

# Package ‘MVTests’

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**Title** Multivariate Hypothesis Tests

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**Description** Multivariate hypothesis tests and the confidence intervals. It can be used to test the hypothesis about mean vector or vectors (one-sample, two independent samples, paired samples), covariance matrix (one or more matrices), and the correlation matrix. Moreover, it can be used for robust Hotelling  $T^2$  test at one sample case in high dimensional data. For this package, we have benefited from the studies Rencher (2003), Nel and Merwe (1986) <[DOI:10.1080/03610928608829342](https://doi.org/10.1080/03610928608829342)>, Tatlidil (1996), Tsagris (2014), Vilasenor Alva and Estrada (2009) <[DOI:10.1080/03610920802474465](https://doi.org/10.1080/03610920802474465)>.

**License** GPL-2

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Bcov

*Bartlett's Test for One Sample Covariance Matrix***Description**

Bcov function tests whether the covariance matrix is equal to a given matrix or not.

**Usage**

```
Bcov(data, Sigma)
```

**Arguments**

data	a data frame.
Sigma	The covariance matrix in NULL hypothesis.

**Details**

This function computes Bartlett's test statistic for the covariance matrix of one sample.

**Value**

a list with 3 elements:

ChiSquare	The value of Test Statistic
df	The Chi-Square statistic's degree of freedom
p.value	p value

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.

**Examples**

```
data(iris)
S<-matrix(c(5.71,-0.8,-0.6,-0.5,-0.8,4.09,-0.74,-0.54,-0.6,
           -0.74,7.38,-0.18,-0.5,-0.54,-0.18,8.33),ncol=4,nrow=4)
result <- Bcov(data=iris[,1:4],Sigma=S)
summary(result)
```

BoxM

*Box's M Test***Description**

BoxM function tests whether the covariance matrices of independent samples are equal or not.

**Usage**

```
BoxM(data, group)
```

**Arguments**

data	a data frame.
group	grouping vector.

**Details**

This function computes Box-M test statistic for the covariance matrices of independent samples. The hypotheses are defined as H0:The Covariance matrices are homogeneous and H1:The Covariance matrices are not homogeneous

**Value**

a list with 3 elements:

ChiSquare	The value of Test Statistic
df	The Chi-Square statistic's degree of freedom
p.value	p value

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.

**Examples**

```
data(iris)
results <- BoxM(data=iris[,1:4],group=iris[,5])
summary(results)
```

Bsper

*Bartlett's Sphericity Test***Description**

Bsper function tests whether a correlation matrix is equal to the identity matrix or not.

**Usage**

```
Bsper(data)
```

**Arguments**

data                    a data frame.

**Details**

This function computes Bartlett's test statistic for Sphericity Test. The hypotheses are  $H_0: R$  is equal to  $I$  and  $H_1: R$  is not equal to  $I$ .

**Value**

a list with 4 elements:

ChiSquare	The value of Test Statistic
df	The Chi-Square statistic's degree of freedom
p.value	p value
R	Correlation matrix

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Tatlidil, H. (1996). Uygulamali Cok Degiskenli Istatistiksel Yontemler. Cem Web.

**Examples**

```
data(iris)
results <- Bsper(data=iris[,1:4])
summary(results)
```

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Coated

*Coated*

---

**Description**

The data set is given in Table 5.3 in Rencher (2003). The data set consists of 2 variables (Depth and Number), 2 treatments and 15 observations. The first column of the data is Location numbers.

**Usage**

Coated

**Format**

A data frame with 15 rows and 5 columns. The columns are as follows:

**Location** The location numbers of observations.

**Coating1.Depth1** The Depth values in the first treatment

**Coating1.Number1** The Number values in the first treatment

**Coating2.Depth2** The Depth values in the second treatment

**Coating2.Number2** The Number values in the second treatment

**Source**

The data set is used in the book entitled Methods of Multivariate Analysis (Rencher,2003).

**References**

Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.

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iris

*Iris Data*

---

**Description**

The Iris dataset is consists of 4 variables, 3 groups and 150 observations. The last column of the data is Iris species.

**Usage**

iris

**Format**

A data frame with 150 rows and 5 columns. The columns are as follows:

**Sepal.Length** The Sepal length values of iris flowers

**Sepal.Width** The Sepal width values of iris flowers

**Petal.Length** The Petal length values of iris flowers

**Petal.Width** The Petal width values of iris flowers

**Species** The species of iris flowers

**Source**

<https://archive.ics.uci.edu/ml/datasets/Iris>

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Mpaired

*Multivariate Paired Test*

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**Description**

Mpaired function computes the value of test statistic based on Hotelling T Square approach in multivariate paired data sets.

**Usage**

```
Mpaired(T1, T2)
```

**Arguments**

T1	The first treatment data.
T2	The second treatment data.

