

Package ‘convo’

September 26, 2018

Type Package

Title Fast Computation of Running Sample Statistics

Version 0.1.0

Description Provides methods for fast computation of running (aka rolling) sample statistics. These include running sample mean of a single numeric sequence, running sample variance of a single numeric sequence, running sample covariance of two numeric sequences. Implementation of the methods uses convolution via Fast Fourier Transform. The complexity of the convolution can be reduced from $O(n^2)$ to $O(n \log n)$ with fast Fourier transform compared to conventional computation.

License GPL-3

Imports stats

Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

Suggests knitr,
rmarkdown,
testthat,
covr

VignetteBuilder knitr

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RunningCor

*Fast Running Correlation Computation***Description**

Computes running correlation between two sequences in a fixed width window, whose length corresponds to the length of the shorter sequence. Uses convolution via Fast Fourier Transform.

Usage

```
RunningCor(x, y, circular = FALSE)
```

Arguments

x	numeric sequence
y	numeric sequence, of equal or shorter length than x sequence
circular	logical; whether running correlation is computed assuming circular nature of x sequence (see Details)

Details

Computes running correlation between two sequences in a fixed width window. The length of a window is equal to the shorter of the two sequences (y), and window "runs" over the length of longer sequence (x).

Parameter circular determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, l_y is the length of shorter sequence y.

If circular equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample correlation between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample correlation between $c(x[1_x], x[1:(l_y - 1)])$ and y.

If circular equals FALSE then

- output sequence length equals $l_x - l_y + 1$,
- first element of the output sequence corresponds to sample correlation between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample correlation between $x[(l_x - l_y + 1):l_x]$.

Value

numeric sequence

Examples

```
x <- sin(seq(0, 1, length.out = 1000) * 2 * pi * 6)
y <- x[1:100]
out1 <- RunningCor(x, y, circular = TRUE)
out2 <- RunningCor(x, y, circular = FALSE)
plot(out1, type = "l"); points(out2, col = "red")
```

RunningCov

*Fast Running Covariance Computation***Description**

Computes running covariance between two sequences in a fixed width window, whose length corresponds to the length of the shorter sequence. Uses convolution via Fast Fourier Transform.

Usage

```
RunningCov(x, y, circular = FALSE)
```

Arguments

x	numeric sequence
y	numeric sequence, of equal or shorter length than x sequence
circular	logical; whether running variance is computed assuming circular nature of x sequence (see Details)

Details

Computes running covariance between two sequences in a fixed width window. The length of a window is equal to the shorter of the two sequences (y), and window "runs" over the length of longer sequence (x).

Parameter circular determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, l_y is the length of shorter sequence y.

If circular equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample covariance between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample covariance between $c(x[1_x], x[1:(l_y - 1)])$ and y.

If circular equals FALSE then

- output sequence length equals $l_x - l_y + 1$,
- first element of the output sequence corresponds to sample covariance between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample covariance between $x[(l_x - l_y + 1):l_x]$.

Value

numeric sequence

Examples

```
x <- sin(seq(0, 1, length.out = 1000) * 2 * pi * 6)
y <- x[1:100]
out1 <- RunningCov(x, y, circular = TRUE)
out2 <- RunningCov(x, y, circular = FALSE)
plot(out1, type = "l"); points(out2, col = "red")
```

RunningL2Norm

Fast Running L2 Norm Computation

Description

Computes running L2 norm between two sequences in a fixed width window, whose length corresponds to the length of the shorter sequence. Uses convolution via Fast Fourier Transform.

Usage

```
RunningL2Norm(x, y, circular = FALSE)
```

Arguments

x	numeric sequence
y	numeric sequence, of equal or shorter length than x sequence
circular	logical; whether running L2 norm is computed assuming circular nature of x sequence (see Details)

Details

Computes running L2 norm between two sequences in a fixed width window. The length of a window is equal to the shorter of the two sequences (y), and window "runs" over the length of longer sequence (x).

Parameter circular determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, l_y is the length of shorter sequence y.

If circular equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample L2 norm between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample L2 norm between $c(x[1_x], x[1:(l_y - 1)])$ and y.

