssertCollection.

Value

Same as documented in check_matrix.

See Also

Other matrix helpers: check_correlation_matrix(), check_covariance_matrix(), cov_to_chol(), diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(), sample_correlation_matr sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
T <- matrix(c(0.8,  0.2,  0.1, 0.1,  0.7,  0.4, 0.1,  0.1,  0.6), nrow = 3)
check_transition_probability_matrix(T)
test_transition_probability_matrix(T)
## Not run:
assert_transition_probability_matrix(T)

## End(Not run)
```

---

chunk_vector *Split a vector into chunks*

Description

This function either

- splits a vector into n chunks of equal size (type = 1),
- splits a vector into chunks of size n (type = 2).

Usage

```
chunk_vector(x, n, type = 1, strict = FALSE)
```

Arguments

x	[atomic()‘] A vector of elements.
n	[integer(1)] A number smaller or equal length(x).
type	[1 | 2] Either • 1 (default) to split x into n chunks of equal size, • or 2 to split x into chunks of size n.
strict	[logical(1)] Set to TRUE to fail if length(x) is not a multiple of n, or FALSE (default), else.

Value

A list.

See Also

Other vector helpers: check_numeric_vector(), check_probability_vector(), insert_vector_entry(), match_numerics(), permutations(), subsets(), vector_occurrence()

Examples

```
x <- 1:12
chunk_vector(x, n = 3, type = 1)
chunk_vector(x, n = 3, type = 2)
try(chunk_vector(x, n = 5, strict = TRUE))
```

---

correlated_regressors *Simulate correlated regressor values*

Description

This function simulates regressor values from various marginal distributions with custom correlations.

Usage

```
correlated_regressors(
  labels,
  n = 100,
  marginals = list(),
  correlation = diag(length(labels)),
  verbose = FALSE
)
```

Arguments

`labels`	[`character()`] Unique labels for the regressors.
`n`	[`integer(1)`] The number of values per regressor.
`marginals`	[`list()`] Optionally marginal distributions for regressors. If not specified, standard normal marginal distributions are used. Each list entry must be named according to a regressor label, and the following distributions are currently supported: **discrete distributions** • Poisson: `list(type = "poisson", lambda = ...)` • categorical: `list(type = "categorical", p = c(...))` **continuous distributions** • normal: `list(type = "normal", mean = ..., sd = ...)` • uniform: `list(type = "uniform", min = ..., max = ...)`
`correlation`	[`matrix()`] A correlation matrix of dimension `length(labels)`, where the `(p, q)`-th entry defines the correlation between regressor `labels[p]` and `labels[q]`.
`verbose`	[`logical(1)`] Print information about the simulated regressors?

Value

A `data.frame` with `n` rows and `length(labels)` columns.

References

This function heavily depends on the `{SimMultiCorrData}` package.

See Also

Other simulation helpers: `ddirichlet_cpp()`, `dmvnorm_cpp()`, `dtnorm_cpp()`, `dwishart_cpp()`, `simulate_markov_chain()`

Examples

```
labels <- c("P", "C", "N1", "N2", "U")
n <- 100
marginals <- list(
  "P" = list(type = "poisson", lambda = 2),
  "C" = list(type = "categorical", p = c(0.3, 0.2, 0.5)),
  "N1" = list(type = "normal", mean = -1, sd = 2),
  "U" = list(type = "uniform", min = -2, max = -1)
)
correlation <- matrix(
  c(1, -0.3, -0.1, 0, 0.5,
    -0.3, 1, 0.3, -0.5, -0.7,
    -0.1, 0.3, 1, -0.3, -0.3,
```

```
    0, -0.5, -0.3, 1, 0.1,
    0.5, -0.7, -0.3, 0.1, 1),
  nrow = 5, ncol = 5
)
data <- correlated_regressors(
  labels = labels, n = n, marginals = marginals, correlation = correlation
)
head(data)
```

---

cov_to_chol *Cholesky root of covariance matrix*

Description

These functions compute the Cholesky root elements of a covariance matrix and, conversely, build a covariance matrix from its Cholesky root elements.

Usage

```
cov_to_chol(cov, unique = TRUE)

chol_to_cov(chol)

unique_chol(chol)
```

Arguments

cov	[matrix()] A covariance matrix.
unique	[logical(1)] Ensure that the Cholesky decomposition is unique by restricting the diagonal elements to be positive?
chol	[numeric()] Cholesky root elements.

Value

For cov_to_chol a numeric vector of Cholesky root elements.

For chol_to_cov a covariance matrix.

See Also

Other matrix helpers: check_correlation_matrix(), check_covariance_matrix(), check_transition_probability_
diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(), sample_correlation_matr
sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
cov <- sample_covariance_matrix(4)
chol <- cov_to_chol(cov)
all.equal(cov, chol_to_cov(chol))
```

`ddirichlet_cpp` *Dirichlet distribution*

Description

The function `ddirichlet()` computes the density of a Dirichlet distribution.

The function `rdirichlet()` samples from a Dirichlet distribution.

The functions with suffix `_cpp` perform no input checks, hence are faster.

Usage

```
ddirichlet_cpp(x, concentration, log = FALSE)

rdirichlet_cpp(concentration)

ddirichlet(x, concentration, log = FALSE)

rdirichlet(n = 1, concentration)
```

Arguments

`x`	[`numeric()`] A probability vector.
`concentration`	[`numeric()`] A concentration vector of the same length as `x`.
`log`	[`logical(1)`] Return the logarithm of the density value?
`n`	[`integer(1)`] The number of samples.

Value

For `ddirichlet()`: The density value.

For `rdirichlet()`: If `n = 1` a `vector` of length `p`, else a `matrix` of dimension `n` times `p` with samples as rows.

See Also

Other simulation helpers: `correlated_regressors()`, `dmvnorm_cpp()`, `dtnorm_cpp()`, `dwishart_cpp()`, `simulate_markov_chain()`

Examples

```
x <- c(0.5, 0.3, 0.2)
concentration <- 1:3

# compute density
ddirichlet(x = x, concentration = concentration)
ddirichlet(x = x, concentration = concentration, log = TRUE)

# sample
rdirichlet(concentration = 1:3)
rdirichlet(n = 4, concentration = 1:2)
```

---

delete_data_frame_columns

Deleting `data.frame` *columns*

Description

This function deletes columns of a `data.frame` by name.

