

Package: dmutate (via r-universe)

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Type Package

Title Mutate Data Frames with Random Variates

Version 0.1.2.9002

Imports dplyr (>= 0.7.4), MASS

Depends methods

Suggests testthat

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Description Work within the 'dplyr' workflow to add random variates to your data frame. Variates can be added at any level of an existing column. Also, bounds can be specified for simulated variates.

URL <https://github.com/kylebmetrum/dmutate>

BugReports <https://github.com/kylebmetrum/dmutate/issues>

License GPL (>= 2)

LazyData TRUE

RoxygenNote 6.0.1

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

RemoteUrl <https://github.com/kylebaron/dmutate>

RemoteRef HEAD

RemoteSha 89927b748158a9f3b3c06593315325d387f1f57d

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as_idata	<i>Create individual data frame from a covset object</i>
----------	--

Description

Create individual data frame from a covset object

Usage

```
as_idata(.covset, .n)
```

Arguments

.covset	a covset object
.n	number of IDs to simulate

Examples

```
cov1 <- covset(Y ~ rbinomial(0.2), Z ~ rnorm(2,2))  
as_idata(cov1, 10)
```

build_covform	<i>Build a object or formula to use with covset.</i>
---------------	--

Description

build_covform formulates then parses a formula that can be used in a covset. build_covobj just assembles the object directly.

Usage

```
build_covform(var, dist, args, lower = NULL, upper = NULL, by = NULL,  
  envir = parent.frame())  
  
build_covobj(var, dist, args, upper = NULL, lower = NULL, by = NULL,  
  envir = parent.frame())
```

Arguments

var	variable name, character
dist	distribution function name
args	character vector of arguments for dist
lower	lower limits for var
upper	upper limits for var
by	grouping variable
envir	environment for resolving symbols in expressions

Details

When length of var is greater than one, both lower and upper must be named vectors when specification is made. However, it is acceptable to specify nothing or to use unnamed limits when the length of var is 1.

Examples

```
build_covform("WT", "rnorm", c("mu = 80", "sd = 40"), lower = 40, upper = 140)
build_covform("WT", "rnorm", "80,40", lower = 40, upper = 140)

build_covobj("WT", "rnorm", "80,40", lower = 40, upper = 140)
```

 covset

Covobj and covset objects.

Description

Covobj and covset objects.
 Create a set of covariates.

Usage

```
new_covobj(x, envir = parent.frame(), ...)

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

## S4 method for signature 'covobj'
as.list(x, ...)

## S4 method for signature 'covset'
as.list(x, ...)

## S3 method for class 'covset'
```

```
print(x, ...)  
covset(..., envir = parent.frame())  
rvset(...)  
as.covset(x)
```

Arguments

x	a formula; may be quoted
envir	for formulae
...	formulae to use for the covset

Details

rvset is an alias for covset.

Examples

```
obj <- new_covobj(Y[0,80] ~ rnorm(20,50))  
  
obj  
  
as.list(obj)  
  
a <- Y ~ runif(0,1)  
b <- Z ~ rbeta(1,1)  
  
set <- covset(a,b)  
  
set  
  
as.list(set)
```

dmutate

mutate a data frame, adding random variables.

Description

mutate a data frame, adding random variables.
Apply formulae to a data frame

Usage

```
dmutate(data, ...)
```

Arguments

data a data frame
 ... formulae and other arguments for `mutate_random`

Examples

```
idata <- dplyr::data_frame(ID = 1:10)

dmutate(idata, y ~ rbinomial(0.5), wt ~ rnorm(mu,sd),
        envir = list(mu = 50, sd = 20))
```

mutate_random	<i>Add random variates to a data frame.</i>
---------------	---

Description

Add random variates to a data frame.

Usage

```
mutate_random(data, input, ...)

## S4 method for signature 'data.frame,formula'
mutate_random(data, input, ...)

## S4 method for signature 'data.frame,character'
mutate_random(data, input,
              envir = parent.frame(), ...)

## S4 method for signature 'data.frame,list'
mutate_random(data, input, ...)

## S4 method for signature 'data.frame,covset'
mutate_random(data, input, ...)

## S4 method for signature 'data.frame,covobj'
mutate_random(data, input,
              envir = parent.frame(), ...)
```

Arguments

data the data.frame to mutate
 input an unquoted R formula; see details
 ... additional inputs
 envir environment for object lookup

Examples

```
data <- data.frame(ID=1:10, GROUP = sample(c(1,2,3),10,replace=TRUE))

mutate_random(data, AGE[40,90] ~ rnorm(55,50))
mutate_random(data, RE ~ rbeta(1,1) | GROUP)

e <- list(lower=40,upper=140,mu=100,sd=100)

egfr <- covset(EGFR[lower,upper] ~ rnorm(mu,sd))

mutate_random(data,egfr,envir=e)
```

rbinomial

Simulate from binomial distribution.

Description

Wrapper for [rbinom](#) with trial size of 1.

Usage

```
rbinomial(n, p, ...)
```

```
rbern(n, p, ...)
```

Arguments

n	number of variates
p	probability of success
...	passed along as appropriate

Details

The size of each trial is always 1.

`rmvnorm`*Simulate from multivariate normal distribution.*

Description

Simulate from multivariate normal distribution.

Usage

```
rmvnorm(n, mu, Sigma)
```

```
r1mvnorm(n, ...)
```

```
rmassnorm(n, ...)
```

```
r1massnorm(n, ...)
```

Arguments

<code>n</code>	number of variates
<code>mu</code>	vector of means
<code>Sigma</code>	variance-covariance matrix with number of columns equal to length of <code>mu</code>
<code>...</code>	arguments passed to <code>rmvnorm</code>

Details

`r1mvnorm` is a multivariate log normal.

`rmassnorm` and `r1massnorm` simulate the multivariate normal using the MASS package.

Value

Returns a matrix of variates with number of rows equal to `n` and number of columns equal to length of `mu`.

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