

Package ‘ChangepointTesting’

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Type Package

Title Change Point Estimation for Clustered Signals

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Description A multiple testing procedure for clustered alternative hypotheses. It is assumed that the p-values under the null hypotheses follow $U(0,1)$ and that the distributions of p-values from the alternative hypotheses are stochastically smaller than $U(0,1)$. By aggregating information, this method is more sensitive to detecting signals of low magnitude than standard methods. Additionally, sporadic small p-values appearing within a null hypotheses sequence are avoided by averaging on the neighboring p-values.

License GPL-2

Depends graphics, methods, stats

NeedsCompilation no

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ChangepointTesting-package

Change Point Estimation for Clustered Signals

Description

A multiple testing procedure for clustered alternative hypotheses. It is assumed that the p-values under the null hypotheses follow $U(0,1)$ and that the distributions of p-values from the alternative hypotheses are stochastically smaller than $U(0,1)$. By aggregating information, this method is more sensitive to detecting signals of low magnitude than standard methods. Additionally, sporadic small p-values appearing within a null hypotheses sequence are avoided by averaging on the neighboring p-values.

Details

| | |
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| Type: | Package |
| Version: | 1.1 |
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Author(s)

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References

Cao, H. and Wu, W. B. (2015) Changepoint estimation: Another look at multiple testing problems. *Biometrika*, 102, 974–980.

changePoint

Change Point Estimation for Clustered Signals

Description

A multiple testing procedure for clustered alternative hypotheses. It is assumed that the p-values under the null hypotheses follow $U(0,1)$ and that the distributions of p-values from the alternative hypotheses are stochastically smaller than $U(0,1)$. By aggregating information, this method is more sensitive to detecting signals of low magnitude than standard methods. Additionally, sporadic small p-values appearing within a null hypotheses sequence are avoided by averaging on the neighboring p-values.

Usage

```
changePoint(pvalues, alpha, km, lm, compare = "BOTH", fdrWindow = 3,
fdrNStep = 300, fdrLambda = 0.1)
```

Arguments

| | |
|-----------|--|
| pvalues | an object of class numeric. A vector of p-values. |
| alpha | an object of class numeric. The significant level for the estimation of the critical value, γ^* . |
| km | an object of class numeric. The size of the window defining the neighborhood in left and right distances. |
| lm | an object of class numeric. The size of the window defining the neighborhood in the long-run variance estimation. |
| compare | one of ("FDRL", "BH", "Both", "None"). In addition to the Cao-Wu method, obtain significance indicators using the FDR_L method (FDRL) (Zhang et al., 2011), the Benjamini-Hochberg method (BH), (Benjamini and Hochberg, 1995), "both" the FDRL and the BH methods, or do not consider alternative methods (none). |
| fdrWindow | an object of class numeric. If FDR_L method requested, the size of the window defining the neighborhood. |
| fdrNStep | an object of class numeric. If FDR_L method requested, the number of threshold values to consider. |
| fdrLambda | and object of class numeric. If FDR_L method requested, the tuning constant. |

Details

The comparison capability is included only for convenience and reproducibility of the original manuscript. The Benjamini-Hochberg and FDR_L methods cannot be accessed outside of the changePoint function.

The following methods retrieve individual results from a changePoint object, x:

BH(x): Retrieves a vector of integer values. An element is 1 if the null hypothesis is rejected by the Benjamini-Hochberg (1995) method.

blocks(x): Retrieves a list, each element of which is a vector of integer values. Each vector contains the indices of an alternative hypothesis block.

CW(x): Retrieves a vector of integer values. An element is 1 if the null hypothesis is rejected by the Cao-Wu change point (2015) method.

changePts(x): Retrieves a vector of integer values. The vector of change points identified by the Cao-Wu (2015) method. If no change points are identified, NULL is returned.

FDRL(x): Retrieves a vector of integer values. Elements are 1 if the null hypothesis is rejected by the FDR_L (Zhang et al. 2011) method.

critical(x): Retrieves the estimated critical value for testing used by the Cao-Wu (2015) method.

numAlt(x): Retrieves the estimated number of alternative hypotheses obtained by the Cao-Wu (2015) method.

`piAlt(x)`: Retrieves the estimated proportion of alternative hypotheses obtained by the Cao-Wu (2015) method.

`plot(x, y, logp, ...)`: Generates plots of $-\log(p)$ vs position or p-value vs position for each alternative hypothesis block obtained by the Cao-Wu (2015) method. `logp` is TRUE/FALSE indicating if $-\log(p)$ /p-values are plotted on the y-axis.