Usage

```
delete_data_frame_columns(df, column_names)
```

Arguments

`df`	[`data.frame`] A `data.frame`.
`column_names`	[`character()`] The name(s) of column(s) of `df` to delete.

Value

The input `df` without the columns defined by `column_names`.

See Also

Other data.frame helpers: `group_data_frame()`

Examples

```
df <- data.frame("label" = c("A", "B"), "number" = 1:10)
delete_data_frame_columns(df = df, column_names = "label")
delete_data_frame_columns(df = df, column_names = "number")
delete_data_frame_columns(df = df, column_names = c("label", "number"))
```

`Dictionary` *Dictionary R6 Object*

Description

Provides a simple key-value interface based on R6.

Active bindings

`keys [character()]`
: Available keys.

`alias [list()]`
: Available keys per alias value.

Methods

Public methods:

- `Dictionary$new()`
- `Dictionary$add()`
- `Dictionary$get()`
- `Dictionary$remove()`
- `Dictionary$print()`

Method `new()`**:** Initializing a new `Dictionary` object.

Usage:

```
Dictionary$new(
  key_name,
  alias_name = NULL,
  value_names = character(),
  value_assert = alist(),
  allow_overwrite = TRUE,
  keys_reserved = character(),
  alias_choices = NULL,
  dictionary_name = NULL
)
```

Arguments:

`key_name [character(1)]`
: The name for the key variable.

`alias_name [NULL | character(1)]`
: Optionally the name for the alias variable.

`value_names [character(0)]`
: The names of the values connected to a key.

`value_assert [alist(1)]`
: For each element in `value_names`, `values_assert` *can* have an identically named element of the form `checkmate::assert_*(...)`, where `...` can be any argument for the assertion function except for the `x` argument.

`allow_overwrite [logical(1)]`
: Allow overwriting existing keys with new values? Duplicate keys are never allowed.

`keys_reserved [character()]`
: Names that must not be used as keys.

`alias_choices [NULL or character()]`
: Optionally possible values for the alias. Can also be `NULL`, then all alias values are allowed.

`dictionary_name [NULL or character()]`
: Optionally the name for the dictionary.

Method `add()`**:** Adding an element to the dictionary.

Usage:

```
Dictionary$add(...)
```

Arguments:

`...` Values for

- the key variable `key_name` (must be a single `character`),
- the alias variable `alias_name` (optionally, must then be a `character vector`),
- all the variables specified for `value_names` (if any, they must comply to the `value_assert` checks).

Method `get()`**:** Getting elements from the dictionary.

Usage:

```
Dictionary$get(key, value = NULL)
```

Arguments:

`key [character(1)]`
: A value for the key variable `key_name`. Use the `$keys` method for available keys.

`value [NULL | character(1)]`
: One of the elements in `value_names`, selecting the required value. Can also be `NULL` (default) for all values connected to the `key`, returned as a `list`.

Method `remove()`**:** Removing elements from the dictionary (and associated alias, if any).

Usage:

```
Dictionary$remove(key)
```

Arguments:

`key [character(1)]`
: A value for the key variable `key_name`. Use the `$keys` method for available keys.

Method `print()`**:** Printing details of the dictionary.

Usage:

```
Dictionary$print()
```

See Also

Other package helpers: Storage, identical_structure(), input_check_response(), match_arg(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

Examples

```
# TODO
```

`diff_cov` *Difference and un-difference covariance matrix*

Description

These functions difference and un-difference a covariance matrix with respect to row `ref`.

Usage

```
diff_cov(cov, ref = 1)

undiff_cov(cov_diff, ref = 1)

delta(ref = 1, dim)
```

Arguments

`cov`, `cov_diff`	[`matrix()`] A (differenced) covariance matrix of dimension `dim` (or `dim` - 1, respectively).
`ref`	[`integer(1)`] The reference row between 1 and `dim` for differencing that maps `cov` to `cov_diff`, see details.
`dim`	[`integer(1)`] The matrix dimension.

Details

For differencing: Let Σ be a covariance matrix of dimension n. Then

$$\tilde{\Sigma} = \Delta_k \Sigma \Delta_k'$$

is the differenced covariance matrix with respect to row $k = 1, \ldots, n$, where Δ_k is a difference operator that depends on the reference row k. More precise, Δ_k the identity matrix of dimension n without row k and with -1s in column k. It can be computed with `delta(ref = k, dim = n)`.

For un-differencing: The "un-differenced" covariance matrix Σ cannot be uniquely computed from $\tilde{\Sigma}$. For a non-unique solution, we add a column and a row of zeros at column and row number k to $\tilde{\Sigma}$, respectively, and add 1 to each matrix entry to make the result a proper covariance matrix.

Value

A (differenced or un-differenced) covariance `matrix`.

See Also

Other matrix helpers: `check_correlation_matrix()`, `check_covariance_matrix()`, `check_transition_probability_matrix()`, `cov_to_chol()`, `insert_matrix_column()`, `matrix_diagonal_indices()`, `matrix_indices()`, `sample_correlation_matrix()`, `sample_covariance_matrix()`, `sample_transition_probability_matrix()`, `stationary_distribution()`

Examples

```
n <- 3
Sigma <- sample_covariance_matrix(dim = n)
k <- 2

# build difference operator
delta(ref = k, dim = n)

# difference Sigma
(Sigma_diff <- diff_cov(Sigma, ref = k))

# un-difference Sigma
undiff_cov(Sigma_diff, ref = k)
```

---

dmvnorm_cpp *Multivariate normal distribution*

Description

The function `dmvnorm()` computes the density of a multivariate normal distribution.

The function `rmvnorm()` samples from a multivariate normal distribution.

The functions with suffix _cpp perform no input checks, hence are faster.

Usage

```
dmvnorm_cpp(x, mean, Sigma, log = FALSE)

rmvnorm_cpp(mean, Sigma, log = FALSE)

dmvnorm(x, mean, Sigma, log = FALSE)

rmvnorm(n = 1, mean, Sigma, log = FALSE)
```

Arguments

x	[numeric()] A quantile vector of length p.
mean	[numeric()] The mean vector of length p.
Sigma	[matrix()] The covariance matrix of dimension p.
log	[logical(1)] Return the logarithm of the density value?
n	[integer(1)] An integer, the number of samples.

