

# Package ‘cpss’

November 23, 2020

**Title** Change-Point Detection by Sample-Splitting Methods

**Version** 0.0.2

**Description** Implements multiple change searching algorithms for a variety of frequently considered parametric change-point models. In particular, it integrates a criterion proposed by Zou, Wang and Li (2020) <doi:10.1214/19-AOS1814> to select the number of change-points in a data-driven fashion. Moreover, it also provides interfaces for users-customized change-point models with their own cost function and estimation routines.

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**LinkingTo** Rcpp, RcppArmadillo

**Imports** Rcpp, magrittr, methods, stats, mvtnorm, Rfast, tibble, dplyr, tidyr, rlang, ggplot2, gridExtra

**Suggests** MASS, L1pack

**NeedsCompilation** yes

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**Repository** CRAN

**Date/Publication** 2020-11-23 09:40:15 UTC

## R topics documented:

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coef , cpss-method	<i>coef method</i>
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### Description

coef method

### Usage

```
## S4 method for signature 'cpss'
coef(object)
```

### Arguments

object	object
cpss	cpss

---

cpss-class	<i>cpss: an S4 class which collects data and information required for further change-point analyses and summaries</i>
------------	---

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### Description

cpss: an S4 class which collects data and information required for further change-point analyses and summaries

**Slots**

dat ANY.  
mdl character.  
algo character.  
algo\_param\_dim numeric.  
SC character.  
ncps integer.  
pelt\_pen numeric.  
cps numeric.  
params list.  
S\_vals numeric.  
SC\_vals matrix.  
call list.  
update.inputs list.

---

cpss.custom

*Detecting changes in users-customized models*

---

**Description**

Detecting changes in users-customized models

**Usage**

```
cpss.custom(  
  dataset,  
  n,  
  g_subdat,  
  g_param,  
  g_cost,  
  algorithm = "BS",  
  dist_min = floor(log(n)),  
  ncps_max = ceiling(n^0.4),  
  pelt_pen_val = NULL,  
  pelt_K = 0,  
  wbs_nintervals = 500,  
  criterion = "CV",  
  times = 2,  
  model = NULL,  
  g_smry = NULL,  
  easy_cost = NULL,  
  param.opt = NULL  
)
```

**Arguments**

dataset	an ANY object that could be of any form such as a vector, matrix, tensor, list, etc.
n	an integer indicating the sample size of the dataset.
g_subdat	a customized R function of two arguments <code>dat</code> and <code>indices</code> , that returns a subset of the <code>dat</code> (inheriting the class from that of <code>dataset</code> ) according to given <code>indices</code> along the observed time orders. The argument <code>indices</code> is a logical vector with <code>TRUE</code> indicating selected indices.
g_param	a customized R function of two arguments, <code>dat</code> and <code>param.opt</code> , that returns estimates of interested parameters that minimizes users-specified cost for a data set <code>dat</code> . The returned object could be of any class such as a numeric value, vector, matrix, list, etc. The argument <code>param.opt</code> might be used in the estimation procedures.
g_cost	a customized R function of two arguments, <code>dat</code> and <code>param</code> , that returns a numeric value of associated cost for a data set <code>dat</code> , under the knowledge of the interested parameters being <code>param</code> . The argument <code>param</code> inherits from the class of the returned object of the function <code>g_param</code> . If <code>param.opt</code> is needed to evaluate the cost, they should be packed into <code>param</code> when defining the function <code>g_param</code> .
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
dist_min	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
ncps_max	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
pelt_pen_val	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
pelt_K	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
wbs_nintervals	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
criterion	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
times	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.
model	a character string indicating the considered change model, and will be set as "custom" if not provided.
g_smry	a customized R function of two arguments <code>dataset</code> and <code>param.opt</code> , which calculates the summary statistics that will be needed in evaluations of the cost. The returned object is a list for convenience.
easy_cost	a customized R function of three arguments <code>data_smry</code> , <code>s</code> and <code>e</code> , that evaluates the cost for a date segment form observed time point <code>\$\$s</code> to <code>\$e\$</code> . The argument <code>data_smry</code> inherits from the returned list of the function <code>g_smry</code> .
param.opt	an ANY object that could be of any form, specifying additional global constant parameters beyond the interested parameters.

**Value**

`cpss.custom` returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries.

