

# Package ‘dmutate’

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**Type** Package

**Title** Mutate Data Frames with Random Variates

**Version** 0.2.0

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

**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/kylebaron/dmutate>

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

**License** GPL (>= 2)

**Encoding** UTF-8

**Language** en-US

**RoxygenNote** 7.3.3

**NeedsCompilation** no

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**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.

**Value**

An idata set data frame.

**Examples**

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

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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

<code>var</code>	variable name, character.
<code>dist</code>	distribution function name.
<code>args</code>	character vector of arguments for <code>dist</code> .
<code>lower</code>	lower limits for <code>var</code> .
<code>upper</code>	upper limits for <code>var</code> .
<code>by</code>	grouping variable.
<code>envir</code>	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.

### Value

A `covobj` object.

### 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*

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**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**

<code>x</code>	a formula; may be quoted.
<code>envir</code>	for formulae.
<code>...</code>	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	<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 `mutate_random()`.

### Examples

```
idata <- 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>
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---

### 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 `stats::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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