

Package ‘dendRoAnalyst’

July 22, 2025

Type Package

Date 2024-03-19

Title A Tool for Processing and Analyzing Dendrometer Data

Version 0.1.5

Maintainer Sugam Aryal <sugam.aryal@fau.de>

Description There are various functions for managing and cleaning data before the application of different approaches. This includes identifying and erasing sudden jumps in dendrometer data not related to environmental change, identifying the time gaps of recordings, and changing the temporal resolution of data to different frequencies. Furthermore, the package calculates daily statistics of dendrometer data, including the daily amplitude of tree growth. Various approaches can be applied to separate radial growth from daily cyclic shrinkage and expansion due to uptake and loss of stem water. In addition, it identifies periods of consecutive days with user-defined climatic conditions in daily meteorological data, then check what trees are doing during that period.

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.3.1

Depends R (>= 3.5.0), stats, base, tidyverse

Imports dplyr, ggplot2, lubridate, readxl, tibble, tidyr, zoo,
forecast, mgcv, minpack.lm, pspline, stringr

Suggests knitr, rmarkdown, testthat (>= 3.0.0)

VignetteBuilder knitr

NeedsCompilation no

Author Sugam Aryal [aut, cre, dtr],
Martin Häusser [aut],
Jussi Grießinger [aut],
Ze-Xin Fan [aut],
Achim Bräuning [aut, dgs]

Repository CRAN

Date/Publication 2024-03-19 16:20:05 UTC

Config/testthat/edition 3

Contents

clim.twd	2
daily.data	3
dendro.resample	4
dendro.truncate	5
dm.detrend	6
dm.fit.gompertz	7
dm.na.interpolation	7
gf_nepa17	8
i.jump.locator	9
jump.locator	9
ktm_rain17	10
mean_detrended.dm	11
mov.cor.dm	11
nepa	12
nepa17	13
nepa2	13
network.interpolation	14
phase.sc	15
phase.zg	16
plot_mov.cor	17
plot_SC_output	18
plot_ZG_output	19
read.dendrometer	20
twd.maxima	20
Index	22

clim.twd	<i>Calculating relative growth change during no-rain periods.</i>
----------	---

Description

This function calculates the number and the location of climatically adverse periods within a climate time series. The user can define a duration and threshold of these conditions. The function also provides the relative radial/circumferencial change during each adverse period for the original or normalized data. See Raffelsbauer et al., (2019) for more details.

Usage

```
clim.twd(  
  df,  
  Clim,  
  dailyValue = "max",  
  thresholdClim = "<10",  
  thresholdDays = ">5",  
  showPlot = TRUE  
)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
Clim	dataframe with the first column containing Date in yyyy-mm-dd and second column containing corresponding climate data.
dailyValue	either 'max', 'min', 'mean', or 'sum' for selecting the daily resampled value. Default is 'max'. See dendro.resample for details.
thresholdClim	string, the threshold for the respective climatic parameter. E.g. if climatic data is precipitation then days, where precipitation is below or equal to this value, are considered as adverse climate. Default is '<10'.
thresholdDays	string, the minimum number of consecutive adverse days to be considered for analysis. For example, thresholdDays=2 means the relative radial/circumferential change is calculated for adverse periods lasting for more than 2 days. Default is '>5'.
showPlot	logical, if TRUE, generates plots.

Value

A dataframe containing the respective periods, relative radial/circumference change for each tree, the ID for each period and their beginning and end.

References

Raffelsbauer V, Spann S, Peña K, Pucha-Cofrep D, Steppe K, Bräuning A (2019) Tree Circumference Changes and Species-Specific Growth Recovery After Extreme Dry Events in a Montane Rainforest in Southern Ecuador. *Front Plant Sci* 10:342. doi:10.3389/fpls.2019.00342

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
data(ktm_rain17)
relative_dry_growth<-clim.twd(df=gf_nepa17, Clim=ktm_rain17, dailyValue='max', showPlot=TRUE)
1

head(relative_dry_growth,10)
```

daily.data

*Calculation of daily statistics for dendrometer data***Description**

This function calculates various statistics of dendrometer data on a daily basis. The daily statistics includes the daily maximum and minimum with their corresponding times and daily amplitude (difference between daily maximum and minimum). See King et al. (2013) for details.

Usage

```
daily.data(df, TreeNum)
```

Arguments

df dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.

TreeNum numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in *df*.

