Package: rolleigen (via r-universe)

August 19, 2024

Type Package
Title Rolling Eigenanalysis
Version 1.0
Date YYYY-MM-DD
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Description Fast and efficient computation of rolling and expanding eigenanalysis for time-series data.
License GPL (>= 2)
URL https://github.com/jasonjfoster/rolleigen
<pre>BugReports https://github.com/jasonjfoster/rolleigen/issues</pre>
Imports Rcpp, RcppParallel
LinkingTo Rcpp, RcppArmadillo, RcppParallel, roll (>= 1.1.7)
SystemRequirements GNU make
Roxygen list(old_usage = TRUE)
RoxygenNote 7.3.1
Encoding UTF-8
Suggests covr, testthat, zoo, pls, roll
Repository https://jasonjfoster.r-universe.dev
RemoteUrl https://github.com/jasonjfoster/rolleigen
RemoteRef HEAD
RemoteSha f4d3eb40742503c08210e8521bc791a41ea6d23c
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2 roll_eigen

Description

Fast and efficient computation of rolling and expanding eigenanalysis for time-series data.

Details

rolleigen is a package that provides fast and efficient computation of rolling and expanding eigenaalysis for time-series data.

Author(s)

Jason Foster

roll_eigen Rolling Eigenvalues and Eigenvectors	
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Description

A function for computing the rolling and expanding eigenvalues and eigenvectors of time-series data

Usage

```
roll_eigen(x, width, weights = rep(1, width), center = TRUE,
    scale = FALSE, order = TRUE, min_obs = width, complete_obs = TRUE,
    na_restore = FALSE, online = TRUE)
```

Arguments

Х	vector or matrix. Rows are observations and columns are variables.
width	integer. Window size.
weights	vector. Weights for each observation within a window.
center	logical. If TRUE then the weighted mean of each variable is used, if FALSE then zero is used.
scale	logical. If TRUE then the weighted standard deviation of each variable is used, if FALSE then no scaling is done.
order	logical. Change sign and order of the components.
min_obs	integer. Minimum number of observations required to have a value within a window, otherwise result is NA.
complete_obs	logical. If TRUE then rows containing any missing values are removed, if FALSE then pairwise is used.
na_restore	logical. Should missing values be restored?
online	logical. Process observations using an online algorithm.

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Value

A list containing the following components:

values An object of the same class and dimension as x with the rolling and expanding

eigenvalues.

vectors A cube with each slice the rolling and expanding eigenvectors.

Examples

```
n <- 15
m <- 3
x <- matrix(rnorm(n * m), nrow = n, ncol = m)
weights <- 0.9 ^ (n:1)

# rolling eigenvalues and eigenvectors with complete windows
roll_eigen(x, width = 5)

# rolling eigenvalues and eigenvectors with partial windows
roll_eigen(x, width = 5, min_obs = 1)

# expanding eigenvalues and eigenvectors with partial windows
roll_eigen(x, width = n, min_obs = 1)

# expanding eigenvalues and eigenvectors with partial windows and weights
roll_eigen(x, width = n, min_obs = 1, weights = weights)</pre>
```

roll_pcr

Rolling Principal Component Regressions

Description

A function for computing the rolling and expanding principal component regressions of time-series data.

Usage

```
roll_pcr(x, y, width, n_comps = ncol(x), weights = rep(1, width),
  intercept = TRUE, center = TRUE, scale = FALSE, min_obs = width,
  complete_obs = TRUE, na_restore = FALSE, online = TRUE)
```

Arguments

X	vector or matrix Rows are observations and columns are the independent variables.
У	vector or matrix. Rows are observations and columns are the dependent variables

width integer. Window size.

roll_pcr

integer. Number of principal components. n_comps vector. Weights for each observation within a window. weights logical. Either TRUE to include or FALSE to remove the intercept. intercept center logical. If TRUE then the weighted mean of each variable is used, if FALSE then zero is used. logical. If TRUE then the weighted standard deviation of each variable is used, if scale FALSE then no scaling is done. min_obs integer. Minimum number of observations required to have a value within a window, otherwise result is NA. complete_obs logical. If TRUE then rows containing any missing values are removed, if FALSE then pairwise is used. na_restore logical. Should missing values be restored? logical. Process observations using an online algorithm. online

Value

A list containing the following components:

coefficients A list of objects with the rolling and expanding coefficients for each y. An object is the same class and dimension (with an added column for the intercept) as x.

r.squared A list of objects with the rolling and expanding r-squareds for each y. An object

is the same class as x.

Examples

```
n <- 15
m <- 3
x <- matrix(rnorm(n * m), nrow = n, ncol = m)
y <- rnorm(n)
weights <- 0.9 ^ (n:1)

# rolling regressions with complete windows
roll_pcr(x, y, width = 5, n_comps = 1)

# rolling regressions with partial windows
roll_pcr(x, y, width = 5, n_comps = 1, min_obs = 1)

# expanding regressions with partial windows
roll_pcr(x, y, width = n, n_comps = 1, min_obs = 1)

# expanding regressions with partial windows and weights
roll_pcr(x, y, width = n, n_comps = 1, min_obs = 1, weights = weights)</pre>
```

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