

# Package ‘wbs’

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**Type** Package

**Title** Wild Binary Segmentation for Multiple Change-Point Detection

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**Author** Rafal Baranowski and Piotr Fryzlewicz

**Maintainer** Rafal Baranowski <package\_maintenance@rbaranowski.com>

**Depends** graphics

**Description** Provides efficient implementation of the Wild Binary Segmentation and Binary Segmentation algorithms for estimation of the number and locations of multiple change-points in the piecewise constant function plus Gaussian noise model.

**License** GPL-2

**NeedsCompilation** yes

**Repository** CRAN

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## Description

The package implements Wild Binary Segmentation, a technique for consistent estimation of the number and locations of multiple change-points in data. It also provides a fast implementation of the standard Binary Segmentation algorithm.

## Details

The main routines of the package are [wbs](#), [sbs](#) and [changepts](#).

## References

P. Fryzlewicz (2014), Wild Binary Segmentation for multiple change-point detection. *Annals of Statistics*, to appear. (<http://stats.lse.ac.uk/fryzlewicz/wbs/wbs.pdf>)

## Examples

```
#an example in which standard Binary Segmentation fails to detect change points
x <- rnorm(300)+ c(rep(0,130),rep(-1,20),rep(1,20),rep(0,130))

s <- sbs(x)
w <- wbs(x)

s.cpt <- changepts(s)
s.cpt

w.cpt <- changepts(w)
w.cpt
# in this example, both algorithms work well
x <- rnorm(300) + c(rep(1,50),rep(0,250))

s <- sbs(x)
w <- wbs(x)

s.cpt <- changepts(s)
s.cpt

w.cpt <- changepts(w)
w.cpt
```

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bic.penalty	<i>Bayesian Information Criterion penalty term</i>
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### Description

The function evaluates the penalty term for the standard Bayesian Information Criterion applied to the change-point detection problem. This routine is typically not called directly by the user; its name can be passed as an argument to [changepoints](#).

### Usage

```
bic.penalty(n, cpt)
```

### Arguments

n	the number of observations
cpt	a vector with localisations of change-points

### Value

the penalty term  $k \log(n)$  where  $k$  denotes the number of elements in `cpt`

### Examples

```
x <- rnorm(300) + c(rep(1,50),rep(0,250))
w <- wbs(x)
w.cpt <- changepoints(w,penalty="bic.penalty")
w.cpt$cpt.ic
x <- rnorm(300) + c(rep(1,50),rep(0,250))
w <- wbs(x)
w.cpt <- changepoints(w,penalty="bic.penalty")
w.cpt$cpt.ic
```

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changepoints	<i>Change-points detected by WBS or BS</i>
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### Description

The function applies user-specified stopping criteria to extract change-points from object generated by [wbs](#) or [sbs](#). For object of class 'sbs', the function returns change-points whose corresponding test statistic exceeds threshold given in `th`. For object of class 'wbs', the change-points can be also detected using information criteria with penalties specified in `penalty`.

**Usage**

```
changepoints(object, ...)

## S3 method for class 'sbs'
changepoints(object, th = NULL, th.const = 1.3,
             Kmax = NULL, ...)

## S3 method for class 'wbs'
changepoints(object, th = NULL, th.const = 1.3,
             Kmax = 50, penalty = c("ssic.penalty", "bic.penalty",
                                   "mbic.penalty"), ...)
```

**Arguments**

object	an object of 'wbs' or 'sbs' class returned by, respectively, <a href="#">wbs</a> and <a href="#">sbs</a> functions
...	further arguments that may be passed to the penalty functions
th	a vector of positive scalars
th.const	a vector of positive scalars
Kmax	a maximum number of change-points to be detected
penalty	a character vector with names of penalty functions used

**Details**

For the change-point detection based on thresholding (object of class 'sbs' or 'wbs'), the user can either specify the thresholds in `th` directly, determine the maximum number `Kmax` of change-points to be detected, or let `th` depend on `th.const`.

When `Kmax` is given, the function automatically sets `th` to the lowest threshold such that the number of detected change-points is lower or equal than `Kmax`. Note that for the BS algorithm it might be not possible to find the threshold such that exactly `Kmax` change-points are found.