`sigmaSq(x)`: Retrieves the estimated variance used to determine the critical value of the Cao-Wu (2015) method.

Value

Returns an object of class `changePoint`.

Author(s)

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References

Benjamini, Y. and Hochberg, Y. (1995). Controlling the false discovery rate: A practical and powerful approach to multiple testing. *Journal of the Royal Statistical Society: Series B*, 57, 289–300.

Cao, H. and Wu, W. B. (2015) Changepoint estimation: Another look at multiple testing problems. *Biometrika*, 102, 974–980.

Zhang, C., Fan, J., and Yu, T. (2011). Multiple testing via FDRL for large-scale imaging data. *Annals of Statistics*, 39, 613–642.

Examples

```
m <- 5000

T <- c(rep(0.1, 75), rep( 0.8, 75), rep(1.8, 100),
      rep(0.0,2250), rep(-1.5,250), rep(0.0,2250)) +
  rnorm(m, mean=0.0, sd = 1.0)

pv <- 2.0*(1.0-pnorm(abs(T)))

res <- changePoint(pvalues = pv,
                  alpha = 0.05,
                  km = {log(m)}^2,
                  lm = m^{1/4},
                  compare = "Both")

print(changePts(res))

print(head(cbind(BH(res),FDRL(res),CW(res))))
```

| | |
|-------------------|---------------------|
| changePoint-class | Class "changePoint" |
|-------------------|---------------------|

Description

Value object returned by call to `changePoint()`.

Objects from the Class

This object should not be created by users.

Slots

CW: Object of class `numeric` or `NULL`. A vector of 1/0 values; 1 indicates that hypothesis was rejected by the Cao-Wu method.

chgPts: Object of class `numeric` or `NULL`. The vector of change points identified by the Cao-Wu method. If no change points are identified, `NULL`.

pi_alt: Object of class `numeric`. The estimated proportion of alternative hypotheses calculated using the Cao-Wu method.

num_alt: Object of class `numeric`. The estimated number of alternative hypotheses calculated using the Cao-Wu method.

FDR_L: Object of class `numeric` or `NULL`. A vector of 1/0 values; 1 indicates that hypothesis was rejected by the `FDR_L` method.

BH: Object of class `numeric` or `NULL`. A vector of 1/0 values; 1 indicates that hypothesis was rejected by the `FDR_L` method.

gammaStar: Object of class `numeric`. The estimated critical value for testing used by the Cao-Wu method.

sigmaSq: Object of class `numeric`. The estimated variance used to determine the critical value of the Cao-Wu method.

pVals: Object of class `numeric`. The original p-values provided as input.

Methods

BH `signature(x = "changePoint")`: Retrieves a vector of integer values. An elements is 1 if the null hypothesis is rejected by the Benjamini-Hochberg (1995) method.

blocks `signature(x = "changePoint")`: Retrieves a list, each element of which is a vector of integer values. Each vector contains the indices of an alternative hypothesis block.

CW `signature(x = "changePoint")`: Retrieves a vector of integer values. An element is 1 if the null hypothesis is rejected by the Cao-Wu change point (2015) method.

changePts `signature(x = "changePoint")`: Retrieves a vector of integer values. The vector of change points identified by the Cao-Wu (2015) method. If no change points are identified, `NULL` is returned.

FDR_L `signature(x = "changePoint")`: Retrieves a vector of integer values. Elements are 1 if the null hypothesis is rejected by the `FDR_L` (Zhang et al. 2011) method.

critical signature(*x* = "changePoint"): Retrieves the estimated critical value for testing used by the Cao-Wu (2015) method.

numAlt signature(*x* = "changePoint"): Retrieves the estimated number of alternative hypotheses obtained by the Cao-Wu (2015) method.

piAlt signature(*x* = "changePoint"): Retrieves the estimated proportion of alternative hypotheses obtained by the Cao-Wu (2015) method.

plot signature(*x* = "changePoint", *y* = "missing", *logp* = FALSE, ...): Generates x-y plots of $-\log(p)$ vs position or p-value vs position for each alternative hypothesis block obtained by the Cao-Wu (2015) method. *logp* is TRUE/FALSE indicating if $-\log(p)$ /p-values are plotted on the y-axis.

sigmaSq signature(*x* = "changePoint"): Retrieves the estimated variance used to determine the critical value of the Cao-Wu (2015) method.

Author(s)

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References

- Benjamini, Y. and Hochberg, Y. (1995). Controlling the false discovery rate: A practical and powerful approach to multiple testing. *Journal of the Royal Statistical Society: Series B*, 57, 289–300.
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Examples

```
showClass("changePoint")
```

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