`dat` an ANY object inheriting from the type of user-input data

`mdl` a character string describing considered change-point model

`algo` a character string indicating user-specified change-point searching algorithm

`algo_param_dim` an integer indicating user-specified maximum number of change-points searched for if the algorithm is chosen among "SN", "BS" and "WBS", or a numeric vector collecting user-specified values for the penalty if the algorithm is "PELT"

`SC` a character string indicating model selection criterion

`ncps` an integer giving estimated number of change-points based on the entire data

`pelt_pen` a numeric value indicating selected penalty value if the "PELT" algorithm is performed based on the entire data

`cps` a numeric vector of detected change-points based on the entire data

`params` a list object, each of whose members is a list containing estimated parameters in the corresponding segment

`S_vals` a numeric vector of candidate model dimensions in terms of a sequence of numbers of change-points or values of penalty

`SC_vals` a numeric matrix, each column of which records the values of criterion based on the validation data under the corresponding model dimension (`S_vals`), and each row of which represents a splitting at each time

**References**

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

**Examples**

```
library("cpss")
if (!requireNamespace("L1pack", quietly = TRUE)) {
  stop("Please install the package \"L1pack\".")
}
set.seed(666)
n <- 1000
tau <- c(250, 500, 750)
tau_ext <- c(0, tau, n)
be0 <- c(1, 1, 0, -1)
be <- c(1, -1, -1, 1)
seg_len <- diff(c(0, tau, n))
x <- rnorm(n)
eta <- unlist(lapply(seq(1, length(tau) + 1), function(k) {
  be0[k] + be[k] * x[(tau_ext[k] + 1):tau_ext[k + 1]]
})))
ep <- L1pack::rlaplace(n)
y <- eta + ep
```

```

g_subdat_l1 <- function(dat, indices) {
  matrix(dat[indices, ], sum(indices), ncol(dat))
}
g_param_l1 <- function(dat, param.opt = NULL) {
  y <- dat[, 1]
  x <- dat[, -1]
  return(L1pack::l1fit(x, y)$coefficients)
}
g_cost_l1 <- function(dat, param) {
  y <- dat[, 1]
  x <- dat[, -1]
  return(sum(abs(y - cbind(1, x) %*% as.matrix(param))))
}
res <- cpss.custom(
  dataset = cbind(y, x), n = n,
  g_subdat = g_subdat_l1, g_param = g_param_l1, g_cost = g_cost_l1,
  algorithm = "BS", dist_min = 10, ncps_max = 10,
  g_smry = NULL, easy_cost = NULL
)
summary(res)
# 250 500 744
do.call(rbind, res@params)
# Intercept      X
# [1,] 0.9327557 0.9558247
# [2,] 0.9868086 -1.0254999
# [3,] -0.0464067 -0.9076744
# [4,] -0.9746133 0.9671701

```

---

cpss.em

*Detecting changes in exponential family*


---

## Description

Detecting changes in exponential family

## Usage

```

cpss.em(
  dataset,
  family,
  size = NULL,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2
)

```

**Arguments**

dataset	a numeric matrix of dimension $n \times d$ , where each row represents an observation and each column stands for a variable. A numeric vector could also be acceptable for univariate observations.
family	a character string indicating the underlying distribution. Currently, detecting changes in binomial ("binom"), multinomial ("multinom"), Poisson ("pois"), exponential ("exp"), geometric ("geom"), dirichlet ("diri"), gamma ("gamma"), beta ("beta"), chi-square ("chisq") and inverse gaussian ("invgauss") distributions are supported.
size	an integer indicating the number of trials if family = "binom" or family = "multinom".
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
dist_min	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
ncps_max	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
pelt_pen_val	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
pelt_K	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
wbs_nintervals	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
criterion	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
times	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.

**Value**

cpss.em returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).

**References**

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

**See Also**

[cpss.meanvar](#) [cpss.mean](#) [cpss.var](#)

**Examples**

```

library("cpss")
set.seed(666)
n <- 1000
tau <- c(100, 300, 700, 900)
tau_ext <- c(0, tau, n)
theta <- c(1, 0.2, 1, 0.2, 1)
seg_len <- diff(c(0, tau, n))
y <- unlist(lapply(seq(1, length(tau) + 1), function(k) {
  rexp(seg_len[k], theta[k])
})))
res <- cpss.em(
  y, family = "exp", algorithm = "WBS",
  dist_min = 10, ncps_max = 10,
  criterion = "MS", times = 10
)
cps(res)
# [1] 100 299 705 901

```

---

cpss.glm

*Detecting changes in GLMs*


---

**Description**

Detecting changes in GLMs

**Usage**

```

cpss.glm(
  formula,
  family,
  data = NULL,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2
)

```

**Arguments**

formula	a formula object describing the change-point model to be fitted.
family	a description of the error distribution and link function to be used in the model, which can be a character string naming a family function or a family function.