When `th` and `Kmax` are omitted, the threshold value is set to

$$th = \sigma \times th.const \sqrt{2 \log(n)},$$

where  $\sigma$  is the Median Absolute Deviation estimate of the noise level and  $n$  is the number of elements in  $x$ .

For the change-point detection based on information criteria (object of class 'wbs' only), the user can specify both the maximum number of change-points (`Kmax`) and a type of the penalty used. Parameter `penalty` should contain a list of characters with names of the functions of at least two arguments ( $n$  and  $cpt$ ). For each penalty given, the following information criterion is minimized over candidate sets of change-points  $cpt$ :

$$\frac{n}{2} \log \hat{\sigma}_k^2 + \text{penalty}(n, cpt),$$

where  $k$  denotes the number of elements in  $cpt$ ,  $\hat{\sigma}_k$  is the corresponding maximum likelihood estimator of the residual variance.

**Value**

sigma	Median Absolute Deviation estimate of the noise level
th	a vector of thresholds
no.cpt.th	the number of change-points detected for each value of th
cpt.th	a list with the change-points detected for each value of th
Kmax	a maximum number of change-points detected
ic.curve	a list with values of the chosen information criteria
no.cpt.ic	the number of change-points detected for each information criterion considered
cpt.ic	a list with the change-points detected for each information criterion considered

**Examples**

```
#we generates gaussian noise + Poisson process signal with 10 jumps on average
set.seed(10)
N <- rpois(1,10)
true.cpt <- sample(1000,N)
m1 <- matrix(rep(1:1000,N),1000,N,byrow=FALSE)
m2 <- matrix(rep(true.cpt,1000),1000,N,byrow=TRUE)
x <- rnorm(1000) + apply(m1>=m2,1,sum)

# we apply the BS and WBS algorithms with default values for their parameters

s <- sbs(x)
w <- wbs(x)

s.cpt <- changepoints(s)
s.cpt

w.cpt <- changepoints(w)
w.cpt

#we can use different stopping criteria, invoking sbs/wbs functions is not necessary

s.cpt <- changepoints(s,th.const=c(1,1.3))
s.cpt
w.cpt <- changepoints(w,th.const=c(1,1.3))
w.cpt
```

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fixed.intervals

*Fixed intervals*


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**Description**

The function generates approximately M intervals with endpoints in 1,2,...,n, without random drawing. This routine can be used inside [wbs](#) function and is typically not called directly by the user.

**Usage**

```
fixed.intervals(n, M)
```

**Arguments**

n                    a number of endpoints to choose from  
M                    a number of intervals to generate

**Details**

Function finds the minimal  $m$  such that  $M \leq \frac{m(m-1)}{2}$ . Then it generates  $m$  approximately equally-spaced positive integers lower than  $n$  and returns all possible intervals consisting of any two of these points.

**Value**

a 2-column matrix with start (first column) and end (second column) points of an interval in each row

**See Also**

[random.intervals wbs](#)

**Examples**

```
fixed.intervals(10,100)
```

---

mbic.penalty

*Modified Bayes Information Criterion penalty term*

---

**Description**

The function evaluates the penalty term for the Modified Bayes Information Criterion proposed in N. Zhang and D. Siegmund (2007). This routine is typically not called directly by the user; its name can be passed as an argument to [changepoints](#).

**Usage**

```
mbic.penalty(n, cpt)
```

**Arguments**

n                    the number of observations  
cpt                  a vector with localisations of change-points

**Value**

the penalty term

$$\frac{3}{2}k \log(n) + \frac{1}{2} \sum_{i=1}^{k+1} \log \frac{l_i}{n},$$

where  $k$  denotes the number of elements in `cpt` and  $l_i$  are the lengths of the intervals between changepoints in `cpt`

**References**

N. Zhang and D. Siegmund (2007), A modified Bayes information criterion with applications to the analysis of comparative genomic hybridization data, *Biometrics*.

