

# Package ‘jcp’

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**Type** Package

**Title** Joint Change Point Detection

**Version** 1.0

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**Description** Procedures for joint detection of changes in both expectation and variance in univariate sequences. Performs a statistical test of the null hypothesis of the absence of change points. In case of rejection performs an algorithm for change point detection. References - Bivariate change point detection (2019+), Michael Messer.

**License** GPL-3

**RoxygenNote** 6.1.1

**NeedsCompilation** no

**Repository** CRAN

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## R topics documented:

jcp . . . . .	1
plot.jcp . . . . .	3
summary.jcp . . . . .	4

<b>Index</b>	<b>6</b>
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jcp	<i>jcp</i>
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## Description

Joint change point detection - expectation and variance - via bivariate moving sum statistics

**Usage**

```
jcp(x, H = NA, q = NA, alpha = 0.05, sim = 1000,
    region = "square")
```

**Arguments**

x	numeric vector. Input sequence of random variables.
H	NA or numeric vector. Window set. If NA (default), then H is automatically set. If not NA, then H must be an increasing vector of positive integers with maximum $\leq \text{length}(x)/2$ .
q	NA or numeric value. Rejection threshold. If NA (default), then the rejection boundary is derived in simulations (from Gaussian process limit) according to sim and alpha. If not NA, then q is considered predefined and must be set as a positive real number.
alpha	numeric value. Significance level. Must be in (0,1), default = 0.05. In case of predefined q, alpha is set to NA.
sim	numeric value. Number of simulations of limit process for approximation of q. Must be positive integer, default = 1000. In case of predefined q, sim is set to NA.
region	character string. Defines rejection region, default = "square". Must be chosen either "square", "circle" or "ellipse".

**Value**

invisible	
changepoints	detected change points (increasingly ordered)
mean_sd	matrix of estimated means and standard deviations
M	test statistic
q	rejection threshold
H	window set
sim	number of simulations of the limit process (approximation of q)
alpha	significance level
region	rejection region
method	derivation of threshold q, either asymptotic or predefined
x	input sequence
EVrho	list containing the auxiliary processes E, V and correlation rho, for each element of H one list entry
CP_meta	matrix containing meta information of estimation. Estimated change points (increasingly ordered), responsible window h, components E, V and rho of joint statistic at estimated change points (regarding responsible window)
SFA	detected change points of single filter algorithms

**Author(s)**

Michael Messer

**References**

Bivariate change point detection (2019+), Michael Messer

**See Also**[plot.jcp](#), [summary.jcp](#)**Examples**

```
# Normal distributed sequence with 3 change points at
# c1=250 (change in expectation),
# c2=500 (change in variance) and
# c3=750 (change in expectation and variance)
set.seed(0)
m      <- c(8,10,10,3);  s  <- c(4,4,10,5)
x      <- rnorm(1000, mean=rep(m,each=250), sd=rep(s,each=250))
result <- jcp(x)
summary(result)
plot(result)

# Set additional parameters (window set)
result2 <- jcp(x,H=c(80,160,240))
summary(result2)
plot(result2)
```

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plot.jcp*plot.jcp*

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**Description**

Plot method for class 'jcp'

**Usage**

```
## S3 method for class 'jcp'
plot(x, cex = 1, cex.main = 1, ...)
```

**Arguments**

x	object of class jcp
cex	numeric, global sizes in plot
cex.main	numeric, size of titles
...	additional arguments

**Author(s)**

Michael Messer

**References**

Bivariate change point detection (2019+), Michael Messer

**See Also**

[jcp](#), [summary.jcp](#)

**Examples**

```
# Normal distributed sequence with 3 change points at
# c1=250 (change in expectation),
# c2=500 (change in variance) and
# c3=750 (change in expectation and variance)
set.seed(0)
m      <- c(8,10,10,3);  s  <- c(4,4,10,5)
x      <- rnorm(1000, mean=rep(m,each=250), sd=rep(s,each=250))
result <- jcp(x)
summary(result)
plot(result)

# Set additional parameters (window set)
result2 <- jcp(x,H=c(80,160,240))
summary(result2)
plot(result2)
```

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summary.jcp

*summary.jcp*


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**Description**

Summary method for class 'jcp'

**Usage**

```
## S3 method for class 'jcp'
summary(object, ...)
```

**Arguments**

object	object of class jcp
...	additional arguments

**Author(s)**

Michael Messer

**References**

Bivariate change point detection (2019+), Michael Messer

**See Also**

[jcp](#), [plot.jcp](#)

**Examples**

```
#' # Normal distributed sequence with 3 change points at
# c1=250 (change in expectation),
# c2=500 (change in variance) and
# c3=750 (change in expectation and variance)
set.seed(0)
m      <- c(8,10,10,3);  s  <- c(4,4,10,5)
x      <- rnorm(1000, mean=rep(m,each=250), sd=rep(s,each=250))
result <- jcp(x)
plot(result)
summary(result)

# Set additional parameters (window set)
result2 <- jcp(x,H=c(80,160,240))
plot(result2)
summary(result2)
```

# Index

jcp, [1](#), [4](#), [5](#)

plot.jcp, [3](#), [3](#), [5](#)

summary.jcp, [3](#), [4](#), [4](#)