

Package: bwd (via r-universe)

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Type Package

Title Backward Procedure for Change-Point Detection

Version 0.1.0

Maintainer Seung Jun Shin <sjshin@korea.ac.kr>

Description Implements a backward procedure for single and multiple change point detection proposed by Shin et al. <[arXiv:1812.10107](https://arxiv.org/abs/1812.10107)>. The backward approach is particularly useful to detect short and sparse signals which is common in copy number variation (CNV) detection.

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Depends R (>= 3.4.0)

Encoding UTF-8

LazyData true

RoxygenNote 6.1.1

NeedsCompilation yes

Author Seung Jun Shin [aut, cre], Yichao Wu [aut], Ning Hao [aut]

Repository CRAN

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bwd*Backward procedure for the change point detection*

Description

Implements backward procedure for detecting single or multiple change points.

Usage

```
bwd(y, alpha = 0.05, kmin = 3, lastkgroup = floor(0.01 * n),  
    mu0 = NULL, normal = T, n.permute = 1000, h = 10)
```

Arguments

<code>y</code>	observed data
<code>alpha</code>	target level that determines stopping criterion. Default is 0.05
<code>kmin</code>	minimum length of segments for checking possible change points
<code>lastkgroup</code>	We can avoid checking possible change points when we have less groups than "lastkgroup" to improve computational efficiency. Default is $0.01 * n$
<code>mu0</code>	Baseline mean value when detecting epidemic change points. Default is NULL
<code>normal</code>	if TRUE normal cutoff values are used, and if FALSE residual permuted cutoff values are used. Default is TRUE
<code>n.permute</code>	number of permutation when computing the permuted cutoff. Default is 1000
<code>h</code>	bandwidth size for variance estimator

Value

`bwd` object that contains information of detected segments and significance levels

Author(s)

Seung Jun Shin, Yicaho Wu, Ning Hao

References

Shin, Wu, and Hao (2018+) A backward procedure for change-point detection with applications to copy number variation detection, arXiv:1812.10107.

See Also

[plot.bwd](#)

Examples

```

# simulated data
set.seed(1)
n <- 1000
L <- 10

mu0 <- -0.5

mu <- rep(mu0, n)
mu[(n/2 + 1):(n/2 + L)] <- mu0 + 1.6
mu[(n/4 + 1):(n/4 + L)] <- mu0 - 1.6
y <- mu + rnorm(n)
alpha <- c(0.01, 0.05)

# BWD
obj1 <- bwd(y, alpha = alpha)

# Modified for epidemic changes with a known baseline mean, mu0.
obj2 <- bwd(y, alpha = alpha, mu0 = 0)

par(mfrow = c(2,1))
plot(obj1, y)
plot(obj2, y)

```

plot.bwd

plot for the backward procedure for the change point detection

Description

A plot of segments estimated by the backward procedure.

Usage

```

## S3 method for class 'bwd'
plot(x, y, ...)

```

Arguments

x	bwd object
y	observed data
...	graphical parameters

Value

plot of estimated segments

Author(s)

Seung Jun Shin, Yicaho Wu, Ning Hao

References

Shin, Wu, and Hao (2018+) A backward procedure for change-point detection with applications to copy number variation detection, arXiv:1812.10107.

See Also

[bwd](#)

Examples

```
# simulated data
set.seed(1)
n <- 1000
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mu0 <- -0.5

mu <- rep(mu0, n)
mu[(n/2 + 1):(n/2 + L)] <- mu0 + 1.6
mu[(n/4 + 1):(n/4 + L)] <- mu0 - 1.6
y <- mu + rnorm(n)
alpha <- c(0.01, 0.05)

# BWD
obj1 <- bwd(y, alpha = alpha)

# Modified for epidemic changes with a known baseline mean, mu0.
obj2 <- bwd(y, alpha = alpha, mu0 = 0)

par(mfrow = c(2,1))
plot(obj1, y)
plot(obj2, y)
```

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