

Package ‘TTAinterfaceTrendAnalysis’

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Type Package

Title Temporal Trend Analysis Graphical Interface

Version 1.5.8

Date 2021-10-06

Description This interface was created to develop a standard procedure to analyse temporal trend in the framework of the OSPAR convention. The analysis process run through 4 successive steps : 1) manipulate your data, 2) select the parameters you want to analyse, 3) build your regulated time series, 4) perform diagnosis and analysis and 5) read the results. Statistical analysis call other package function such as Kendall tests or cusum() function.

License GPL (>= 2)

Depends R (>= 3.3.0), base, stats, grDevices

Imports pastecs, reshape, e1071, relimp, multcomp, rkt, stlplus, nlme, tcltk, tcltk2, mvtnorm, zoo, methods

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TTAinterface-package *Interface Package for Temporal Trend Analysis*

Description

A friendly interface to perform Temporal Trend Analyses (MannKendall tests). Just follow the successive step from the data formatting to the results sorting.

Details

Package: TTAinterface
 Type: Package
 Version: 1.5.7
 Date: 2021-05-27
 License: GPL (>=2)

Author(s)

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References

Devreker, D. and Lefebvre, A. (2014), TTAinterfaceTrendAnalysis: An R GUI for routine temporal trend analysis and diagnostics. *Journal of Oceanography, Research and Data*, 1(7), 1-18.

about *About !*

Description

Display logo, version and developers name and email of the package.

Usage

about()

Details

Display logo, version and developers name of the package.

Envir

A temporary environment to stock data and objects

Description

The function create an environment where the data, arguments and objects that are used between the differents function of the package will be stock for better exchange processes.

Usage

Envir()

Details

Objects passed through the environment 'Envir' are called in the other function as Envir\$objects

fixdata

Fixdata function

Description

Simply modify your datase through the interface

Usage

fixdata()

Value

The edited database that is automaticaly read by the interface to replace former values

Note

fixdata() call the function fix() that act on the rawdata base. The fix() function itself call the function edit() from the package utils

See Also

[fix edit](#)

 FULLoption

Main function

Description

This is the core function of the interface. It receive arguments from the interface (see the function <TTAinterface>) and build regularized time series , perform diagnostics and analyses.

Usage

```
FULLoption(param, depth=NULL, sal = NULL, site=NULL, rawdata="NO", select="NO",
resume.reg="NO", test.normality="NO", plotB = "NO", selectBox="ByYears", log.trans="NO",
plotZ="NO", datashow="NO", help.timestep = "NO", auto.timestep = "NO", time.step = NULL,
help.aggreg = "NO", auto.aggreg = "NO", aggreg = NULL, mix = "YES", outliers.re = "NO",
na.replace="NO", start = NULL, end = NULL, months = c(1:12), norm = "NO", npsu = 30,
test.on.remaider = "NO", autocorr = "NO", spectrum="NO", anomaly="NO", a.barplot="NO",
zsmooth="NO", local.trend = "NO", test= "MK")
```

Arguments

param	The name of the parameter you want to analyse it must be the name of the column where are your data. Have to be enter like this : "yourparam".
depth	If existing, the depth interval where your data will be analyse. If values are different from depth max and depth min, missing value are exclude Depth column must be name as 'DEPTH'. Enter the value like this : c(a,b). For analysis at one specific depth you can enter c(a,a).
sal	Same thing as for the depth Salinity column must be name as 'S'.
site	Labels of sampling site as they appears in the database Enter the value like this c("S1", "S2").
rawdata	Peform descriptive statistics on raw database, can be "YES" or "NO" (the default).
select	Peform descriptive statistics on selected parameter and site, can be "YES" or "NO" (the default).
resume.reg	Peform descriptive statistics on regularized time series, can be "YES" or "NO" (the default).
test.normality	Perform a Shapiro-Wilk normality test on selected parameter, can be "YES" or "NO" (the default).
plotB	Display a boxplot of rawdata with outliers identified as cirle, can be "YES" or "NO" (the default).
selectBox	Options for plotB: allow to choose between boxplot by years or by months.
log.trans	This option transform your data in log(x+1) prior to perform analysis.
plotZ	Display a plot of the regularized time series, can be "YES" or "NO" (the default).
datashow	Show a table of the regularized data, can be "YES" or "NO" (the default).