Value

For dmvnorm(): The density value.

For rmvnorm(): If n = 1 a vector of length p, else a matrix of dimension n times p with samples as rows.

See Also

Other simulation helpers: correlated_regressors(), ddirichlet_cpp(), dtnorm_cpp(), dwishart_cpp(), simulate_markov_chain()

Examples

```
x <- c(0, 0)
mean <- c(0, 0)
Sigma <- diag(2)

# compute density
dmvnorm(x = x, mean = mean, Sigma = Sigma)
dmvnorm(x = x, mean = mean, Sigma = Sigma, log = TRUE)

# sample
rmvnorm(n = 3, mean = mean, Sigma = Sigma)
rmvnorm(mean = mean, Sigma = Sigma, log = TRUE)
```

---

do.call_timed *Measure computation time*

Description

This function measures the computation time of a call.

Usage

```
do.call_timed(what, args, units = "secs")
```

Arguments

what, args	Passed to do.call.
units	Passed to difftime.

Details

This function is a wrapper for do.call.

Value

A list of the two elements "result" (the results of the do.call call) and "time" (the computation time).

See Also

Other function helpers: function_arguments(), function_body(), function_defaults(), quiet(), timed(), try_silent(), variable_name()

Examples

```
## Not run:
what <- function(s) {
  Sys.sleep(s)
  return(s)
}
args <- list(s = 1)
do.call_timed(what = what, args = args)

## End(Not run)
```

---

dtnorm_cpp *Truncated normal distribution*

Description

The function dtnorm() computes the density of a truncated normal distribution.

The function rtnorm() samples from a truncated normal distribution.

The function dttnorm() and rttnorm() compute the density and sample from a two-sided truncated normal distribution, respectively.

The functions with suffix _cpp perform no input checks, hence are faster.

Usage

```
dtnorm_cpp(x, mean, sd, point, above, log = FALSE)

dttnorm_cpp(x, mean, sd, lower, upper, log = FALSE)

rtnorm_cpp(mean, sd, point, above, log = FALSE)

rttnorm_cpp(mean, sd, lower, upper, log = FALSE)

dtnorm(x, mean, sd, point, above, log = FALSE)

dttnorm(x, mean, sd, lower, upper, log = FALSE)

rtnorm(mean, sd, point, above, log = FALSE)

rttnorm(mean, sd, lower, upper, log = FALSE)
```

Arguments

x	[numeric(1)] A quantile.
mean	[numeric(1)] The mean.
sd	[numeric(1)] The non-negative standard deviation.
point, lower, upper	[numeric(1)] The truncation point.
above	[logical(1)] Truncate from above? Else, from below.
log	[logical(1)] Return the logarithm of the density value?

Value

For `dtnorm()` and `dttnorm()`: The density value.

For `rtnorm()` and `rttnorm()`: The random draw

See Also

Other simulation helpers: `correlated_regressors()`, `ddirichlet_cpp()`, `dmvnorm_cpp()`, `dwishart_cpp()`, `simulate_markov_chain()`

Examples

```
x <- c(0, 0)
mean <- c(0, 0)
```

```
Sigma <- diag(2)

# compute density
dmvnorm(x = x, mean = mean, Sigma = Sigma)
dmvnorm(x = x, mean = mean, Sigma = Sigma, log = TRUE)

# sample
rmvnorm(n = 3, mean = mean, Sigma = Sigma)
rmvnorm(mean = mean, Sigma = Sigma, log = TRUE)
```

`dwishart_cpp` *Wishart distribution*

Description

The function `dwishart()` computes the density of a Wishart distribution.

The function `rwishart()` samples from a Wishart distribution.

The functions with suffix _cpp perform no input checks, hence are faster.

Usage

```
dwishart_cpp(x, df, scale, log = FALSE, inv = FALSE)

rwishart_cpp(df, scale, inv = FALSE)

dwishart(x, df, scale, log = FALSE, inv = FALSE)

rwishart(df, scale, inv = FALSE)
```

Arguments

`x`	[`matrix()`] A covariance matrix of dimension `p`.
`df`	[`integer()`] The degrees of freedom greater of equal `p`.
`scale`	[`matrix()`] The scale covariance matrix of dimension `p`.
`log`	[`logical(1)`] Return the logarithm of the density value?
`inv`	[`logical(1)`] Use this inverse Wishart distribution?

Value

For `dwishart()`: The density value.

For `rwishart()`: A `matrix`, the random draw.

See Also

Other simulation helpers: correlated_regressors(), ddirichlet_cpp(), dmvnorm_cpp(), dtnorm_cpp(), simulate_markov_chain()

Examples

```
x <- diag(2)
df <- 4
scale <- diag(2)

# compute density
dwishart(x = x, df = df, scale = scale)
dwishart(x = x, df = df, scale = scale, log = TRUE)
dwishart(x = x, df = df, scale = scale, inv = TRUE)

# sample
rwishart(df = df, scale = scale)
rwishart(df = df, scale = scale, inv = TRUE)
```

---

function_arguments *Get function arguments*

Description

This function returns the names of function arguments.

Usage

```
function_arguments(f, with_default = TRUE, with_ellipsis = TRUE)
```

Arguments

f	[function] A function.
with_default	[logical(1)] Include function arguments that have default values?
with_ellipsis	[logical(1)] Include the "..." argument if present?

Value

A character vector.

See Also

Other function helpers: do.call_timed(), function_body(), function_defaults(), quiet(), timed(), try_silent(), variable_name()

Examples

```
f <- function(a, b = 1, c = "", ...) { }
function_arguments(f)
function_arguments(f, with_default = FALSE)
function_arguments(f, with_ellipsis = FALSE)
```

`function_body` *Extract function body*

Description

This function extracts the body of a function as a single `character`.

Usage

```
function_body(fun, braces = FALSE, nchar = getOption("width") - 4)
```

Arguments

`fun`	[`function`] A `function`.
`braces`	[`logical(1)`] Remove `"{"` and `"}"` at start and end (if any)?
`nchar`	[`integer(1)`] The maximum number of characters before abbreviation, at least `3`.