**Examples**

```
x <- rnorm(300) + c(rep(1,50),rep(0,250))
w <- wbs(x)
w.cpt <- changepoints(w,penalty="mbic.penalty")
w.cpt$cpt.ic
```

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means.between.cpt	<i>Means between change-points</i>
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**Description**

The function finds the average of the input vector `x` between change-points given in `cpt`.

**Usage**

```
means.between.cpt(x, cpt = NULL, ...)
```

**Arguments**

<code>x</code>	a vector
<code>cpt</code>	a vector of integers with localisations of change-points
<code>...</code>	further arguments passed to mean method

**Value**

a vector of the same length as `x`, piecewise constant and equal to the mean between change-points given in `cpt`

## Examples

```
x <- rnorm(100)+c(rep(-1,50),rep(1,50))
cpt <- 50
means.between.cpt(x,cpt)
w <- wbs(x)
cpt <- changepoints(w)
means.between.cpt(x,cpt=cpt$cpt.ic$sbic)
```

---

plot.sbs

*Plot for an 'sbs' object*

---

## Description

Plots the input vector used to generate 'sbs' object x with fitted piecewise constant function, equal to the mean between change-points specified in cpt.

## Usage

```
## S3 method for class 'sbs'
plot(x, cpt, ...)
```

## Arguments

x	an object of class 'sbs', returned by <a href="#">sbs</a>
cpt	a vector of integers with localisations of change-points
...	other parameters which may be passed to plot and changepoints

## Details

When cpt is omitted, the function automatically finds change-points using changepoints function with a default value of the threshold.

## See Also

[sbs changepoints](#)



---

plot.wbs	<i>Plot for a 'wbs' object</i>
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---

### Description

Plots the input vector used to generate 'wbs' object x with fitted piecewise constant function, equal to the mean between change-points specified in cpt.

### Usage

```
## S3 method for class 'wbs'  
plot(x, cpt, ...)
```

### Arguments

x	an object of class 'wbs', returned by <a href="#">wbs</a>
cpt	a vector of integers with localisations of change-points
...	other parameters which may be passed to plot and changepoints

### Details

When cpt is omitted, the function automatically finds change-points using changepoints function with strengthened Schwarz Information Criterion as a stopping criterion for the WBS algorithm.

### See Also

[wbs](#) [changepoints](#) [ssic](#).[penalty](#)

---

print.sbs	<i>Print for an 'sbs' object</i>
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---

### Description

Print for an 'sbs' object

### Usage

```
## S3 method for class 'sbs'  
print(x, ...)
```

### Arguments

x	an object of class 'sbs'
...	further arguments passed to print method

### See Also

[sbs](#)

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print.wbs	<i>Print for a 'wbs' object</i>
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**Description**

Print for a 'wbs' object

**Usage**

```
## S3 method for class 'wbs'  
print(x, ...)
```

**Arguments**

x	an object of class 'wbs'
...	further arguments passed to print method

**See Also**

[wbs](#)

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random.intervals	<i>Random intervals</i>
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**Description**

The function generates M intervals, whose endpoints are drawn uniformly without replacements from 1,2,..., n. This routine can be used inside [wbs](#) function and is typically not called directly by the user.

**Usage**

```
random.intervals(n, M)
```

**Arguments**

n	a number of endpoints to choose from
M	a number of intervals to generate

**Value**

a M by 2 matrix with start (first column) and end (second column) points of an interval in each row

**See Also**

[fixed.intervals](#) [wbs](#)

**Examples**

```
random.intervals(10,100)
```

---

sbs

---

*Change-point detection via standard Binary Segmentation*


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**Description**

The function applies the Binary Segmentation algorithm to identify potential locations of the change-points in the mean of the input vector  $x$ . The object returned by this routine can be further passed to the `changepoints` function, which finds the final estimate of the change-points based on thresholding.