<code>data</code>	an optional data frame, list or environment containing the variables in the model.
<code>algorithm</code>	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
<code>dist_min</code>	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
<code>ncps_max</code>	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
<code>pelt_pen_val</code>	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
<code>pelt_K</code>	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
<code>wbs_nintervals</code>	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
<code>criterion</code>	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
<code>times</code>	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.

### Value

`cpss.glm` returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).

### References

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

### See Also

[cpss.lm](#)

### Examples

```
library("cpss")
set.seed(666)
n <- 200
size <- rpois(n, 20 - 1) + 1
tau <- c(75, 100, 175)
tau_ext <- c(0, tau, n)
be <- list(c(0, 0.5), c(0, -0.5), c(0.5, -0.5), c(-0.5, -0.5))
seg_len <- diff(c(0, tau, n))
x <- rnorm(n)
eta <- lapply(seq(1, length(tau) + 1), function(k) {
  be[[k]][1] + be[[k]][2] * x[(tau_ext[k] + 1):tau_ext[k + 1]]
})
```

```

eta <- do.call(c, eta)
p <- 1 / (1 + exp(-eta))
y <- rbinom(n, size = size, prob = p)

pelt_pen_val <- (log(n))^seq(0.5, 2, by = 0.1)
res <- cpss.glm(
  formula = cbind(y, size - y) ~ x, family = binomial(),
  algorithm = "PELT", pelt_pen_val = pelt_pen_val,
  dist_min = 5, ncps_max = 10
)
summary(res)
# 75 105 175
coef(res)
# [1,] 0.02540872 0.08389551 0.5284425 -0.4980768
# [2,] 0.57222684 -0.45430385 -0.5203319 -0.4581678

```

---

cpss.lm

*Detecting changes in linear models*


---

## Description

Detecting changes in linear models

## Usage

```

cpss.lm(
  formula,
  data = NULL,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2
)

```

## Arguments

formula	a formula object describing the change-point model to be fitted.
data	an optional data frame, list or environment containing the variables in the model.
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.

<code>dist_min</code>	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
<code>ncps_max</code>	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
<code>pelt_pen_val</code>	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
<code>pelt_K</code>	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
<code>wbs_nintervals</code>	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
<code>criterion</code>	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
<code>times</code>	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.

### Value

`cpss.lm` returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).

### References

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

### See Also

[cpss.glm](#)

### Examples

```
library("cpss")
set.seed(666)
n <- 400
tau <- c(80, 200, 300)
tau_ext <- c(0, tau, n)
be <- list(c(0, 1), c(1, 0.5), c(0, 1), c(-1, 0.5))
seg_len <- diff(c(0, tau, n))
x <- rnorm(n)
mu <- lapply(seq(1, length(tau) + 1), function(k) {
  be[[k]][1] + be[[k]][2] * x[(tau_ext[k] + 1):tau_ext[k + 1]]
})
mu <- do.call(c, mu)
sig <- unlist(lapply(seq(1, length(tau) + 1), function(k) {
  rep(be[[k]][2], seg_len[k])
}))
y <- rnorm(n, mu, sig)
res <- cpss.lm(
  formula = y ~ x,
```

```

algorithm = "BS",
dist_min = 5, ncps_max = 10
)
summary(res)
# 80 202 291
coef(res)
# $coef
#           [,1]      [,2]      [,3]      [,4]
# [1,] -0.00188792 1.0457718 -0.03963209 -0.9444813
# [2,]  0.91061557 0.6291965  1.20694409  0.4410036
#
# $sigma
# [1] 0.8732233 0.4753216 0.9566516 0.4782329

```

---

cpss.mean

*Detecting changes in mean*


---

## Description

Detecting changes in mean

## Usage

```

cpss.mean(
  dataset,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2,
  Sigma = NULL
)

```

## Arguments

dataset	a numeric matrix of dimension $n \times d$ , where each row represents an observation and each column stands for a variable. A numeric vector could also be acceptable for univariate observations.
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
dist_min	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .

ncps_max	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
pelt_pen_val	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
pelt_K	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
wbs_nintervals	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
criterion	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
times	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.
Sigma	if a numeric matrix (or constant) is supplied, it would be taken as the value of known overall covariance (or variance). By default it is set as NULL, and the common covariance of the data is estimated based on the difference method, i.e.,

$$\hat{\Sigma} = \frac{1}{2(n-1)} \sum_{i=1}^{n-1} (Y_i - Y_{i+1})(Y_i - Y_{i+1})'$$

### Value

cpss.mean returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).

