

Package ‘rshift’

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Type Package

Title Paleoecology Functions for Regime Shift Analysis

Version 3.0.0

Description Contains a variety of functions, based around regime shift analysis of paleoecological data.

Citations:

Rodionov() from Rodionov (2004) <[doi:10.1029/2004GL019448](https://doi.org/10.1029/2004GL019448)>

Lanzante() from Lanzante (1996) <[doi:10.1002/\(SICI\)1097-0088\(199611\)16:11%3C1197::AID-JOC89%3E3.0.CO;2-L](https://doi.org/10.1002/(SICI)1097-0088(199611)16:11%3C1197::AID-JOC89%3E3.0.CO;2-L)>

Hellinger_trans from Numerical Ecology, Legendre & Legendre (ISBN 9780444538680)

rolling_autoc from Liu, Gao & Wang (2018) <[doi:10.1016/j.scitotenv.2018.06.276](https://doi.org/10.1016/j.scitotenv.2018.06.276)>

Sample data sets lake_data & lake_RSI processed from Bush, Silman & Urrego (2004) <[doi:10.1126/science.1090795](https://doi.org/10.1126/science.1090795)>

Sample data set January_PDO from NOAA: <<https://www.ncei.noaa.gov/access/monitoring/pdo/>>.

Suggests R.rsp

VignetteBuilder R.rsp

Depends R (>= 3.5.0)

Imports grid, tibble, dplyr, ggplot2

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NeedsCompilation yes

SystemRequirements rustc & cargo if building from source

URL <https://github.com/alexhroom/rshift>

BugReports <https://github.com/alexhroom/rshift/issues>

Encoding UTF-8

LazyData true

RoxygenNote 7.2.3

Author Alex H. Room [aut, cre, cph] (<<https://orcid.org/0000-0002-5314-2331>>),
Felipe Franco-Gaviria [ctb, fnd]
(<<https://orcid.org/0000-0003-4799-1457>>),

Dunia H. Urrego [ctb, fnd] (<<https://orcid.org/0000-0001-7938-5529>>),
 The authors of the dependency Rust crates [ctb] (see inst/AUTHORS file
 for details)

Maintainer Alex H. Room <alex.room@btinternet.com>

Repository CRAN

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R topics documented:

absolute_to_percentage	2
Hellinger_trans	3
January_PDO	3
lake_data	4
lake_RSI	4
Lanzante	5
Rodionov	5
rolling_autoc	6
RSI_graph	7
rust_rodionov	7

Index	8
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absolute_to_percentage

*Converts absolute abundance data to a percentage of total abundance
 for each site*

Description

Converts absolute abundance data to a percentage of total abundance for each site

Usage

```
absolute_to_percentage(data, col, site)
```

Arguments

- data The dataframe to be used.
- col The column that change is being measured on.
- site The column containing the site of each sample.

Value

The ‘data’ dataframe with an added ‘percentage’ column.

<i>Hellinger_trans</i>	<i>Hellinger transform</i>
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Description

Hellinger transforms data (Legendre and Legendre, Numerical Ecology)

Usage

```
Hellinger_trans(data, col, site)
```

Arguments

<code>data</code>	The dataframe to be used.
<code>col</code>	The column that change is being measured on.
<code>site</code>	The column containing the site of each sample.

Value

The ‘data’ dataframe with an added ‘hellinger_trans_vals’ column.

<i>January_PDO</i>	<i>Pacific Decadal Oscillation in January</i>
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Description

A dataset containing January PDO values. Subset of the data from NOAA: <<https://www.ncei.noaa.gov/access/monitoring/pd>>

Usage

```
data(January_PDO)
```

Format

A data frame with 104 rows and 2 variables

Details

- PDO - Pacific Decadal Oscillation in January for the given year.
- Age - the year for which the PDO was measured.

lake_data*DCA-ordinated pollen data from Lake Consuelo*

Description

A dataset containing pre-processed DCA-ordinated data from Bush, Silman & Urrego (2004) <doi:10.1126/science.1090795>

Usage

```
data(lake_data)
```

Format

A data frame with 39 rows and 2 variables

Details

- DCA1 - DCA values for each timepoint from the raw dataset.
- Age - timepoint of each sample that has been DCA-ordinated.

lake_RSI*DCA-ordinated pollen data from Lake Consuelo with RSI values*

Description

A dataset containing pre-processed DCA-ordinated data from Bush, Silman & Urrego (2004) <doi:10.1126/science.1090795>
This data has been processed using Rodionov(lake_data, "DCA1", "Age", l=5, merge=TRUE)

Usage

```
data(lake_RSI)
```

Format

A data frame with 39 rows and 3 variables

Details

- DCA1 - DCA values for each timepoint from the raw dataset.
- Age - timepoint of each sample that has been DCA-ordinated.
- RSI - Regime Shift Index (see docs for Rodionov()) for each timepoint.

