

# Package ‘dmutate’

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**Type** Package

**Title** Mutate Data Frames with Random Variates

**Version** 0.1.3

**Imports** dplyr (>= 0.7.4), MASS

**Depends** methods

**Suggests** testthat

**Maintainer** Kyle T Baron <kylebtwin@imap.cc>

**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/kylebaron/dmutate>

**BugReports** <https://github.com/kylebaron/dmutate/issues>

**License** GPL (>= 2)

**RoxygenNote** 7.1.1

**Encoding** UTF-8

**NeedsCompilation** no

**Author** Kyle T Baron [aut, cre, cph]

**Repository** CRAN

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as_idata	<i>Create individual data frame from a covset object</i>
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**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>
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**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)
```

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dmutate	<i>mutate a data frame, adding random variables.</i>
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**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 <a href="#">mutate_random</a>

**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.

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rmvnorm	<i>Simulate from multivariate normal distribution.</i>
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**Description**

Simulate from multivariate normal distribution.

**Usage**

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

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

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

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

**Arguments**

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

**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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