Value

A `character`, the body of `f`.

See Also

Other function helpers: `do.call_timed()`, `function_arguments()`, `function_defaults()`, `quiet()`, `timed()`, `try_silent()`, `variable_name()`

Examples

```
fun <- mean.default
function_body(fun)
function_body(fun, braces = TRUE)
function_body(fun, nchar = 30)
```

function_defaults *Get default function arguments*

Description

This function returns the default function arguments (if any).

Usage

```
function_defaults(f, exclude = NULL)
```

Arguments

f	[function] A function.
exclude	[NULL | character()] Argument names to exclude. Can be NULL (default) to not exclude any argument names.

Value

A named list.

See Also

Other function helpers: do.call_timed(), function_arguments(), function_body(), quiet(), timed(), try_silent(), variable_name()

Examples

```
f <- function(a, b = 1, c = "", ...) { }
function_defaults(f)
function_defaults(f, exclude = "b")
```

group_data_frame *Grouping of a* data.frame

Description

This function groups a data.frame according to values of one column.

Usage

```
group_data_frame(df, by, keep_by = TRUE)
```

Arguments

df	[data.frame] A data.frame.
by	[character(1)] The name of a column of df to group by.
keep_by	[logical(1)] Keep the grouping column by?

Value

A list of data.frames, subsets of df.

See Also

Other data.frame helpers: delete_data_frame_columns()

Examples

```
df <- data.frame("label" = c("A", "B"), "number" = 1:10)
group_data_frame(df = df, by = "label")
group_data_frame(df = df, by = "label", keep_by = FALSE)
```

---

identical_structure *Check if two objects have identical structure*

Description

This function determines whether two objects have the same structure,

- which includes the mode, class and dimension
- but does *not* include concrete values or attributes.

Usage

```
identical_structure(x, y)
```

Arguments

x, y	[any] Two objects.

Value

Either TRUE if x and y have the same structure, and FALSE, else.

References

Inspired by https://stackoverflow.com/a/45548885/15157768.

See Also

Other package helpers: Dictionary, Storage, input_check_response(), match_arg(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

Examples

```
identical_structure(integer(1), 1L)
identical_structure(diag(2), matrix(rnorm(4), 2, 2))
identical_structure(diag(2), data.frame(diag(2)))
```

---

input_check_response *Standardized response to an input check*

Description

This function provides standardized responses to input checks, ensuring consistency.

Usage

```
input_check_response(
  check,
  var_name = NULL,
  error = TRUE,
  prefix = "Input {.var {var_name}} is bad:"
)
```

Arguments

check	[TRUE | character(1)] Matches the return value of the check* functions from the {checkmate} package, i.e., either TRUE if the check was successful, or a character (the error message) else.
var_name	[NULL | character(1)] Optionally specifies the name of the input being checked. This name will be used for the default value of the prefix argument.
error	[logical(1)] If check is not TRUE, throw an error?
prefix	[character(1)] A prefix for the thrown error message, if check is not TRUE and error is TRUE.

Value

TRUE if check is TRUE. If check is not TRUE, depending on error:

- If error is TRUE, throws an error.
- If error is FALSE, returns FALSE.

See Also

Other package helpers: Dictionary, Storage, identical_structure(), match_arg(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

Examples

```
x <- "1"
y <- 1

### check is successful
input_check_response(
  check = checkmate::check_character(x),
  var_name = "x",
  error = TRUE
)

### standardized check response
## Not run:
input_check_response(
  check = checkmate::check_character(y),
  var_name = "y",
  error = TRUE
)

## End(Not run)
```

---

insert_matrix_column *Insert column in matrix*

Description

This function inserts a column into a matrix.

Usage

```
insert_matrix_column(A, x, p)
```

Arguments

A	[matrix()] A matrix.
x	[atomic()] The column to be added, of length nrow(A). Can also be a single value.
p	[integer())] The position(s) where to add the column, one or more of:

- p = 0 appends the column left
- p = ncol(A) appends the column right
- p = n inserts the column between the n-th and (n + 1)-th column of A.

Value

A matrix.

See Also

Other matrix helpers: check_correlation_matrix(), check_covariance_matrix(), check_transition_probability_matrix(), cov_to_chol(), diff_cov(), matrix_diagonal_indices(), matrix_indices(), sample_correlation_matrix(), sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
A <- diag(3)
x <- 1:3
insert_matrix_column(A, x, 0)
insert_matrix_column(A, x, 1)
insert_matrix_column(A, x, 2)
insert_matrix_column(A, x, 3)

### also single value
x <- 2
insert_matrix_column(A, x, 0)

### also multiple positions
insert_matrix_column(A, x, 0:3)

### also trivial case
insert_matrix_column(matrix(nrow = 0, ncol = 0), integer(), integer())
```

---

insert_vector_entry *Insert entry in vector*

Description

This function inserts a value into a vector.

Usage

```
insert_vector_entry(v, x, p)
```

Arguments

v	[atomic()] A vector.
x	[atomic(1)] The entry to be added.
p	[integer())] The position(s) where to add the value, one or more of: • p = 0 appends the value left • p = length(v) appends the value right • p = n inserts the value between the n-th and (n + 1)-th entry of v.

Value

A vector.

See Also

Other vector helpers: check_numeric_vector(), check_probability_vector(), chunk_vector(), match_numerics(), permutations(), subsets(), vector_occurrence()

Examples

```
v <- 1:3
x <- 0
insert_vector_entry(v, x, 0)
insert_vector_entry(v, x, 1)
insert_vector_entry(v, x, 2)
insert_vector_entry(v, x, 3)

### also multiple positions
insert_vector_entry(v, x, 0:3)

### also trivial case
insert_vector_entry(integer(), integer(), integer())
```

---

match_arg *Argument matching*

Description

This function matches function arguments and is a modified version of match.arg.

Usage