### References

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

### See Also

[cpss.meanvar](#) [cpss.var](#)

### Examples

```
library("cpss")
set.seed(666)
n <- 2048
tau <- c(205, 267, 308, 472, 512, 820, 902, 1332, 1557, 1598, 1659)
seg_len <- diff(c(0, tau, n))
mu <- rep(c(0, 14.64, -3.66, 7.32, -7.32, 10.98, -4.39, 3.29, 19.03, 7.68, 15.37, 0), seg_len)
ep <- 7 * rnorm(n)
y <- mu + ep

res <- cpss.mean(y, algorithm = "SN", dist_min = 10, ncps_max = 20)
summary(res)
# 205 267 307 471 512 820 897 1332 1557 1601 1659
plot(res, type = "scatter")
```

```

plot(res, type = "path")
out <- update(res, dim_update = 12)
out$cps_update
# 205 267 307 471 512 820 897 1332 1557 1601 1659 1769
out$params_update

```

---

cpss.meanvar

*Detecting changes in mean and (co)variance*


---

## Description

Detecting changes in mean and (co)variance

## Usage

```

cpss.meanvar(
  dataset,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2
)

```

## Arguments

dataset	a numeric matrix of dimension $n \times d$ , where each row represents an observation and each column stands for a variable. A numeric vector could also be acceptable for univariate observations.
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
dist_min	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
ncps_max	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
pelt_pen_val	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
pelt_K	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).

- `wbs_nintervals` an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
- `criterion` a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
- `times` an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.

### Value

`cpss.meanvar` returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).

### References

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

### See Also

[cpss.mean](#) [cpss.var](#)

### Examples

```
library("cpss")
if (!requireNamespace("MASS", quietly = TRUE)) {
  stop("Please install the package \"MASS\".")
}
set.seed(666)
n <- 1000
tau <- c(200, 400, 600, 800)
mu <- list(rep(0, 2), rep(1, 2), rep(1, 2), rep(0, 2), rep(0, 2))
Sigma <- list(diag(2), diag(2), matrix(c(1,-1,-1, 4), 2), matrix(c(1, 0.5, 0.5, 1), 2), diag(2))
seg_len <- diff(c(0, tau, n))
y <- lapply(seq(1, length(tau) + 1), function(k) {
  MASS::mvrnorm(n = seg_len[k], mu = mu[[k]], Sigma = Sigma[[k]])
})
y <- do.call(rbind, y)
res <- cpss.meanvar(y, algorithm = "BS", dist_min = 20)
cps(res)
# [1] 211 402 598 804
plot(res, type = "coef")
```

---

cpss.var

*Detecting changes in (co)variance*

---

### Description

Detecting changes in (co)variance

**Usage**

```
cpss.var(
  dataset,
  algorithm = "BS",
  dist_min = floor(log(n)),
  ncps_max = ceiling(n^0.4),
  pelt_pen_val = NULL,
  pelt_K = 0,
  wbs_nintervals = 500,
  criterion = "CV",
  times = 2,
  mu = NULL
)
```

**Arguments**

dataset	a numeric matrix of dimension $n \times d$ , where each row represents an observation and each column stands for a variable. A numeric vector could also be acceptable for univariate observations.
algorithm	a character string specifying the change-point searching algorithm, one of four state-of-the-art candidates "SN" (segment neighborhood), "BS" (binary segmentation), "WBS" (wild binary segmentation) and "PELT" (pruned exact linear time) algorithms.
dist_min	an integer indicating the minimum distance between two successive candidate change-points, with a default value $\text{floor}(\log(n))$ .
ncps_max	an integer indicating the maximum number of change-points searched for, with a default value $\text{ceiling}(n^{0.4})$ .
pelt_pen_val	a numeric vector specifying the collection of candidate values of the penalty if the "PELT" algorithm is used.
pelt_K	a numeric value to adjust the pruning tactic, usually is taken to be 0 if negative log-likelihood is used as a cost; more details can be found in Killick et al. (2012).
wbs_nintervals	an integer indicating the number of random intervals drawn in the "WBS" algorithm and a default value 500 is used.
criterion	a character string indicating which model selection criterion, "cross-validation" ("CV") or "multiple-splitting" ("MS"), is used.
times	an integer indicating how many times of sample-splitting should be performed; if "CV" criterion is used, it should be set as 2.
mu	if a numeric vector or constant is supplied, it would be taken as the value of known overall mean. By default it is set as NULL, and the common mean of the data is estimated by the sample mean based on the entire data set.

