

# Package ‘condMVNorm’

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**Title** Conditional Multivariate Normal Distribution

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**Description** Computes conditional multivariate normal probabilities, random deviates and densities.

**Imports** stats

**Depends** R(>= 1.9.0), mvtnorm

**License** GPL-2

**Author** Ravi Varadhan [aut, cre]

**Maintainer** Ravi Varadhan <ravi.varadhan@jhu.edu>

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cmvnorm *Conditional Multivariate Normal Density and Random Deviates*

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## Description

These functions provide the density function and a random number generator for the conditional multivariate normal distribution, [Y given X], where  $Z = (X, Y)$  is the fully-joint multivariate normal distribution with mean equal to mean and covariance matrix `sigma`.

**Usage**

```
dcmvnorm(x, mean, sigma, dependent.ind, given.ind,
X.given, check.sigma=TRUE, log = FALSE)
rcmvnorm(n, mean, sigma, dependent.ind, given.ind,
X.given, check.sigma=TRUE,
method=c("eigen", "svd", "chol"))
```

**Arguments**

x	vector or matrix of quantiles of Y. If x is a matrix, each row is taken to be a quantile.
n	number of random deviates.
mean	mean vector, which must be specified.
sigma	a symmetric, positive-definite matrix of dimension n x n, which must be specified.
dependent.ind	a vector of integers denoting the indices of dependent variable Y.
given.ind	a vector of integers denoting the indices of conditioning variable X.
X.given	a vector of reals denoting the conditioning value of X. When both given.ind and X.given are missing, the distribution of Y becomes Z[dependent.ind]
check.sigma	logical; if TRUE, the variance-covariance matrix is checked for appropriateness (symmetry, positive-definiteness). This could be set to FALSE if the user knows it is appropriate.
log	logical; if TRUE, densities d are given as log(d).
method	string specifying the matrix decomposition used to determine the matrix root of sigma. Possible methods are eigenvalue decomposition ("eigen", default), singular value decomposition ("svd"), and Cholesky decomposition ("chol"). The Cholesky is typically fastest, not by much though.

**See Also**

[pcmvnorm](#), [pmvnorm](#), [dmvnorm](#), [qmvnorm](#)

**Examples**

```
# 10-dimensional multivariate normal distribution
n <- 10
A <- matrix(rnorm(n^2), n, n)
A <- A %*% t(A)

# density of Z[c(2,5)] given Z[c(1,4,7,9)]=c(1,1,0,-1)
dcmvnorm(x=c(1.2,-1), mean=rep(1,n), sigma=A,
dependent.ind=c(2,5), given.ind=c(1,4,7,9),
X.given=c(1,1,0,-1))

dcmvnorm(x=-1, mean=rep(1,n), sigma=A, dep=3, given=c(1,4,7,9,10), X=c(1,1,0,0,-1))

dcmvnorm(x=c(1.2,-1), mean=rep(1,n), sigma=A, dep=c(2,5))
```

```
# gives an error since `x' and `dep' are incompatibe
# dcmvnorm(x=-1, mean=rep(1,n), sigma=A, dep=c(2,3),
# given=c(1,4,7,9,10), X=c(1,1,0,0,-1))

rcmvnorm(n=10, mean=rep(1,n), sigma=A, dep=c(2,5),
given=c(1,4,7,9,10), X=c(1,1,0,0,-1),
method="eigen")

rcmvnorm(n=10, mean=rep(1,n), sigma=A, dep=3,
given=c(1,4,7,9,10), X=c(1,1,0,0,-1),
method="chol")
```

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condMVN

*Conditional Mean and Variance of Multivariate Normal Distribution*


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### Description

These functions provide the conditional mean and variance-covariance matrix of [Y given X], where  $Z = (X, Y)$  is the fully-joint multivariate normal distribution with mean equal to mean and covariance matrix sigma.

### Usage

```
condMVN(mean, sigma, dependent.ind, given.ind, X.given, check.sigma=TRUE)
```

### Arguments

mean	mean vector, which must be specified.
sigma	a symmetric, positive-definite matrix of dimension $n \times n$ , which must be specified.
dependent.ind	a vector of integers denoting the indices of dependent variable Y.
given.ind	a vector of integers denoting the indices of conditioning variable X.
X.given	a vector of reals denoting the conditioning value of X. When both given.ind and X.given are missing, the distribution of Y becomes $Z[\text{dependent.ind}]$
check.sigma	logical; if TRUE, the variance-covariance matrix is checked for appropriateness (symmetry, positive-definiteness). This could be set to FALSE if the user knows it is appropriate.

### See Also

[dcmvnorm](#), [pcmvnorm](#), [pmvnorm](#), [dmvnorm](#), [qmvnorm](#)

**Examples**

```
# 10-dimensional multivariate normal distribution
n <- 10
A <- matrix(rnorm(n^2), n, n)
A <- A %*% t(A)

condMVN(mean=rep(1,n), sigma=A, dependent=c(2,3,5), given=c(1,4,7,9),X.given=c(1,1,0,-1))

condMVN(mean=rep(1,n), sigma=A, dep=3, given=c(1,4,7,9), X=c(1,1,0,-1))
```

pcmvnorm

*Conditional Multivariate Normal Distribution***Description**

Computes the distribution function of the conditional multivariate normal, [Y given X], where  $Z = (X, Y)$  is the fully-joint multivariate normal distribution with mean equal to mean and covariance matrix sigma.

**Usage**

```
pcmvnorm(lower=-Inf, upper=Inf, mean, sigma,
  dependent.ind, given.ind, X.given,
  check.sigma=TRUE, algorithm = GenzBretz(), ...)
```

**Arguments**

lower	the vector of lower limits of length n.
upper	the vector of upper limits of length n.
mean	the mean vector of length n.
sigma	a symmetric, positive-definite matrix, of dimension n x n, which must be specified.
dependent.ind	a vector of integers denoting the indices of the dependent variable Y.
given.ind	a vector of integers denoting the indices of the conditioning variable X.
X.given	a vector of reals denoting the conditioning value of X. When both given.ind and X.given are missing, the distribution of Y becomes $Z[\text{dependent.ind}]$
check.sigma	logical; if TRUE, the variance-covariance matrix is checked for appropriateness (symmetry, positive-definiteness). This could be set to FALSE if the user knows it is appropriate.
algorithm	an object of class <a href="#">GenzBretz</a> , <a href="#">Miwa</a> or <a href="#">TVPACK</a> specifying both the algorithm to be used as well as the associated hyper parameters.
...	additional parameters (currently given to GenzBretz for backward compatibility issues).

**Details**

This program involves the computation of multivariate normal probabilities with arbitrary correlation matrices.

**Value**

The evaluated distribution function is returned with attributes

error	estimated absolute error and
msg	status messages.

**See Also**

[dcmvnorm](#), [rcmvnorm](#), [pmvnorm](#).

**Examples**

```
n <- 10
A <- matrix(rnorm(n^2), n, n)
A <- A %*% t(A)
```

```
pcmvnorm(lower=-Inf, upper=1, mean=rep(1,n), sigma=A, dependent.ind=3, given.ind=c(1,4,7,9,10),
X.given=c(1,1,0,0,-1))
```

```
pcmvnorm(lower=-Inf, upper=c(1,2), mean=rep(1,n),
sigma=A, dep=c(2,5), given=c(1,4,7,9,10),
X=c(1,1,0,0,-1))
```

```
pcmvnorm(lower=-Inf, upper=c(1,2), mean=rep(1,n), sigma=A,
dep=c(2,5))
```

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