Value

A dataframe with the daily statistics of the dendrometer data that contains:

Columns	Description
DATE	The day of year in "yyyy-mm-dd".
Min	The minimum value record for the corresponding day.
Time_min	The time when minimum value recorded for the corresponding day.
Max	The maximum value record for the corresponding day.
Time_max	The time when maximum value recorded for the corresponding day.
mean	The daily average value of the dendrometer reading.
median	The daily median value of the dendrometer reading.
amplitude	The difference between daily maximum and daily minimum.
Remarks	"*" if Time_max > Time_min otherwise "".

References

King G, Fonti P, Nievergelt D, Büntgen U, Frank D (2013) Climatic drivers of hourly to yearly tree radius variations along a 6°C natural warming gradient. *Agricultural and Forest Meteorology* 168:36–46. doi:10.1016/j.agrformet.2012.08.002

Examples

```
library(dendRoAnalyst)
data(nepa17)
daily_stats<-daily.data(df=nepa17, TreeNum=1)
head(daily_stats,10)
```

dendro.resample

Resampling temporal resolution of dendrometer and climate data

Description

This function is designed to change the temporal resolution of data. Depending on the objective, the user can define either maximum, minimum, or mean values to resample data in hourly, daily, weekly or monthly frequency.

Usage

```
dendro.resample(df, by, value)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS.
by	either <i>H</i> , <i>D</i> , <i>W</i> or <i>M</i> to resample data into hourly, daily, weekly or monthly resolution.
value	either <i>max</i> , <i>min</i> , <i>mean</i> or <i>sum</i> for the resampling value.

Value

Dataframe with resampled data.

Examples

```
library(dendRoAnalyst)
data(nepa17)
# To resample monthly with maximum value
resample_M<-dendro.resample(df=gf_nepa17, by='M', value='max')
head(resample_M,10)
```

dendro.truncate	<i>Truncation of the dendrometer data</i>
-----------------	---

Description

This function is helpful to truncate dendrometer data for a user-defined period.

Usage

```
dendro.truncate(df, CalYear, DOY)
```

Arguments

df	dataframe with the first column named date and time in the format yyyy-mm-dd HH:MM:SS.
CalYear	numerical value or array of two elements for the desired year of calculation.
DOY	numerical value or array of two elements representing the day of year. If we provide an array instead of a single value for CalYear and a single value for DOY, it truncates data from the DOY of the first CalYear to the same DOY of the second CalYear. Conversely, if we provide one value for CalYear and an array of two elements for DOY truncates the data form first DOY to second DOY within the same CalYear. Finally, if we provide an array with two values for both DOY and CalYear, it truncates data from the first DOY of the first CalYear to the second DOY of second CalYear.

Value

A dataframe with the truncated data for the defined periods.

Examples

```
library(dendRoAnalyst)
data(nepa)
#Extracting data from doy 20 to 50 in 2017.
trunc1<-dendro.truncate(df=nepa, CalYear=2017, DOY=c(20,50))
head(trunc1,10)
```

dm.detrend	<i>Detrend the dendrometer data</i>
------------	-------------------------------------

Description

This function detrends the dendrometer data either using first difference or using gam function from [mgcv](#) package or the Gompertz function.

Usage

```
dm.detrend(df, method = "gam", CalYear)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
method	either 'f_diff', 'gam' or 'gomp' indicating the method to detrend the dendrometer data column in <i>df</i> .
CalYear	numeric for year of calculation. If df has more than one year, assigning CalYear truncates the data of only that year.

Value

A dataframe with the detrended dendrometer series.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
detrended<-dm.detrend(df=nepa17, method='f_diff', CalYear=2017)
head(detrended,10)
```

dm.fit.gompertz	<i>Fitting gompertz function on annual dendrometer data</i>
-----------------	---

Description

This function models the annual growth of dendrometer data using gompertz function.

Usage

```
dm.fit.gompertz(df, CalYear, TreeNum, f_derivative = F)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
CalYear	numeric for year of calculation. If df has more than one year, assigning CalYear truncates the data of only that year.
TreeNum	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .
f_derivative	logical if yes returns first derivative of gompertz curve.

Value

A dataframe with the modelled dendrometer series.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
gomp_fitted<-dm.fit.gompertz(df=gf_nepa17, TreeNum = 1, CalYear=2017)
head(gomp_fitted,10)
```

dm.na.interpolation	<i>Detection and interpolation of missing values in dendrometer data.</i>
---------------------	---

Description

This function detects gap(s) in time series, inserts the missing rows based on the provided temporal resolution and assigns NA values to the corresponding value. If required the NA values can be replaced by spline interpolation using [na.spline](#) of the package **zoo** or seasonal interpolation considering the seasonality of the daily pattern using [na.interp](#) of the package **forecast**.

Usage

```
dm.na.interpolation(df, resolution, fill = FALSE, method = "spline")
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and following columns with dendrometer data for the same temporal resolution and time period.
resolution	integer, indicating the resolution of dendrometer data in minutes .
fill	logical, if TRUE it fills the NA values using spline interpolation. Default is FALSE.
method	string, 'spline' for the spline interpolation or 'seasonal' for the seasonal interpolation.