If circular equals FALSE then

- output sequence length equals $l_x - l_y + 1$,
- first element of the output sequence corresponds to sample L2 norm between $x[1:l_y]$ and y,
- last element of the output sequence corresponds to sample L2 norm between $x[(l_x - l_y + 1):l_x]$.

Value

numeric sequence

Examples

```
x <- sin(seq(0, 1, length.out = 1000) * 2 * pi * 6)
y1 <- x[1:100] + rnorm(100)
y2 <- rnorm(100)
out1 <- RunningL2Norm(x, y1)
out2 <- RunningL2Norm(x, y2)
plot(out1, type = "l"); points(out2, col = "blue")
```

RunningMean	<i>Fast Running Mean Computation</i>
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Description

Computes running sample mean of a sequence in a fixed width window. Uses convolution via Fast Fourier Transform.

Usage

```
RunningMean(x, W, circular = FALSE)
```

Arguments

x	numeric sequence
W	numeric; width of x sequence window
circular	logical; whether running sample mean is computed assuming circular nature of x sequence (see Details)

Details

Parameter circular determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, W is a fixed length of x sequence window.

If circular equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample mean of $x[1:W]$,
- last element of the output sequence corresponds to sample mean of $c(x[1_x], x[1:(W - 1)])$.

If circular equals FALSE then

- output sequence length equals $l_x - W + 1$,
- first element of the output sequence corresponds to sample mean of $x[1:W]$,
- last element of the output sequence corresponds to sample mean of $x[(l_x - W + 1):l_x]$.

Value

numeric sequence

Examples

```
x <- rnorm(1000)
RunningMean(x, 100)
length(RunningMean(x, 100, circular = FALSE))
length(RunningMean(x, 100, circular = TRUE))
```

RunningSd*Fast Running Standard Deviation Computation*

Description

Computes running sample standard deviation of a sequence in a fixed width window. Uses convolution via Fast Fourier Transform.

Usage

```
RunningSd(x, W, circular = FALSE)
```

Arguments

x	numeric sequence
W	numeric; width of x sequence window
circular	logical; whether running sample standard deviation is computed assuming circular nature of x sequence (see Details)

Details

Parameter `circular` determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, W is a fixed length of x sequence window.

If `circular` equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample standard deviation of $x[1:W]$,
- last element of the output sequence corresponds to sample standard deviation of $c(x[l_x], x[1:(W - 1)])$.

If `circular` equals FALSE then

- output sequence length equals $l_x - W + 1$,
- first element of the output sequence corresponds to sample standard deviation of $x[1:W]$,
- last element of the output sequence corresponds to sample standard deviation of $x[(l_x - W + 1):l_x]$.

Value

numeric sequence

Examples

```
x <- rnorm(1000)
RunningSd(x, 100)
length(RunningSd(x, 100, circular = FALSE))
length(RunningSd(x, 100, circular = TRUE))
```

RunningVar

*Fast Running Variance Computation***Description**

Computes running sample variance of a sequence in a fixed width window. Uses convolution via Fast Fourier Transform.

Usage

```
RunningVar(x, W, circular = FALSE)
```

Arguments

x	numeric sequence
W	numeric; width of x sequence window
circular	logical; whether running sample variance is computed assuming circular nature of x sequence (see Details)

Details

Parameter circular determines whether x sequence is assumed to have a circular nature. Assume l_x is the length of sequence x, W is a fixed length of x sequence window.

If circular equals TRUE then

- output sequence length equals l_x ,
- first element of the output sequence corresponds to sample variance of $x[1:W]$,
- last element of the output sequence corresponds to sample variance of $c(x[1_x], x[1:(W - 1)])$.

If circular equals FALSE then

- output sequence length equals $l_x - W + 1$,
- first element of the output sequence corresponds to sample variance of $x[1:W]$,
- last element of the output sequence corresponds to sample variance of $x[(1_x - W + 1):1_x]$.

Value

numeric sequence

Examples

```
x <- rnorm(1000)
RunningVar(x, 100)
length(RunningVar(x, 100, circular = FALSE))
length(RunningVar(x, 100, circular = TRUE))
```

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