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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	<i>Box's M Test</i>
------	---------------------

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)
```

`ccc`*Concordance Correlation Coefficient*

Description

Classical Concordance Correlation Coefficient

Usage

```
ccc(x, y)
```

Arguments

<code>x</code>	the vector which contains the first variable values
<code>y</code>	the vector which contains the second variable values

Details

ccc function calculates directly classical concordance correlation coefficient.

Value

a list with 1 elements:

<code>coef</code>	The value of concordance correlation coefficient
-------------------	--

Author(s)

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

References

Bulut, H (2025). A Robust Concordance Correlation Coefficient. (Unpublished)

Lin, L. I. "A Concordance Correlation-Coefficient to Evaluate Reproducibility." Biometrics 45, no. 1 (1989): 255-68.

Examples

```
x<-rnorm(50)
y<-2+3*x+rnorm(50,mean = 3)
ccc(x,y)
```

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.

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>

Mhg

Pair-Wise comparison between hth and gth sample

Description

Pair-Wise comparison of covariance matrices between hth and gth sample

Usage

Mhg(Sh, Sg, S, nh, ng, n)

Arguments

Sh	the robust covariance matrix of the hth sample
Sg	the robust covariance matrix of the gth sample
S	the robust pooled covariance matrix.
nh	the sample size of the hth sample
ng	the sample size of the gth sample
n	the sample size of the full data

Details

Mhg function computes proposed Mgh values as defined in the paper.

Value

a list with 1 elements:

Mhg	Mgh value
-----	-----------

Author(s)

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

References

Bulut, H (2024). A robust permutational test to compare covariance matrices in high dimensional data. (Unpublished)

Examples

```
if (requireNamespace("rrcov", quietly=TRUE)) {
  x1<-rmvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = diag(20))
  x2<-rmvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = 2*diag(20))
  x3<-rmvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = 3*diag(20))
  data<-rbind(x1,x2,x3)
  group_label<-c(rep(1,10),rep(2,10),rep(3,10))
  n <- nrow(data)
  p <- ncol(data)
  nk <- table(group_label)
  g <- length(nk)
  Levels <- unique(group_label)
  Si.matrices<-lapply(1:g, function(i) rrcov::CovMrcd(data[(group_label==Levels[i]),],
    alpha=0.9)@cov)
  Spool <- Reduce("+", Map("*", nk, Si.matrices)) / n
  #for the first and second groups
  Mhg(Sh = Si.matrices[[1]], Sg = Si.matrices[[2]],S = Spool, nh = nk[1], ng = nk[2], n = n)}
```

Mpaired

Multivariate Paired Test

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	<i>One Sample Hotelling T^2 Test</i>
--------------	---

Description

OneSampleHT2 computes one sample Hotelling T^2 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^2 statistics that is used to test whether population mean vector is equal to a vector given by a user. When H_0 is rejected, this function computes confidence intervals for all variables.

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
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)
```

rccc

Robust Concordance Correlation Coefficient (rCCC)

Description

Computes a robust concordance correlation coefficient using Minimum Covariance Determinant (MCD) estimates.

Usage

```
rccc(x, y, alpha = 0.75)
```

Arguments

x	Numeric vector; first variable.
y	Numeric vector; second variable.
alpha	Numeric in (0.5, 1]; MCD subset size proportion. Default 0.75.

Details

The rCCC replaces means and (co)variances in Lin's CCC with their MCD counterparts: $\rho_c = \frac{2\sigma_{xy}}{\sigma_x^2 + \sigma_y^2 + (\mu_x - \mu_y)^2}$.

Value

A list with one element:

coef	Robust concordance correlation coefficient
------	--

Author(s)

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

References

Bulut, H. (2025). A Robust Concordance Correlation Coefficient. (Unpublished)

Examples

```
if (requireNamespace("robustbase", quietly = TRUE)) {
  set.seed(1)
  x <- rnorm(50)
  y <- 2 + 3*x + rnorm(50, mean = 3)
  rccc(x, y)
}
```

Description

Robust Hotelling T² Test for One Sample in high Dimensional Data

Usage

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

Arguments

<code>data</code>	the data. It must be matrix or data.frame.
<code>mu0</code>	the mean vector which will be used to test the null hypothesis.
<code>alpha</code>	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.
<code>d</code>	the constant in Equation (11) in the study by Bulut (2021).
<code>q</code>	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^2 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:

<code>T2</code>	The Robust Hotelling T^2 value in high dimensional data
<code>Fval</code>	The F value based on <code>T2</code>
<code>pval</code>	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

```
if (requireNamespace("rrcov", quietly = TRUE)) {
  utils::data("octane", package = "rrcov")
  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)}
```

Rob_CovTest	<i>Robust Test for Covariance Matrices</i>
-------------	--

Description

Robust Test for Covariance Matrices in High Dimensional Data

Usage

Rob_CovTest(x, group, alpha = 0.75)

Arguments

- x the data matrix
- group the grouping vector. It must be factor.
- 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.

Details

Rob_CovTest function computes the calculated value of the test statistic for covariance matrices of two or more independent samples in high dimensional data based on the minimum regularized covariance determinant estimators.