Value

A dataframe containing the dendrometer data including gaps filled with either NA or interpolated values.

Examples

```
library(dendRoAnalyst)
data(nepa17)
gf_nepa17<-dm.na.interpolation(df=nepa17, resolution=60)
head(gf_nepa17,10)
```

gf_nepa17

Dendrometer data of Kathmandu for 2017 with gap filled

Description

The dendrometer data from three Chir pine tree collected in hourly resolution for 2017.

Usage

```
gf_nepa17
```

Format

A data frame with 8760 rows and 3 variables:

Time datetime time of data recording

T2 double reading for first tree

T3 double reading for second tree

i.jump.locator	<i>Removing artefacts due to manual adjustments of dendrometers interactively</i>
----------------	---

Description

Dendrometers generally have limited memory capacity beyond which it stops recording. To keep the measurement ongoing, they should be adjusted periodically, which can cause positive or negative jumps in the data. This function locates these artefacts and interactively adjusts them one by one.

Usage

```
i.jump.locator(df, TreeNum, v)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
TreeNum	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .
v	numerical value which is considered as artefact. E.g. v=1 implies that if the difference to the consecutive data point is more than 1 or less than -1, it will be considered as an artefact.

Value

A dataframe containing jump-free dendrometer data.

jump.locator	<i>Removing artefacts due to manual adjustments of dendrometers automatically for more than one dendrometers</i>
--------------	--

Description

Dendrometers generally have limited memory capacity beyond which it stops recording. To keep the measurement ongoing, they should be adjusted periodically, which can cause positive or negative jumps in the data. This function locates these artefacts and adjusts them. Unlike in [i.jump.locator](#), it can handle dataset with more than one dendrometers.

Usage

```
jump.locator(df, v)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
v	numerical value which is considered as artefact. E.g. v=1 implies that if the difference to the consecutive data point is more than 1 or less than -1, it will be considered as an artefact.

Value

A dataframe containing jump-free dendrometer data.

Examples

```
library(dendRoAnalyst)
data(nepa)
jump_free_nepa<-jump.locator(df=nepa ,v=1)
head(jump_free_nepa,10)
```

ktm_rain17

Daily rainfall data of Kathmandu for 2017.

Description

This file contains daily rainfall data of Kathmandu. The source of this data is 'Government of Nepal, Department of Hydrology and Meteorology'.

Usage

```
ktm_rain17
```

Format

A data frame with 365 rows and 2 variables:

TIME Date in YYYY-MM-DD format.

rainfall double rainfall in millimeters

Source

<http://www.mfd.gov.np/city?id=31/>

mean_detrended.dm	<i>Calculate mean of detrended dendrometer data.</i>
-------------------	--

Description

This function calculate the mean detrended series of dendrometer data. It is usefull to make a single averaged detrended dendrometer series of a species in a site. Further, it provides option for removing first order autocorrelation from dendrometer series using `auto.arima` function of "forecast" package.

Usage

```
mean_detrended.dm(detrended.dm, ac1.remove = T, robust.mean = T)
```

Arguments

<code>detrended.dm</code>	dataframe output data frame of <code>dm.detrend</code> function.
<code>ac1.remove</code>	logical if TRUE removes first order autocorrelation.
<code>robust.mean</code>	logical if TRUE calculates robust mean.

Value

A data frame with the mean of detrended dendrometer series.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
detrended<-dm.detrend(df=nepa17, method='f_diff', CalYear=2017)
m_detre <- mean_detrended.dm(detrended)
head(m_detre,10)
```

<code>mov.cor.dm</code>	<i>Calculating running correlation between dendrometer data and daily climate.</i>
-------------------------	--

Description

This function calculates running correlation between dendrometer series and provided climate variables. Users can choose methods such as pearson, kendall and spearman, see `cor.test` for further information.

Usage

```
mov.cor.dm(df, Clim, TreeNum, win_size, cor_method = "pearson")
```

Arguments

<code>df</code>	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and the dendrometer data in following columns.
<code>Clim</code>	dataframe with the first column containing Date in yyyy-mm-dd and second column containing corresponding climate data.
<code>TreeNum</code>	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .
<code>win_size</code>	numerical, the running days windows of which the correlation is to be calculated. The minimum value is 18.
<code>cor_method</code>	string, method to be applied during correlation calculation. One of the following: pearson, kendall and spearman

Value

A list of dataframes containing the correlation and significant value between dendrometer series and provided climates.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
data(ktm_rain17)
out_corr<-mov.cor.dm(df=gf_nepa17, Clim=ktm_rain17, TreeNum=1, win_size=21)
head(out_corr[1],10)
```

nepa

Dendrometer data from Kathmandu

Description

Dendrometer data from three Chir pine trees collected in hourly resolution for 2 years.