**Details**

This function computes one sample Hotelling  $T^2$  statistics for paired data sets.

**Value**

a list with 7 elements:

HT2	The value of Hotelling $T^2$ Test Statistic
F	The value of F Statistic
df	The F statistic's degree of freedom
p.value	p value
Descriptive1	The descriptive statistics of the first treatment
Descriptive2	The descriptive statistics of the second treatment
Descriptive.Difference	The descriptive statistics of the differences

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.

**Examples**

```
data(Coated)
X<-Coated[,2:3]; Y<-Coated[,4:5]
result <- Mpaired(T1=X,T2=Y)
summary(result)
```

---

 OneSampleHT2

*One Sample Hotelling T<sup>2</sup> Test*


---

**Description**

OneSampleHT2 computes one sample Hotelling T<sup>2</sup> statistics and gives confidence intervals

**Usage**

```
OneSampleHT2(data, mu0, alpha = 0.05)
```

**Arguments**

data	a data frame.
mu0	mean vector that is used to test whether population mean parameter is equal to it.
alpha	Significance Level that will be used for confidence intervals. default alpha=0.05.

**Details**

This function computes one sample Hotelling T<sup>2</sup> statistics that is used to test whether population mean vector is equal to a vector given by a user. When H<sub>0</sub> is rejected, this function computes confidence intervals for all variables.

**Value**

a list with 7 elements:

HT2	The value of Hotelling T <sup>2</sup> Test Statistic
F	The value of F Statistic
df	The F statistic's degree of freedom
p.value	p value

CI	The lower and upper limits of confidence intervals obtained for all variables
alpha	The alpha value using in confidence intervals
Descriptive	Descriptive Statistics

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

- Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.  
 Tatlidil, H. (1996). Uygulamali Cok Degiskenli Istatistiksel Yontemler. Cem Web.

**Examples**

```
data(iris)

mean0<-c(6,3,1,0.25)
result <- OneSampleHT2(data=iris[1:50,-5],mu0=mean0,alpha=0.05)
summary(result)
```

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RHT2

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*Robust Hotelling T<sup>2</sup> Test for One Sample in High Dimensional Data*


---

**Description**

Robust Hotelling T<sup>2</sup> Test for One Sample in high Dimensional Data

**Usage**

```
RHT2(data, mu0, alpha = 0.75, d, q)
```

**Arguments**

data	the data. It must be matrix or data.frame.
mu0	the mean vector which will be used to test the null hypothesis.
alpha	numeric parameter controlling the size of the subsets over which the determinant is minimized. Allowed values are between 0.5 and 1 and the default is 0.75.
d	the constant in Equation (11) in the study by Bulut (2021).
q	the second degree of freedom value of the approximate F distribution in Equation (11) in the study by Bulut (2021).

**Details**

RHT2 function performs a robust Hotelling T<sup>2</sup> test in high dimensional test based on the minimum regularized covariance determinant estimators. This function needs the q and d values. These values can be obtained simRHT2 function. For more detailed information, you can see the study by Bulut (2021).



**Value**

a list with 3 elements:

T2	The Robust Hotelling $T^2$ value in high dimensional data
Fval	The F value based on T2
pval	The p value based on the approximate F distribution

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Bulut, H (2021). A robust Hotelling test statistic for one sample case in high dimensional data, Communication in Statistics: Theory and Methods.

**Examples**

```
library(rrcov)
data(octane)
mu.clean<-colMeans(octane[-c(25,26,36,37,38,39),])

RHT2(data=octane,mu0=mu.clean,alpha=0.84,d=1396.59,q=1132.99)
```

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RobCat	<i>Robust CAT Algorithm</i>
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**Description**

RobCat computes p value based on robust CAT algorithm to compare two means vectors under multivariate Behrens-Fisher problem.

**Usage**

```
RobCat(X, Y, M = 1000, alpha = 0.75)
```

**Arguments**

X	a matrix or data frame for first group.
Y	a matrix or data frame for second group.
M	iteration number and the default is 1000.
alpha	numeric parameter controlling the size of the subsets over which the determinant is minimized; roughly $\alpha \cdot n$ , observations are used for computing the determinant. Allowed values are between 0.5 and 1 and the default is 0.75.

**Details**

This function computes p value based on robust CAT algorithm to compare two means vectors under multivariate Behrens-Fisher problem. When p value < 0.05, it means the difference of two mean vectors is significant statistically.

**Value**

a list with 2 elements:

Cstat	Calculated value of test statistic
pval	The p value

**Author(s)**

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**Examples**

```
data(iris)
RobCat(X=iris[1:20,-5],Y=iris[81:100,-5])
```

---

simRHT2

*Monte Carlo Simulation to obtain d and q constants for RHT2 function*

---

**Description**

Monte Carlo Simulation to obtain d and q constants for RHT2 function

**Usage**

```
simRHT2(n, p, nrep = 500)
```

**Arguments**

n	the sample size
p	the number of variables
nrep	the number of iteration. The default value is 500.