```
match_arg(arg, choices, several.ok = FALSE, none.ok = FALSE)
```

Arguments

arg	[character()] The function argument.
choices	[character()] Allowed values for arg.
several.ok	[logical(1)] Is arg allowed to have more than one element?
none.ok	[logical(1)] Is arg allowed to have zero elements?

Value

The un-abbreviated version of the exact or unique partial match if there is one. Otherwise, an error is signaled if several.ok is FALSE or none.ok is FALSE. When several.ok is TRUE and (at least) one element of arg has a match, all un-abbreviated versions of matches are returned. When none.ok is TRUE and arg has zero elements, character(0) is returned.

See Also

Other package helpers: Dictionary, Storage, identical_structure(), input_check_response(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

---

match_numerics *Best-possible match of two numeric vectors*

Description

This function matches the indices of two numeric vectors as good as possible (that means with the smallest possible sum of deviations).

Usage

```
match_numerics(x, y)
```

Arguments

x, y	[numeric()] Two vectors of the same length.

Value

An integer vector of length length(x) with the positions of y in x.

See Also

Other vector helpers: check_numeric_vector(), check_probability_vector(), chunk_vector(), insert_vector_entry(), permutations(), subsets(), vector_occurrence()

Examples

```
x <- c(-1, 0, 1)
y <- c(0.1, 1.5, -1.2)
match_numerics(x, y)
```

`matrix_diagonal_indices` *Get indices of matrix diagonal*

Description

This function returns the indices of the diagonal elements of a quadratic matrix.

Usage

```
matrix_diagonal_indices(n, triangular = NULL)
```

Arguments

`n`	[`integer(1)`] The matrix dimension.
`triangular`	[`NULL` or `character(1)`] If `NULL` (default), all elements of the matrix are considered. If `"lower"` (`"upper"`), only the lower- (upper-) triangular matrix is considered.

Value

An `integer vector`.

See Also

Other matrix helpers: `check_correlation_matrix()`, `check_covariance_matrix()`, `check_transition_probability_matrix()`, `cov_to_chol()`, `diff_cov()`, `insert_matrix_column()`, `matrix_indices()`, `sample_correlation_matrix()`, `sample_covariance_matrix()`, `sample_transition_probability_matrix()`, `stationary_distribution()`

Examples

```
# indices of diagonal elements
n <- 3
matrix(1:n^2, n, n)
matrix_diagonal_indices(n)

# indices of diagonal elements of lower-triangular matrix
L <- matrix(0, n, n)
L[lower.tri(L, diag=TRUE)] <- 1:((n * (n + 1)) / 2)
L
matrix_diagonal_indices(n, triangular = "lower")
```

```
# indices of diagonal elements of upper-triangular matrix
U <- matrix(0, n, n)
U[upper.tri(U, diag=TRUE)] <- 1:((n * (n + 1)) / 2)
U
matrix_diagonal_indices(n, triangular = "upper")
```

`matrix_indices` *Get* `matrix` *indices*

Description

This function returns `matrix` indices as `character`.

Usage

```
matrix_indices(x, prefix = "", exclude_diagonal = FALSE)
```

Arguments

`x`	[`matrix`] A `matrix`.
`prefix`	[`character(1)`] A prefix for the indices.
`exclude_diagonal`	[`logical(1)`] Exclude indices where row equals column?

Value

A `character` `vector`.

See Also

Other matrix helpers: `check_correlation_matrix()`, `check_covariance_matrix()`, `check_transition_probability_`
`cov_to_chol()`, `diff_cov()`, `insert_matrix_column()`, `matrix_diagonal_indices()`, `sample_correlation_matrix(`
`sample_covariance_matrix()`, `sample_transition_probability_matrix()`, `stationary_distribution()`

Examples

```
M <- diag(3)
matrix_indices(M)
matrix_indices(M, "M_")
matrix_indices(M, "M_", TRUE)
```

merge_lists *Merge named lists*

Description

This function merges `lists` based on their element names. Elements are only included in the final output `list`, if no former `list` has contributed an element with the same name.

Usage

```
merge_lists(...)
```

Arguments

`...`	One or more named `list`(s).

Value

A `list`.

See Also

Other list helpers: `check_list_of_lists()`

Examples

```
merge_lists(list("a" = 1, "b" = 2), list("b" = 3, "c" = 4, "d" = NULL))
```

package_logo *Creating a basic logo for an R package*

Description

This function creates a basic R package logo. The logo has a white background and the package name (with or without curly brackets) in the center. The font size for the package name is scaled such that it fits inside the logo. Type `?oeli` to see an example.

Usage

```
package_logo(package_name, brackets = TRUE, use_logo = FALSE)
```

Arguments

package_name	[character(1)] The package name.
brackets	[logical(1)] Curly brackets around the package name?
use_logo	[logical(1)] Run use_logo in the end?

Details

The function optionally calls use_logo if use_logo = TRUE to set up the logo for a package.

Value

A ggplot object.

See Also

Other package helpers: Dictionary, Storage, identical_structure(), input_check_response(), match_arg(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

Examples

```
package_logo("my_package", brackets = TRUE, use_logo = FALSE)
```

---

permutations *Build permutations*

Description

This function creates all permutations of a given vector.

Usage

```
permutations(x)
```

Arguments

x	[atomic()] Any vector.

Value

A list of all permutations of x.

References

Modified version of https://stackoverflow.com/a/20199902/15157768.

See Also

Other vector helpers: check_numeric_vector(), check_probability_vector(), chunk_vector(), insert_vector_entry(), match_numerics(), subsets(), vector_occurrence()

Examples

```
permutations(1:3)
permutations(LETTERS[1:3])
```

---

print_matrix *Print (abbreviated)* matrix

Description

This function prints a (possibly abbreviated) matrix.

Usage

```
print_matrix(
  x,
  rowdots = 4,
  coldots = 4,
  digits = 2,
  label = NULL,
  simplify = FALSE,
  details = !simplify
)
```

Arguments

x	[atomic() | matrix] The object to be printed.
rowdots	[integer(1)] The row number which is replaced by ....
coldots	[integer(1)] The column number which is replaced by ....
digits	[integer(1)] The number of printed decimal places if input x is numeric.
label	[character(1)] A label for x. Only printed if simplify = FALSE.
simplify	[logical(1)] Simplify the output?
details	[logical(1)] Print the type and dimension of x?

Value

Invisibly returns `x`.

References

This function is a modified version of `ramify::pprint()`.

See Also

Other package helpers: `Dictionary`, `Storage`, `identical_structure()`, `input_check_response()`, `match_arg()`, `package_logo()`, `renv_development_packages()`, `system_information()`, `unexpected_error()`, `user_confirm()`

Examples