Lanzante

*Lanzante L-test***Description**

performs the L-method for detection of regime shifts (Lanzante, 1996)

Usage

```
Lanzante(data, col, time, p = 0.05, merge = FALSE)
```

Arguments

data	The dataframe to be used.
col	The column we are measuring change on.
time	The column containing time units (e.g. age of a subsample)
p	The largest p-value you want to check regime shifts for. Defaults to p = 0.05.
merge	Sets the result to be either a regime-shift only table (if FALSE), or an addition to the original table (if TRUE)

Value

If merge = FALSE (default), produces a 2-column table of time (the time value for each regime shift) and p (the p-value for each regime shift). If merge = TRUE, returns the original dataset with an extra p-value column, giving the p-value for each time unit - 0 for non-shift years.

Examples

```
Lanzante(lake_data, "DCA1", "Age")
Lanzante(lake_data, "DCA1", "Age", p=0.10, merge=TRUE)
```

Rodionov

*Rodionov (2004)'s STARS algorithm***Description**

performs STARS analysis (Rodionov, 2004) on a dataset

Usage

```
Rodionov(data, col, time, l, prob = 0.05, startrow = 1, merge = FALSE)
```

Arguments

<code>data</code>	The dataframe to be used.
<code>col</code>	The column we are measuring change on.
<code>time</code>	The column containing time units (e.g. age of a subsample)
<code>l</code>	The cut-off length of a regime; affects sensitivity (see Rodionov, 2004)
<code>prob</code>	The p-value for significance of a regime shift. Defaults to <code>p = 0.05</code> .
<code>startrow</code>	What row the analysis starts at. Defaults to 1.
<code>merge</code>	Sets the result to be either a regime-shift only table (if FALSE), or an addition to the original table (if TRUE)

Value

If `merge = FALSE` (default), produces a 2-column table of time (the time value for each regime shift) and RSI (the regime shift index for each regime shift). If `merge = TRUE`, returns the original dataset with an extra RSI column, giving the regime shift index for each time unit - 0 for non-shift years.

Examples

```
Rodionov(lake_data, "DCA1", "Age", l=5)
Rodionov(lake_data, "DCA1", "Age", l=5, prob=0.01, startrow=2, merge=TRUE)
```

<code>rolling_autoc</code>	<i>Rolling autocorrelation</i>
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Description

finds lag-1 autocorrelation in a rolling window; can be used to predict resilience (Liu, Gao, & Wang, 2018)

Usage

```
rolling_autoc(data, col, l)
```

Arguments

<code>data</code>	The dataframe that will be used.
<code>col</code>	The column we are measuring change on.
<code>l</code>	The time interval (no. of columns) used in the autocorrelation.

Value

A table of rolling lag-1 autocorrelation values.

<code>RSI_graph</code>	<i>Regime Shift Index graph</i>
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Description

creates two graphs, one of data and one of the RSI, as seen in Rodionov (2004)

Usage

```
RSI_graph(data, col, time, rsi)
```

Arguments

<code>data</code>	The dataframe that will be used.
<code>col</code>	The column we are measuring change on.
<code>time</code>	The column containing time units (e.g. age of a subsample)
<code>rsi</code>	The column containing RSI values (for best visualisation use Rodionov() with merge=TRUE)

Value

Two graphs, one on top of the other; one of col against time and one of RSI against time.

Examples

```
RSI_graph(lake_RSI, "DCA1", "Age", "RSI")
```

<code>rust_rodionov</code>	<i>Calculate STARS RSI points and return to R as a vector</i>
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Description

Calculate STARS RSI points and return to R as a vector

Usage

```
rust_rodionov(vals, t_crit, 1)
```

Arguments

<code>vals</code>	The column we are measuring change on
<code>t_crit</code>	The critical value of a t-distribution at the desired p-value
<code>1</code>	The cut-off length of a regime; affects sensitivity

Index

* datasets

January_PDO, [3](#)

lake_data, [4](#)

lake_RSI, [4](#)

absolute_to_percentage, [2](#)

Hellinger_trans, [3](#)

January_PDO, [3](#)

lake_data, [4](#)

lake_RSI, [4](#)

Lanzante, [5](#)

Rodionov, [5](#)

rolling_autoc, [6](#)

RSI_graph, [7](#)

rust_rodionov, [7](#)