help.timestep	Display an advice for time step selection, base on the mean time that separate two successive measurements. Can be "YES" or "NO" (the default).
auto.timestep	Autoexecute the advice without display it.
time.step	Choice of the time step for data aggregation during the build of the time series, can be "Fortnight", "Semi-fortnight", "Mensual", "Annual" or "Mono-mensual" for an aggregation of the data of a month of all years (e.g. all January data).
help.aggreg	Display an advice for method of aggregation selection, base on Wilcoxon p.value between rawdata and the different method. Can be "YES" or "NO" (the default).
auto.aggreg	Autoexecute the advice without display it.
aggreg	Choice of the method of aggregation during the build of the time series, can be "Mean", "Median", "Max" for maximal value selection or "Quantile" for selection of the quantile 90
mix	Allow to mix the data of all sampling site during analysis. Permanently set to "YES" and removed from the GUI since version 1.5.
outliers.re	Remove the outliers from the rawdata, the outliers list is save in a .csv file. (for outliers visual identification see boxplot section).
na.replace	Replace missing values with median of the corresponding cycle (week, month...) for lags longer than 3 days and linear regression for shorter missed period. Can be "YES" or "NO" (the default).
start	Define the first year of data analysis (by default the first of the database).
end	Define the last year of data analysis (by default the last of the database).
months	Define the months of data analysis (by default the twelve months).
norm	Compute normalised values of nutrients at the salinity npsu for each years, can be "YES" or "NO" (the default).
npsu	Compute normalised values of nutrients at the salinity npsu for each years, 30 by default.
test.on.remaider	Extract the reminders from the data series using the stl package functions to perform statistical analysis.
autocorr	Display the autocorrelation diagramme of the regularized time series using the acf function with arguments : lag.max = ((nrow(TimeSerie))/2), na.action = na.pass. Can be "YES" or "NO" (the default)
spectrum	Display the Fourier spectrum of the regularized time series using a Smoothed Periodogram (spec.pgram). Can be "YES" or "NO" (the default).
anomaly	Display a color box (function filled.contour) plot by year each time.step (months or weeks) minus the mean of the time.step of all years. Red colors show positive anomalies and blue colors negative anomalies. Can be "YES" or "NO" (the default).
a.barplot	Display an anomaly barplot as a function of the time.step. Red colors show positive anomalies and blue colors negative anomalies. Can be "YES" or "NO" (the default).
zsmooth	Display a detrended plot of the time series using the stl function with arguments s.window="periodic", na.action=na.fail. Can be "YES" or "NO" (the default).

<code>local.trend</code>	Display the interactive cusum plot of the time series (<code>local.trend</code> of the <code>pastecs</code> package) that allow to manually identify the period of change in the tendency using the function <code>identify</code> and perform a Kendall family test on each identified period (see <code>test</code> section). Can be "YES" or "NO" (the default).
<code>test</code>	Perform a test to evaluate the presence and the magnitude of a temporal trend on the time series. Can be "MK" for Seasonal Mann-Kendall test (the default), "SMK" for the same test with detail for each time step, "LOESS" that fit a polynomial surface determined by one or more numerical predictors, using local fitting; a MK is perform on this fitting.

Value

Results are return as `.png` figures or `.txt` files Results are also directly readable directly in the right part of the interface.

`Savepath` can be choose using the option 'Select directory' (see the function `selectdirectory` more more informations)

Name of saved filed follow the nomenclature : `Original.file.name_analysis.name_parameter.txt/.png`

or for multiple period analysis (see `cusum` for more details) : `Original.file.name_analysis.name_parameter_starting.year_endi analysis.names` are :

`_Boxplot_` for boxplot figure (`.png`). `_Outliers_` for the save of removed outliers (`.txt`). `_TimeSeries_` for the plot of the regularized time series (`.png`). `_Regularised_data_` for the table of regularized time series (`.txt`). `_Autocor_` for the autocorelation diagram (`.png`). `_Spectrum_` for the Fourier spectrum plot (`.png`) . `_ColorPlot_` for the anomaly color.plot (`.png`). `_Anomaly BarPlot_` for the anomaly barplot (`.png`) `_Detrended_` for detrended plot (`.png`). `_Local_Global Trend_` for result of Seasonal Mann Kendall apply to local trend (`.txt`). `_Local_Seasonal Trend_` same as above with detail for each time step (`.txt`). `_Global Trend_` for result of Seasonal Mann Kendall (`.txt`). `_Seasonal Trend_` same as above with detail for each time step (`.txt`). `_LOESSplot_` for loess plot (`.png`). `_NormalNutri_` for analysis of normalized values of nutrients (`.png`).

See values output of corresponding functions.