```
print_matrix(x = 1, label = "single numeric")
print_matrix(x = LETTERS[1:26], label = "letters")
print_matrix(x = 1:3, coldots = 2)
print_matrix(x = matrix(rnorm(99), ncol = 1), label = "single column matrix")
print_matrix(x = matrix(1:100, nrow = 1), label = "single row matrix")
print_matrix(x = matrix(LETTERS[1:24], ncol = 6), label = "big matrix")
print_matrix(x = diag(5), coldots = 2, rowdots = 2, simplify = TRUE)
```

---

quiet *Silence R code*

Description

This function silences warnings, messages and any `cat()` or `print()` output from R expressions or functions.

Usage

```
quiet(x, print_cat = TRUE, message = TRUE, warning = TRUE)
```

Arguments

`x`	[expression] Any function or expression or value assignment expression.
`print_cat`	[logical(1)] Silence `print()` and `cat()` outputs?
`message`	[logical(1)] Silence messages?
`warning`	[logical(1)] Silence warnings?

Value

Invisibly the expression `x`.

References

This function is a modified version of `quiet`.

See Also

Other function helpers: `do.call_timed()`, `function_arguments()`, `function_body()`, `function_defaults()`, `timed()`, `try_silent()`, `variable_name()`

Examples

```
f <- function() {
  warning("warning")
  message("message")
  cat("cat")
  print("print")
}
quiet(f())
```

---

renv_development_packages

Using development packages when working with `{renv}`

Description

This function creates a file that loads development packages so that `{renv}` can detect and write them to the lockfile.

Usage

```
renv_development_packages(
  packages = c("covr", "devtools", "DT", "markdown", "R.utils", "yaml"),
  file_name = "development_packages"
)
```

Arguments

`packages`	`[character()]` Package names.
`file_name`	`[character(1)]` The name for the `.R` file.

Value

No return value, but it writes a file to the project root and adds an entry to the `.Rbuildignore` file.

See Also

Other package helpers: Dictionary, Storage, identical_structure(), input_check_response(), match_arg(), package_logo(), print_matrix(), system_information(), unexpected_error(), user_confirm()

`sample_correlation_matrix` *Sample correlation matrix*

Description

This function samples a correlation matrix by sampling a covariance matrix from an inverse Wishart distribution and transforming it to a correlation matrix.

Usage

```
sample_correlation_matrix(dim, df = dim, scale = diag(dim))
```

Arguments

`dim`	[integer(1)] The dimension.
`df`	[integer(1)] The degrees of freedom of the inverse Wishart distribution greater or equal `dim`.
`scale`	[matrix()] The scale covariance matrix of the inverse Wishart distribution of dimension `dim`.

Value

A correlation `matrix`.

See Also

Other matrix helpers: check_correlation_matrix(), check_covariance_matrix(), check_transition_probability_matrix(), cov_to_chol(), diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(), sample_covariance_matrix(), sample_transition_probability_matrix(), stationary_distribution()

Examples

```
sample_correlation_matrix(dim = 3)
```

`sample_covariance_matrix` *Sample covariance matrix*

Description

This function samples a covariance matrix from an inverse Wishart distribution.

Usage

```
sample_covariance_matrix(dim, df = dim, scale = diag(dim), diag = FALSE)
```

Arguments

`dim`	[`integer(1)`] The dimension.
`df`	[`integer(1)`] The degrees of freedom of the inverse Wishart distribution greater or equal `dim`.
`scale`	[`matrix()`] The scale covariance matrix of the inverse Wishart distribution of dimension `dim`.
`diag`	[`logical(1)`] Diagonal matrix?

Value

A covariance `matrix`.

See Also

Other matrix helpers: `check_correlation_matrix()`, `check_covariance_matrix()`, `check_transition_probability_matrix()`, `cov_to_chol()`, `diff_cov()`, `insert_matrix_column()`, `matrix_diagonal_indices()`, `matrix_indices()`, `sample_correlation_matrix()`, `sample_transition_probability_matrix()`, `stationary_distribution()`

Examples

```
sample_covariance_matrix(dim = 3)
```

sample_transition_probability_matrix *Sample transition probability matrices*

Description

This function returns a random, squared matrix of dimension `dim` that fulfills the properties of a transition probability matrix.

Usage

```
sample_transition_probability_matrix(dim, state_persistent = TRUE)
```

Arguments

`dim`	[integer(1)] The dimension.
`state_persistent`	[logical(1)] Put more probability on the diagonal?

Value

A transition probability `matrix`.

See Also

Other matrix helpers: `check_correlation_matrix()`, `check_covariance_matrix()`, `check_transition_probability_`
`cov_to_chol()`, `diff_cov()`, `insert_matrix_column()`, `matrix_diagonal_indices()`, `matrix_indices()`,
`sample_correlation_matrix()`, `sample_covariance_matrix()`, `stationary_distribution()`

Examples

```
sample_transition_probability_matrix(dim = 3)
```

simulate_markov_chain *Simulate Markov chain*

Description

This function simulates a Markov chain.

Usage

```
simulate_markov_chain(Gamma, T, delta = oeli::stationary_distribution(Gamma))
```

Arguments

Gamma	[matrix()] A transition probability matrix.
T	[integer(1)] The length of the Markov chain.
delta	[numeric()] A probability vector, the initial distribution. By default, delta is the stationary distribution of Gamma.

Value

A numeric vector of length T with states.

See Also

Other simulation helpers: correlated_regressors(), ddirichlet_cpp(), dmvnorm_cpp(), dtnorm_cpp(), dwishart_cpp()

Examples

```
Gamma <- sample_transition_probability_matrix(dim = 3)
simulate_markov_chain(Gamma = Gamma, T = 10)
```

---

stationary_distribution

Stationary distribution

Description

This function computes the stationary distribution corresponding to a transition probability matrix.

Usage

