

Package: holodeck (via r-universe)

August 22, 2024

Title A Tidy Interface for Simulating Multivariate Data

Version 0.2.2.9000

Description Provides pipe-friendly (`%>%`) wrapper functions for `MASS::mvrnorm()` to create simulated multivariate data sets with groups of variables with different degrees of variance, covariance, and effect size.

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Encoding UTF-8

Imports dplyr, tibble, MASS, purrr, rlang, assertthat

RoxygenNote 7.2.3

URL <https://github.com/Aariq/holodeck>

BugReports <https://github.com/Aariq/holodeck/issues>

Suggests testthat, covr, knitr, rmarkdown, mice, ggplot2

VignetteBuilder knitr

Repository <https://aariq.r-universe.dev>

RemoteUrl <https://github.com/Aariq/holodeck>

RemoteRef HEAD

RemoteSha 638f46e6c7d6726837f7ec4df2380238f7c5cf7f

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|-----------------|----------------------------|
| <code>:=</code> | <i>Definition operator</i> |
|-----------------|----------------------------|

Description

Internally, this package uses the definition operator, `:=`, to make assignments that require computing on the LHS.

Arguments

| | |
|-----------------------|--|
| <code>x</code> | An object to test. |
| <code>lhs, rhs</code> | Expressions for the LHS and RHS of the definition. |

| | |
|-----------------------|---|
| <code>set_diag</code> | <i>Pipe friendly wrapper to 'diag(x) <- value'</i> |
|-----------------------|---|

Description

Pipe friendly wrapper to `'diag(x) <- value'`

Usage

```
set_diag(x, value)
```

Arguments

| | |
|--------------------|---|
| <code>x</code> | a matrix |
| <code>value</code> | either a single value or a vector of length equal to the diagonal of <code>'x'</code> . |

Value

a matrix

Examples

```
library(dplyr)
matrix(0,3,3) %>%
  set_diag(1)
```

| | |
|---------|----------------------------------|
| sim_cat | <i>Simulate categorical data</i> |
|---------|----------------------------------|

Description

This is a simple wrapper that creates a tibble of length 'n_obs' with a single column 'groups'. It will warn if there are fewer than three replicates per group.

Usage

```
sim_cat(.data = NULL, n_obs = NULL, n_groups, name = "group")
```

Arguments

| | |
|----------|---|
| .data | An optional dataframe. If a dataframe is supplied, simulated categorical data will be added to the dataframe. Either '.data' or 'n_obs' must be supplied. |
| n_obs | Total number of observations/rows to simulate if '.data' is not supplied. |
| n_groups | How many groups or treatments to simulate. |
| name | The column name for the grouping variable. Defaults to "group". |

Details

To-do:

- Make this optionally create multiple categorical variables as being nested or crossed or random

Value

a tibble

See Also

[sim_covar](#), [sim_discr](#)

Other multivariate normal functions: [sim_covar\(\)](#), [sim_discr\(\)](#)

Examples

```
df <- sim_cat(n_obs = 30, n_groups = 3)
```

`sim_covar`*Simulate co-varying variables*

Description

Adds a group of variables (columns) with a given variance and covariance to a data frame or tibble

Usage

```
sim_covar(.data = NULL, n_obs = NULL, n_vars, var, cov, name = NA, seed = NA)
```

Arguments

| | |
|---------------------|---|
| <code>.data</code> | An optional dataframe. If a dataframe is supplied, simulated categorical data will be added to the dataframe. Either <code>‘.data‘</code> or <code>‘n_obs‘</code> must be supplied. |
| <code>n_obs</code> | Total number of observations/rows to simulate if <code>‘.data‘</code> is not supplied. |
| <code>n_vars</code> | Number of variables to simulate. |
| <code>var</code> | Variance used to construct variance-covariance matrix. |
| <code>cov</code> | Covariance used to construct variance-covariance matrix. |
| <code>name</code> | An optional name to be appended to the column names in the output. |
| <code>seed</code> | An optional seed for random number generation. If <code>‘NA‘</code> (default) a random seed will be used. |

Value

a tibble

See Also

[sim_cat](#), [sim_discr](#)

Other multivariate normal functions: [sim_cat\(\)](#), [sim_discr\(\)](#)

Examples

```
library(dplyr)
sim_cat(n_obs = 30, n_groups = 3) %>%
  sim_covar(n_vars = 5, var = 1, cov = 0.5, name = "correlated")
```

| | |
|-----------|--|
| sim_discr | <i>Simulate co-varying variables with different means by group</i> |
|-----------|--|

Description

To-do: make this work with 'dplyr::group_by()' instead of 'group ='

Usage

```
sim_discr(.data, n_vars, var, cov, group_means, name = NA, seed = NA)
```

Arguments

| | |
|-------------|--|
| .data | A dataframe containing a grouping variable column. |
| n_vars | Number of variables to simulate. |
| var | Variance used to construct variance-covariance matrix. |
| cov | Covariance used to construct variance-covariance matrix. |
| group_means | A vector of the same length as the number of grouping variables. |
| name | An optional name to be appended to the column names in the output. |
| seed | An optional seed for random number generation. If 'NA' (default) a random seed will be used. |

Value

a tibble

See Also

[sim_cat](#), [sim_covar](#)

Other multivariate normal functions: [sim_cat\(\)](#), [sim_covar\(\)](#)

Examples

```
library(dplyr)
sim_cat(n_obs = 30, n_groups = 3) %>%
  group_by(group) %>%
  sim_discr(n_vars = 5, var = 1, cov = 0.5, group_means = c(-1, 0, 1), name = "descr")
```

| | |
|-------------|--------------------------------|
| sim_missing | <i>Simulate missing values</i> |
|-------------|--------------------------------|

Description

Takes a data frame and randomly replaces a user-supplied proportion of values with 'NA'.

Usage

```
sim_missing(.data, prop, seed = NA)
```

Arguments

| | |
|-------|--|
| .data | A dataframe. |
| prop | Proportion of values to be set to 'NA'. |
| seed | An optional seed for random number generation. If 'NA' (default) a random seed will be used. |

Value

a dataframe with NAs

Examples

```
library(dplyr)
df <- sim_cat(n_obs = 10, n_groups = 2) %>%
  sim_covar(n_vars = 10, var = 1, cov = 0.5) %>%
  sim_missing(0.05)
```

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