Value

- a list with 1 elements:
 - TM The calculated value of test statistics based on raw data

Author(s)

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

References

Bulut, H (2024). A robust permutational test to compare covariance matrices in high dimensional data. (Unpublished)

Examples

```
if (requireNamespace("rrcov", quietly=TRUE)) {
  x1<-rmvtnorm::rmvnorm(n = 8,mean = rep(0,10),sigma = diag(10))
  x2<-rmvtnorm::rmvnorm(n = 8,mean = rep(0,10),sigma = 2*diag(10))
  x3<-rmvtnorm::rmvnorm(n = 8,mean = rep(0,10),sigma = 3*diag(10))
  data<-rbind(x1,x2,x3)
  group_label<-c(rep(1,8),rep(2,8),rep(3,8))
  Rob_CovTest(x=data, group=group_label)}
```

RobCat*Robust CAT Algorithm*

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 $p \text{ value} < 0.05$, 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)

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

Examples

```
data(iris)
if (requireNamespace("robustbase", quietly=TRUE)) {
  RobCat(X=iris[1:20,-5],Y=iris[81:100,-5])
}
```

RobPer_CovTest	<i>Robust Permutation Test for Covariance Matrices</i>
----------------	--

Description

Robust Permutation Test for Covariance Matrices in High Dimensional Data

Usage

RobPer_CovTest(x, group, N = 100, alpha = 0.75)

Arguments

- x the data matrix
- group the grouping vector. It must be factor.
- N the permutation number and the default value is 100.
- 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.

Details

RobPer_CovTest function calculates directly p-value based on the calculated value of test statistics and the permutational distribution of test statistics for covariance matrices of two or more independent samples in high dimensional data based on the minimum regularized covariance determinant estimators.

Value

a list with 3 elements:

- pval p-value of the robust permutation test process
- TM The calculated value of test statistics based on raw data
- Permutations_TM The calculated values of test statistics based on each permutational data

Author(s)

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

References

Bulut, H (2024). A robust permutational test to compare covariance matrices in high dimensional data. (Unpublished)

Examples

```
if (requireNamespace("rrcov", quietly=TRUE)) {
  x1<-mvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = diag(20))
  x2<-mvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = 2*diag(20))
  x3<-mvtnorm::rmvnorm(n = 10,mean = rep(0,20),sigma = 3*diag(20))
  data<-rbind(x1,x2,x3)
  group_label<-c(rep(1,10),rep(2,10),rep(3,10))
  RobPer_CovTest(x=data, group=group_label)}
```

RperT2	<i>Robust Permutation Hotelling T^2 Test in High Dimensional Data</i>
--------	---

Description

Robust Permutation Hotelling T^2 Test for Two Independent Samples in high Dimensional Data

Usage

```
RperT2(X1, X2, alpha = 0.75, N = 100)
```

Arguments

X1	the data matrix for the first group. It must be matrix or data.frame.
X2	the data matrix for the first group. It must be matrix or data.frame.
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.
N	the permutation number

Details

RperT2 function performs a robust permutation Hotelling T^2 test for two independent samples in high dimensional test based on the minimum regularized covariance determinant estimators.

Value

a list with 2 elements:

T2	The calculated value of Robust Hotelling T^2 statistic based on MRCD estimations
p.value	p value obtained from test process

Author(s)

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

References

Bulut et al. (2024). A Robust High-Dimensional Test for Two-Sample Comparisons, Axioms.

Examples

```
if (requireNamespace("rrcov", quietly=TRUE)) {
  x<-rmvnorm::rmvnorm(n=10,sigma=diag(20),mean=rep(0,20))
  y<-rmvnorm::rmvnorm(n=10,sigma=diag(20),mean=rep(1,20))
  RperT2(X1=x,X2=y)$p.value}
```

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, alpha = 0.75)
```

Arguments

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

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.

summary.MVTests*Summarizing Results in MVTests Package*

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)
```

TR2	<i>Robust Hotelling T^2 Test Statistic</i>
-----	---

Description

Robust Hotelling T^2 Test Statistic for Two Independent Samples in high Dimensional Data

Usage

```
TR2(x1, x2, alpha = 0.75)
```

Arguments

x1	the data matrix for the first group. It must be matrix or data.frame.
x2	the data matrix for the first group. It must be matrix or data.frame.
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.

Details

TR2 function calculates the robust Hotelling T^2 test statistic for two independent samples in high dimensional data based on the minimum regularized covariance determinant estimators.

Value

a list with 2 elements:

TR2	The calculated value of Robust Hotelling T^2 statistic based on MRCD estimations
-----	--

Author(s)

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

References

Bulut et al. (2024). A Robust High-Dimensional Test for Two-Sample Comparisons, Axioms

Examples

```
if (requireNamespace("rrcov", quietly=TRUE)) {
  x<-rmvnorm::rmvnorm(n=10,sigma=diag(20),mean=rep(0,20))
  y<-rmvnorm::rmvnorm(n=10,sigma=diag(20),mean=rep(1,20))
  TR2(x1=x,x2=y)}
```

TwoSamplesHT2	<i>Two Independent Samples Hotelling T^2 Test</i>
---------------	---

Description

TwoSamplesHT2 function computes Hotelling T^2 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 H0 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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