Package: pecora (via r-universe)

October 16, 2026

Type Package

Title Permutation Conditional Random Tests

Version 0.1.3

Date 2026-07-14

Description It provides functions to perform permutation conditional random one-sample and two-samples t-tests in a multivariate framework.

License GPL (>= 2)

Imports Rcpp (>= 1.0.3), matrixStats, stats

LinkingTo Rcpp, RcppArmadillo

RoxygenNote 7.1.1

Encoding UTF-8

Language en-US

NeedsCompilation yes

Author Angela Andreella [aut, cre] (Main author, ORCID: <https://orcid.org/0000-0002-1141-3041>)

Maintainer Angela Andreella <angela.andreella@unive.it>

Repository https://cran.r-universe.dev

Date/Publication 2026-07-16 10:30:02 UTC

RemoteUrl https://github.com/cran/pecora

RemoteRef HEAD

RemoteSha 4b5e1239b97d1d0d913bb423c48c6d27ca4672d3

Contents

pecora-package 2
oneSample 2
t2p 3
twoSamples 4

Index **6**

pecora-package *pecora-package*

Description

The library is devoted to permutation-based inferential methods.

The pecora (permutation conditional random) package provides functions to perform the one-sample and two-samples t-tests using permutations/sign-flipping.

The tests comprised are: the one and two samples t-tests.

Author(s)

Angela Andreella.

Maintainer: Angela Andreella. <angela.andreella@unipd.it>

References

For the general framework of univariate and multivariate permutation tests see: Pesarin, F. (2001) Multivariate Permutation Tests with Applications in Biostatistics. Wiley, New York.

Examples

```
X <- matrix(rnorm(100*20), nrow=20)
out <- oneSample(X = X)
pv <- t2p(Test = out, alternative = "two.sided")

X <- matrix(rnorm(100*20), nrow=20)
rownames(X) <- c(rep(0, 10), rep(1,10))
out<- twoSamples(X = X)
pv <- t2p(Test = out, alternative = "two.sided")
```

oneSample *Permutatation-based one sample t-test*

Description

Performs sign-flipped one-sample t-tests.

Usage

```
oneSample(X, B = 1000, seed = 1234, permReturn = TRUE)
```

Arguments

X	data matrix where columns represent the m variables and rows the n observations.
B	numeric value, number of permutations to be performed, including the identity. Default is 1000.
seed	numeric value, specify seed. Default is 1234.
permReturn	logical value, TRUE to return the t-tests permutation distribution. Default is TRUE.

Value

Returns an object matrix:

tv Matrix with dimensions mxB of permuted one-sample t-tests. The first column is the observed one-sample t-tests.

if permReturn = TRUE otherwise returns:

tv Vector of m observed one-sample t-tests

Author(s)

Angela Andreella

Examples

```
X <- matrix(rnorm(100*20), nrow=20)
out <- oneSample(X = X)
```

---

t2p *From t-tests to p-values*

Description

Use permutation distribution of a test statistic to get p-values.

Usage

```
t2p(Test, alternative = "two.sided", rankBased = TRUE, permReturn = TRUE, df = Inf)
```

Arguments

Test	can be a matrix or a vector. In the first case the columns represent the B permutations and rows the m tests statistic. The observed test statistic is in the first column and the permutation distribution in the remaining columns. In the second case, it is a vector of length m of observed tests statistics. If rankBased = TRUE, you must provide the first option (matrix of permuted statistical tests).
alternative	character string referring to the alternative hypothesis ("greater", "lower", or "two.sided"). is "two.sided".

rankBased	logical value, TRUE to compute p-values by permutation distribution. Default @TRUE.
permReturn	logical value, TRUE to return the t-tests and p-values permutation distribution. Default @TRUE.
df	numerical value. Degrees of freedom (> 0, maybe non-integer). Default df = Inf

Value

Returns an object matrix:

pv Matrix with dimensions mxB of permuted one-sample p-values. The first column is the p-values for the observed one-sample t-tests.

if permReturn = TRUE otherwise returns:

pv Vector of m p-values for the observed one-sample t-tests

Author(s)

Angela Andreella

Examples

```
X <- matrix(rnorm(100*20), nrow=20)
out <- oneSample(X = X)
pv <- t2p(Test = out)
```

---

twoSamples *Permutatation-based two sample t-test*

Description

Performs two-sample t-tests by permutations.

Usage

```
twoSamples(X, B = 1000, seed = 1234,
permReturn = TRUE, label = NULL)
```

Arguments

X	data matrix where columns represent the m variables and rows the n observations. The columns' name defines the groups' label.
B	numeric value, number of permutations to be performed, including the identity. Default is 1000.
seed	numeric value, specify seed. Default is 1234.

permReturn	logical value, TRUE to return the t-tests and p-values permutation distribution. Default is TRUE.
label	by default label = NULL. Labels of the observations, if NULL the rows's name are considered. D

Value

Returns a matrix objects:

Test Matrix with dimensions mxB of permuted two-samples t-tests. The first column is the observed one-sample t-tests.

if permReturn = TRUE otherwise returns:

Test Vector of m observed two-samples t-tests

Author(s)

Angela Andreella

Examples

```
X <- matrix(rnorm(100*20), nrow=20)
rownames(X) <- c(rep(0, 10), rep(1,10))
out<- twoSamples(X = X)
```

Index

∗ **package**
pecora-package, 2

oneSample, 2

pecora-package, 2
pecora.package *(*pecora-package*)*, 2
pecorapackage *(*pecora-package*)*, 2

t2p, 3
twoSamples, 4