```
stationary_distribution(tpm, soft_fail = FALSE)
```

Arguments

tpm	[matrix()] A transition probability matrix.
soft_fail	[logical(1)] Return the discrete uniform distribution if the computation of the stationary distribution fails for some reason? Else, throw an error.

Value

A numeric vector.

See Also

Other matrix helpers: check_correlation_matrix(), check_covariance_matrix(), check_transition_probability_
cov_to_chol(), diff_cov(), insert_matrix_column(), matrix_diagonal_indices(), matrix_indices(),
sample_correlation_matrix(), sample_covariance_matrix(), sample_transition_probability_matrix()

Examples

```
tpm <- matrix(0.05, nrow = 3, ncol = 3)
diag(tpm) <- 0.9
stationary_distribution(tpm)
```

Storage *Storage R6 Object*

Description

Provides a simple indexing interface for list elements based on R6. Basically, it allows to store items in a list and to regain them based on identifiers defined by the user.

Value

The output depends on the method:

- `$new()` returns a `Storage` object.
- `$add()`, `$remove()`, and `$print()` invisibly return the `Storage` object (to allow for method chaining)
- `$get()` returns the requested element(s)
- `$number()` returns an `integer`
- `$indices()` return an `integer vector`

Setting identifiers

An identifier is a `character`, typically a binary property. Identifiers can be negated by placing an exclamation mark (`"!"`) in front of them. Identifiers that have been assigned to other elements previously do not need to be specified again for new elements; instead, a default value can be used. This default value can be defined either globally for all cases (via the `$missing_identifier` field) or separately for each specific case (via the method argument).

User confirmation

If desired, the user can be asked for confirmation when adding, extracting, or removing elements using identifiers. This behavior can be set globally through the `$confirm` field or customized separately for each specific case via the method argument.

Active bindings

`identifier [character()]`
: The identifiers used.

`confirm [logical(1)]`
: The default value for confirmations.

`missing_identifier [logical(1)]`
: The default value for not specified identifiers.

`hide_warnings [logical(1)]`
: Hide warnings (for example if unknown identifiers are selected)?

Methods

Public methods:

- `Storage$new()`
- `Storage$add()`
- `Storage$get()`
- `Storage$remove()`
- `Storage$number()`
- `Storage$indices()`
- `Storage$print()`

Method `new()`**:** Initializing a `Storage` object.

Usage:

```
Storage$new()
```

Method `add()`**:** Adding an element.

Usage:

```
Storage$add(
  x,
  identifier,
  confirm = interactive() & self$confirm,
  missing_identifier = self$missing_identifier
)
```

Arguments:

`x [any()]`
: An object to be saved.

`identifier [character()]`
: Pne or more identifiers (the identifier `"all"` is reserved to select all elements).

`confirm [logical(1)]`
: Prompted for confirmation?

`missing_identifier [logical(1)|NA]`
: The value for not specified identifiers.

Method `get()`**:** Getting elements.

Usage:

```
Storage$get(
  identifier = character(),
  ids = integer(),
  logical = "and",
  confirm = interactive() & self$confirm,
  missing_identifier = self$missing_identifier,
  id_names = FALSE
)
```

Arguments:

`identifier [character()]`
: Pne or more identifiers (the identifier `"all"` is reserved to select all elements).

`ids [integer()]`
: One or more ids.

`logical [character(1)]`
: In the case that multiple identifiers are selected, how should they be combined? Options are:
  - `"and"` (the default): the identifiers are combined with logical and (all identifiers must be `TRUE`)
  - `"or"`: the identifiers are combined with logical or (at least one identifier must be `TRUE`)

`confirm [logical(1)]`
: Prompted for confirmation?

`missing_identifier [logical(1)|NA]`
: The value for not specified identifiers.

`id_names [logical(1)]`
: Name the elements according to their ids?

Method `remove()`**:** removing elements

Usage:

```
Storage$remove(
  identifier = character(),
  ids = integer(),
  logical = "and",
  confirm = interactive() & self$confirm,
  missing_identifier = self$missing_identifier,
  shift_ids = TRUE
)
```

Arguments:

`identifier [character()]`
: Pne or more identifiers (the identifier `"all"` is reserved to select all elements).

`ids [integer()]`
: One or more ids.

`logical [character(1)]`
: In the case that multiple identifiers are selected, how should they be combined? Options are:

- "and" (the default): the identifiers are combined with logical and (all identifiers must be TRUE)
- "or": the identifiers are combined with logical or (at least one identifier must be TRUE)

confirm [logical(1)]
: Prompted for confirmation?

missing_identifier [logical(1) | NA]
: The value for not specified identifiers.

shift_ids [logical(1)]
: Shift ids when in-between elements are removed?

Method number(): Computing the number of identified elements.

Usage:

```
Storage$number(
  identifier = "all",
  missing_identifier = self$missing_identifier,
  logical = "and",
  confirm = FALSE
)
```

Arguments:

identifier [character()]
: Pne or more identifiers (the identifier "all" is reserved to select all elements).

missing_identifier [logical(1) | NA]
: The value for not specified identifiers.

logical [character(1)]
: In the case that multiple identifiers are selected, how should they be combined? Options are:
  - "and" (the default): the identifiers are combined with logical and (all identifiers must be TRUE)
  - "or": the identifiers are combined with logical or (at least one identifier must be TRUE)

confirm [logical(1)]
: Prompted for confirmation?

Method indices(): Returning indices based on defined identifiers.

Usage:

```
Storage$indices(
  identifier = "all",
  logical = "and",
  confirm = interactive() & self$confirm
)
```

Arguments:

identifier [character()]
: Pne or more identifiers (the identifier "all" is reserved to select all elements).

logical [character(1)]
: In the case that multiple identifiers are selected, how should they be combined? Options are:

- "and" (the default): the identifiers are combined with logical and (all identifiers must be TRUE)
- "or": the identifiers are combined with logical or (at least one identifier must be TRUE)

`confirm [logical(1)]`
Prompted for confirmation?

Method `print()`**:** Printing details of the saved elements.

Usage:

```
Storage$print(...)
```

Arguments:

`...` Currently not used.

See Also

Other package helpers: Dictionary, identical_structure(), input_check_response(), match_arg(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error(), user_confirm()

Examples

```
### 1. Create a `Storage` object:
my_storage <- Storage$new()