**Value**

cpss.var returns an object of an S4 class, called "cpss", which collects data and information required for further change-point analyses and summaries. See [cpss.custom](#).



**References**

Killick, R., Fearnhead, P., and Eckley, I. A. (2012). Optimal Detection of Changepoints With a Linear Computational Cost. *Journal of the American Statistical Association*, 107(500):1590–1598.

**See Also**

[cpss.meanvar](#) [cpss.mean](#)

**Examples**

```
library("cpss")
if (!requireNamespace("MASS", quietly = TRUE)) {
  stop("Please install the package \"MASS\".")
}
set.seed(666)
n <- 1000
tau <- c(200, 500, 750)
mu <- list(rep(0, 2), rep(0, 2), rep(0, 2), rep(0, 2))
Sigma <- list(diag(2), matrix(c(1, 0, 0, 4), 2), matrix(c(1, -0.5, -0.5, 4), 2), diag(2))
seg_len <- diff(c(0, tau, n))
y <- lapply(seq(1, length(tau) + 1), function(k) {
  MASS::mvrnorm(n = seg_len[k], mu = mu[[k]], Sigma = Sigma[[k]])
})
y <- do.call(rbind, y)
res <- cpss.var(y, algorithm = "BS", dist_min = 20)
cps(res)
# [1] 215 515 751
```

---

plot,cpss-method

*plot method*

---

**Description**

plot method

**Usage**

```
## S4 method for signature 'cpss'
plot(obj, type, x = c(), y = c(), ...)
```

**Arguments**

obj	obj
type	type
x	x
y	y
...	...
cpss	cpss

summary, cpss-method    *summary method*

---

**Description**

summary method

**Usage**

```
## S4 method for signature 'cpss'  
summary(object)
```

**Arguments**

object	object
cpss	cpss

---

tool1                    *tool1*

---

**Description**

tool1

**Usage**

```
dat(x)  
mdl(x)  
algo(x)  
algo_param_dim(x)  
SC(x)  
ncps(x)  
pelt_pen(x)  
cps(x)  
params(x)  
S_vals(x)
```

SC\_vals(x)

upcalle.inputs(x)

### Arguments

x                    x

---

tool2

*tool2*

---

### Description

tool2

### Usage

dat(x) <- value

mdl(x) <- value

algo(x) <- value

algo\_param\_dim(x) <- value

SC(x) <- value

ncps(x) <- value

pelt\_pen(x) <- value

cps(x) <- value

params(x) <- value

S\_vals(x) <- value

SC\_vals(x) <- value

upcalle.inputs(x) <- value

### Arguments

x                    x

value                value

---

tool3

*tool3*

---

### Description

tool3

### Usage

```
## S4 method for signature 'cpss'  
dat(x)
```

```
## S4 method for signature 'cpss'  
mdl(x)
```

```
## S4 method for signature 'cpss'  
algo(x)
```

```
## S4 method for signature 'cpss'  
algo_param_dim(x)
```

```
## S4 method for signature 'cpss'  
SC(x)
```

```
## S4 method for signature 'cpss'  
ncps(x)
```

```
## S4 method for signature 'cpss'  
pelt_pen(x)
```

```
## S4 method for signature 'cpss'  
cps(x)
```

```
## S4 method for signature 'cpss'  
params(x)
```

```
## S4 method for signature 'cpss'  
S_vals(x)
```

```
## S4 method for signature 'cpss'  
SC_vals(x)
```

```
## S4 method for signature 'cpss'  
upcalle.inputs(x)
```

### Arguments

x

x

cpss            cpss

---

tool4            *tool4*

---

## Description

tool4

## Usage

```
## S4 replacement method for signature 'cpss'  
dat(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
mdl(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
algo(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
algo_param_dim(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
SC(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
ncps(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
pelt_pen(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
cps(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
params(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
S_vals(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
SC_vals(x) <- value
```

```
## S4 replacement method for signature 'cpss'  
upcalle.inputs(x) <- value
```

**Arguments**

x	x
value	value
cpss	cpss

---

update, cpss-method	<i>update method</i>
---------------------	----------------------

---

**Description**

update method

**Usage**

```
## S4 method for signature 'cpss'  
update(object, dim_update)
```

**Arguments**

object	object
dim_update	dim_update
cpss	cpss

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