**Usage**

```
sbs(x, ...)
```

```
## Default S3 method:
```

```
sbs(x, ...)
```

**Arguments**

<code>x</code>	a numeric vector
<code>...</code>	not in use

**Value**

an object of class "sbs", which contains the following fields

<code>x</code>	the vector provided
<code>n</code>	the length of $x$
<code>res</code>	a 6-column matrix with results, where 's' and 'e' denote start- end points of the intervals in which change-points candidates 'cpt' have been found; column 'CUSUM' contains corresponding value of CUSUM statistic; 'min.th' is the smallest threshold value for which given change-point candidate would be not added to the set of estimated change-points; the last column is the scale at which the change-point has been found

**Examples**

```
x <- rnorm(300) + c(rep(1,50),rep(0,250))
s <- sbs(x)
s.cpt <- changepoints(s)
s.cpt
th <- c(s.cpt$th,0.7*s.cpt$th)
s.cpt <- changepoints(s,th=th)
s.cpt
```

---

ssic.penalty	<i>Strengthened Schwarz Information Criterion penalty term</i>
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### Description

The function evaluates the penalty term for the strengthened Schwarz Information Criterion proposed in P. Fryzlewicz (2014). This routine is typically not called directly by the user; its name can be passed as an argument to `changepoints`.

### Usage

```
ssic.penalty(n, cpt, alpha = 1.01, ssic.type = c("log", "power"))
```

### Arguments

<code>n</code>	the number of observations
<code>cpt</code>	a vector with localisations of change-points
<code>alpha</code>	a scalar greater than one
<code>ssic.type</code>	a string ("log" or "power")

### Value

the penalty term  $k(\log(n))^{\alpha}$  for `ssic.penalty="log"` or  $kn^{\alpha}$  for `ssic.penalty="power"`, where  $k$  denotes the number of elements in `cpt`

### References

P. Fryzlewicz (2014), Wild Binary Segmentation for multiple change-point detection. *Annals of Statistics*, to appear. (<http://stats.lse.ac.uk/fryzlewicz/wbs/wbs.pdf>)

### Examples

```
x <- rnorm(300) + c(rep(1,50),rep(0,250))
w <- wbs(x)
w.cpt <- changepoints(w,penalty="ssic.penalty")
w.cpt$cpt.ic
```

**Description**

The function applies the Wild Binary Segmentation algorithm to identify potential locations of the change-points in the mean of the input vector  $x$ . The object returned by this routine can be further passed to the [changepoints](#) function, which finds the final estimate of the change-points based on chosen stopping criteria.

**Usage**

```
wbs(x, ...)
```

```
## Default S3 method:
wbs(x, M = 5000, rand.intervals = TRUE,
    integrated = TRUE, ...)
```

**Arguments**

<code>x</code>	a numeric vector
<code>...</code>	not in use
<code>M</code>	a number of intervals used in the WBS algorithm
<code>rand.intervals</code>	a logical variable; if <code>rand.intervals=TRUE</code> intervals used in the procedure are random, thus the output of the algorithm may slightly vary from run to run; for <code>rand.intervals=FALSE</code> the intervals used depend on <code>M</code> and the length of <code>x</code> only, hence the output is always the same for given input parameters
<code>integrated</code>	a logical variable indicating the version of Wild Binary Segmentation algorithm used; when <code>integrated=TRUE</code> , augmented version of WBS is launched, which combines WBS and BS into one

**Value**

an object of class "wbs", which contains the following fields

<code>x</code>	the input vector provided
<code>n</code>	the length of <code>x</code>
<code>M</code>	the number of intervals used
<code>rand.intervals</code>	a logical variable indicating type of intervals
<code>integrated</code>	a logical variable indicating type of WBS procedure
<code>res</code>	a 6-column matrix with results, where 's' and 'e' denote start- end points of the intervals in which change-points candidates 'cpt' have been found; column 'CUSUM' contains corresponding value of CUSUM statistic; 'min.th' is the smallest threshold value for which given change-point candidate would be not added to the set of estimated change-points; the last column is the scale at which the change-point has been found

**Examples**

```
x <- rnorm(300) + c(rep(1,50),rep(0,250))
w <- wbs(x)
plot(w)
w.cpt <- changepoints(w)
w.cpt
th <- c(w.cpt$th,0.7*w.cpt$th)
w.cpt <- changepoints(w,th=th)
w.cpt$cpt.th
```

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