**Details**

simRHT2 function computes d and q constants to construct an approximate F distribution of robust Hotelling  $T^2$  statistic in high dimensional data. These constants are used in RHT2 function. For more detailed information, you can see the study by Bulut (2021).

**Value**

a list with 2 elements:

q	The q value
d	The d value

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**References**

Bulut, H (2021). A robust Hotelling test statistic for one sample case in highdimensional data, *Communication in Statistics: Theory and Methods*.

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summary.MVTests	<i>Summarizing Results in MVTests Package</i>
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**Description**

summary.MVTests function summarizes of results of functions in this package.

**Usage**

```
## S3 method for class 'MVTests'  
summary(object, ...)
```

**Arguments**

object	an object of class MVTests.
...	additional parameters.

**Details**

This function prints a summary of the results of multivariate hypothesis tests in the MVTests package.

**Value**

the input object is returned silently.

**Author(s)**

Hasan BULUT <hasan.bulut@omu.edu.tr>

**Examples**

```
# One Sample Hotelling T Square Test
data(iris)
X<-iris[1:50,1:4]
mean0<-c(6,3,1,0.25)
result.onesample <- OneSampleHT2(data=X,mu0=mean0,alpha=0.05)
summary(result.onesample)

#Two Independent Sample Hotelling T Square Test
data(iris)
G<-c(rep(1,50),rep(2,50))
result.twosamples <- TwoSamplesHT2(data=iris[1:100,1:4],group=G,alpha=0.05)
summary(result.twosamples)

#Box's M Test
data(iris)
result.BoxM <- BoxM(data=iris[,1:4],group=iris[,5])
summary(result.BoxM)

#Barlett's Test of Sphericity
data(iris)
result.Bsper <- Bsper(data=iris[,1:4])
summary(result.Bsper)

#Bartlett's Test for One Sample Covariance Matrix
data(iris)
S<-matrix(c(5.71,-0.8,-0.6,-0.5,-0.8,4.09,-0.74,-0.54,-0.6,-0.74,
            7.38,-0.18,-0.5,-0.54,-0.18,8.33),ncol=4,nrow=4)
result.bcov<- Bcov(data=iris[,1:4],Sigma=S)
summary(result.bcov)
```

---

TwoSamplesHT2

*Two Independent Samples Hotelling T<sup>2</sup> Test*


---

**Description**

TwoSamplesHT2 function computes Hotelling T<sup>2</sup> statistic for two independent samples and gives confidence intervals.

**Usage**

```
TwoSamplesHT2(data, group, alpha = 0.05, Homogeneity = TRUE)
```

**Arguments**

data	a data frame.
group	a group vector consisting of 1 and 2 values.
alpha	Significance Level that will be used for confidence intervals. default=0.05

Homogeneity a logical argument. If sample covariance matrices are homogeneity, then Homogeneity=TRUE. Otherwise Homogeneity=FALSE The homogeneity of covariance matrices can be investigated with BoxM function.

### Details

This function computes two independent samples Hotelling  $T^2$  statistics that is used to test whether two population mean vectors are equal to each other. When  $H_0$  is rejected, this function computes confidence intervals for all variables to determine variable(s) affecting on rejection decision. Moreover, when covariance matrices are not homogeneity, the approach proposed by D. G. Nel and V. D. Merwe (1986) is used.

### Value

a list with 8 elements:

HT2	The value of Hotelling $T^2$ Test Statistic
F	The value of F Statistic
df	The F statistic's degree of freedom
p.value	p value
CI	The lower and upper limits of confidence intervals obtained for all variables
alpha	The alpha value using in confidence intervals
Descriptive1	Descriptive Statistics for the first group
Descriptive2	Descriptive Statistics for the second group

### Author(s)

Hasan BULUT <hasan.bulut@omu.edu.tr>

### References

Rencher, A. C. (2003). Methods of multivariate analysis (Vol. 492). John Wiley & Sons.  
 Tatlidil, H. (1996). Uygulamali Cok Degiskenli Istatistiksel Yontemler. Cem Web.  
 D.G. Nel & C.A. Van Der Merwe (1986) A solution to the multivariate behrens fisher problem, Communications in Statistics:Theory and Methods, 15:12, 3719-3735

### Examples

```
data(iris)
G<-c(rep(1,50),rep(2,50))
# When covariances matrices are homogeneity
results1 <- TwoSamplesHT2(data=iris[1:100,1:4],group=G,alpha=0.05)
summary(results1)
# When covariances matrices are not homogeneity
results2 <- TwoSamplesHT2(data=iris[1:100,1:4],group=G,Homogeneity=FALSE)
summary(results2)
```

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