Author(s)

David Devreker

See Also

[boxplot](#) [impute](#) [shapiro.test](#) [summary](#) [acf](#) [spectrum](#) [filled.contour](#) [stl](#) [local.trend](#) [mannKen](#) [seasonTrend](#) [seaKen](#) [loess](#)

interpTs

Interpolate or substitute missing time series values (code of the former wq package)

Description

Imperpolates or substitutes missing data in a time series for gaps up to a specified size.

Usage

```
interpTs(x, type = c("linear", "series.median", "series.mean", "cycle.median",
"cycle.mean"), gap = NULL)
```

Arguments

x	object of class "ts" or "mts"
type	method of interpolation or substitution
gap	maximum gap to be replaced

Value

The time series with some or all missing values replaced.

Author(s)

Alan D. Jassby and James E. Cloern

mannKen	<i>Mann-Kendall trend test and the Sen slope (code modified from the former wq package)</i>
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Description

Applies Kendall's tau test for the significance of a monotonic time series trend. Also calculates the Sen slope as an estimate of this trend.

Usage

```
mannKen(x, ...)
```

Arguments

x	A numeric vector, matrix or data frame
...	Other arguments to pass to plotting function

Value

A list of the following if x is a vector:

sen.slope.Sen	slope
sen.slope.rel	Relative Sen slope
p.value	Significance of slope
S	Kendall's S
varS	Variance of S
miss	Fraction of missing slopes connecting first and last fifths of x or a matrix with corresponding columns if x is a matrix or data frame.

Author(s)

Alan D. Jassby and James E. Cloern

See Also

[seaKen seasonTrend](#)

seaKen	<i>Seasonal and Regional Kendall trend test (code modified from the former wq package)</i>
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Description

Calculates the Seasonal or Regional Kendall test of trend significance, including an estimate of the Sen slope.

Usage

```
seaKen(x, ...)
```

Arguments

x	A numeric vector, matrix or data frame made up of seasonal time series
...	Other arguments to pass to plotting function

Value

A list of the following if x is a vector: seaKen returns a list with the following members:

sen.slope	Sen slope
sen.slope.pct	Sen slope as percent of mean
p.value	significance of slope
miss	for each season, the fraction missing of slopes connecting first and last 20 percent of the years or a matrix with corresponding columns if x is a matrix or data frame.

Author(s)

Alan D. Jassby and James E. Cloern

See Also

[mannKen](#)

seasonTrend	<i>Determine seasonal trends (code modified from the former wq package)</i>
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Description

Finds the trend for each season and each variable in a time series.

Usage

```
seasonTrend(x, ...)
```

Arguments

x	Time series vector, or time series matrix with column names
...	Further options to pass to plotting function

Value

A data frame with the following fields:

series	series names
season	season number
sen.slope	Sen slope in original units per year
sen.slope.rel	Sen slope divided by median for that specific season and series
p	p-value for the trend according to the Mann-Kendall test.
missing	Proportion of slopes joining first and last fifths of the data that are missing

Author(s)

Alan D. Jassby and James E. Cloern

See Also

[mannKen](#)

selectdirectory	<i>Saved path selection</i>
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Description

Allow to chose the directory where results (.txt and .png files) will be saved.

Usage

```
selectdirectory()
```

Details

It select the main save directory; the package will create appropriate sub-folder as function of selected parameters, statsitics, methods etc. Then you will be able to perform successive analyses wihtout overwriting the previous results.

SRNDunkerque	<i>Coastal survey near the Gravelines power plant form 1995 to 2010</i>
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Description

Variation in temperature, salinity and chlorophyll-a concentration (microg/l) monthly measured between 1995 and 2010 at three different stations distributed onshore to offshore (North See) near the city of Dunkerque (north of France) for the SRN monitoring program (Ifremer). This database contain many missing values.

Format

A data.frame (TXT) containing 1561 measurments of temperature, salinity and chlorophyll-a concentration

Source

The Ifremer QUADRIGE_2 meta-database

TTAinterface

Graphic Interface For Temporal Trend Analysis

Description

A friendly user graphic interface to perform temporal trend analysis. The interface offer multiple options to select parameters and build time series that the user can follow step by step. Some options are selected by default to let the hurry user to do really quick analysis. Some diagnostic tools are also present.

Usage

TTAinterface()

Value

Results are saved in .txt files or .png figures in the desire directory (see selectdirectory). See 'FULLoption' values fore more details.

Author(s)

David Devreker

See Also

[FULLoption](#) [fixdata](#) [selectdirectory](#)

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