# 2. Add elements along with identifiers:
my_storage$
  add(42, c("number", "rational"))$
  add(pi, c("number", "!rational"))$
  add("fear of black cats", c("text", "!rational"))$
  add("wearing a seat belt", c("text", "rational"))$
  add(mean, "function")

# 3. What elements are stored?
print(my_storage)

# 4. Extract elements based on identifiers:
my_storage$get("rational")
my_storage$get("!rational")
my_storage$get(c("text", "!rational"))
my_storage$get("all") # get all elements
my_storage$get(c("text", "!text"))
my_storage$get(c("text", "!text"), logical = "or")

# 5. Extract elements based on ids:
my_storage$get(ids = 4:5)
my_storage$get(ids = 4:5, id_names = TRUE) # add the ids as names
```

subsets *Generate vector subsets*

Description

This function generates subsets of a vector.

Usage

```
subsets(v, n = seq_along(v))
```

Arguments

v	[atomic()‘] A vector of elements.
n	[integer(1)‘] The requested subset sizes.

Value

A `list`, each element is a subset of `v`.

See Also

Other vector helpers: `check_numeric_vector()`, `check_probability_vector()`, `chunk_vector()`, `insert_vector_entry()`, `match_numerics()`, `permutations()`, `vector_occurrence()`

Examples

```
v <- 1:3
subsets(v)
subsets(v, c(1, 3)) # only subsets of length 1 or 3
subsets(integer())  # trivial case works
```

system_information *General system level information*

Description

This function returns a `list` of general system level information.

Usage

```
system_information()
```

Value

A `list` with elements:

- `maschine`, the model name of the device
- `cores`, the number of cores
- `ram`, the size of the RAM
- `os`, the operating system
- `rversion`, the R version used

See Also

Other package helpers: `Dictionary`, `Storage`, `identical_structure()`, `input_check_response()`, `match_arg()`, `package_logo()`, `print_matrix()`, `renv_development_packages()`, `unexpected_error()`, `user_confirm()`

Examples

```
system_information()
```

---

timed *Interrupt long evaluations*

Description

This function interrupts an evaluation after a certain number of seconds. Note the limitations documented in `setTimeLimit`.

Usage

```
timed(expression, seconds = Inf, on_time_out = "silent")
```

Arguments

`expression`	[`expression`] An R expression to be evaluated.
`seconds`	[`numeric(1)`] The number of seconds.
`on_time_out`	[`character(1)`] Defines what action to take if the evaluation time exceeded, either: • `"error"` to throw an error exception • `"warning"` to return `NULL` along with a warning • `"silent"` (the default) to just return `NULL`

Value

The value of `expression` or, if the evaluation time exceeded, whatever is specified for `on_time_out`.

See Also

Other function helpers: do.call_timed(), function_arguments(), function_body(), function_defaults(), quiet(), try_silent(), variable_name()

Examples

```
foo <- function(x) {
  for (i in 1:10) Sys.sleep(x / 10)
  return(x)
}
timed(foo(0.5), 1)
timed(foo(1.5), 1)
```

---

try_silent *Try an expression silently*

Description

This function tries to execute expr and returns a string with the error message if the execution failed.

Usage

```
try_silent(expr)
```

Arguments

expr	[expression] An R expression to be evaluated.

Details

This function is a wrapper for try.

Value

Either the value of expr or in case of a failure an object of class fail, which contains the error message.

See Also

Other function helpers: do.call_timed(), function_arguments(), function_body(), function_defaults(), quiet(), timed(), variable_name()

Examples

```
## Not run:
try_silent(1 + 1)
try_silent(1 + "1")

## End(Not run)
```

`unexpected_error` *Handling of an unexpected error*

Description

This function reacts to an unexpected error by throwing an error and linking to an issue site with the request to submit an issue.

Usage

```
unexpected_error(
  msg = "Ups, an unexpected error occured.",
  issue_link = "https://github.com/loelschlaeger/oeli/issues"
)
```

Arguments

`msg`	[`character(1)`] An error message.
`issue_link`	[`character(1)`] The URL to an issues site.

Value

No return value, but it throws an error.

See Also

Other package helpers: `Dictionary`, `Storage`, `identical_structure()`, `input_check_response()`, `match_arg()`, `package_logo()`, `print_matrix()`, `renv_development_packages()`, `system_information()`, `user_confirm()`

user_confirm *User confirmation*

Description

This function asks in an interactive question a binary question.

Usage

```
user_confirm(question = "Question?", default = FALSE)
```

Arguments

question	[character(1)] The binary question to ask. It should end with a question mark.
default	[logical(1)] The default decision.

Value

Either TRUE or FALSE.

See Also

Other package helpers: Dictionary, Storage, identical_structure(), input_check_response(), match_arg(), package_logo(), print_matrix(), renv_development_packages(), system_information(), unexpected_error()

variable_name *Determine variable name*

Description

This function tries to determine the name of a variable passed to a function.

Usage

```
variable_name(variable, fallback = "unnamed")
```

Arguments

variable	[any] Any object.
fallback	[character(1)] A fallback name if for some reason the actual variable name (which must be a single character) cannot be determined.

Value

A character, the variable name.

See Also

Other function helpers: do.call_timed(), function_arguments(), function_body(), function_defaults(), quiet(), timed(), try_silent()

Examples

```
variable_name(a)
f <- function(x) variable_name(x)
f(x = a)
```

---

vector_occurrence *Find the positions of first or last occurrence of unique vector elements*

Description

This function finds the positions of first or last occurrence of unique vector elements.

Usage

```
vector_occurrence(x, type = "first")
```

Arguments

x	[atomic()] A vector.
type	[character(1)] Either "first" for the first or "last" for the last occurrence.

Value

An integer vector, the positions of the unique vector elements. The ordering corresponds to unique(x), i.e., the i-th element in the output is the (first or last) occurrence of the i-th element from unique(x).

See Also

Other vector helpers: check_numeric_vector(), check_probability_vector(), chunk_vector(), insert_vector_entry(), match_numerics(), permutations(), subsets()

Examples

```
x <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
unique(x)
vector_occurrence(x, "first")
vector_occurrence(x, "last")
```

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