Usage

```
nepa
```

Format

A data frame with 14534 rows and 3 variables:

Time datetime time of data recording

T2 double reading for first tree

T3 double reading for second tree

nepa17	<i>Dendrometer data of Kathmandu for 2017</i>
--------	---

Description

Dendrometer data from three Chir pine tree collected in hourly resolution for 2017.

Usage

nepa17

Format

A data frame with 8753 rows and 3 variables:

Time datetime time of data recording

T2 double reading for first tree

T3 double reading for second tree

nepa2	<i>Dendrometer data from Kathmandu version 2</i>
-------	--

Description

Dendrometer data from three Chir pine trees collected in hourly resolution for 2 years with separated time.

Usage

nepa2

Format

A data frame with 14534 rows and 8 variables:

year numeric year of data recording

month numeric months of data recording

day numeric days of data recording

hours numeric hours of data recording

minutes numeric minutes of data recording

seconds numeric seconds of data recording

T2 double reading for first tree

T3 double reading for second tree

network.interpolation *Interpolation of NA values using the dendrometer network*

Description

A function to interpolate the missing data of a dendrometer with the help of other dendrometers from the same site, provided they have the same measurement period and temporal resolution.

Usage

```
network.interpolation(df, referenceDF, niMethod)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS and dendrometer data in the second column and onward. The data gaps must be filled with NA using the gap.interpolation function.
referenceDF	dataframe with other dendrometers to be used as reference for the interpolation. The more dendrometers are included, the more robust will be the interpolation.
niMethod	string, either <i>'linear'</i> or <i>'proportional'</i> for interpolation method.

Value

A dataframe with NA values replaced by interpolated data.

Examples

```
library(dendRoAnalyst)
data("gf_nepa17")
df1<-gf_nepa17
# Creating an artificial reference dataset.
df2<-cbind(gf_nepa17,gf_nepa17[,2:3],gf_nepa17[,2:3])
colnames(df2) <- c('Time','T1','T2','T3','T4','T5','T6')
# Creating gaps in dataset by replacing some of the reading with NA in dataset.
df1[40:50,3]<-NA
# Using proportional interpolation method.
df1_NI<-network.interpolation(df=df1, referenceDF=df2, niMethod='proportional')
head(df1_NI,10)
```

phase.sc	<i>Application of the stem-cycle approach to calculate different phases, their duration and to plot them.</i>
----------	---

Description

This function analyses the dendrometer data using Stem-cycle approach (Downs et al. 1999; Deslauriers et al. 2011). A function that defines three phases: 1) Shrinkage, when the dendrometer reading is less than previous reading, 2) Expansion, when current reading is more than previous reading and 3) Increment, when current reading is higher than the previous maximum. Additionally, it calculates various statistics for each phase.

Usage

```
phase.sc(df, TreeNum, smoothing = NULL)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS. It should contain data with constant temporal resolution for best results.
TreeNum	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .
smoothing	numerical value from 1 to 12 which indicates the length of the smoothing spline, i.e. 1 = 1 hour and 12 = 12 hours. Default is NULL for no smoothing. The function smooth.Pspline is used for smoothing the data.

Value

A list of two dataframes. The first dataframe *SC_cycle* with cyclic phases along with various statistics and the second dataframe *SC_phase* with assigned phases for each data point. The dataframe *SC_cycle* contains the beginning, end, duration, magnitude and rate of each phase. The dataframe *SC_phase* contains time and corresponding phases during that time. The contents of *SC_cycle* are:

Columns	Description
Phase	Cyclic phases. 1, 2, and 3 for Shrinkage, Expansion, and Increment respectively.
start	Time when the corresponding phase starts.
end	Time when the corresponding phase ends.
Duration_h	Duration of the corresponding phase in hours.
Duration_m	Duration of the corresponding phase in minutes.
Magnitude	Radial/circumferential change during the corresponding phase in millimeters.
rate	Rate of Radial/circumferential change in micrometers per hour.
DOY	Day of year for the corresponding phase.

References

Deslauriers A, Rossi S, Turcotte A, Morin H, Krause C (2011) A three-step procedure in SAS to analyze the time series from automatic dendrometers. *Dendrochronologia* 29:151–161. doi:10.1016/j.dendro.2011.01.008

Downes G, Beadle C, Worledge D (1999) Daily stem growth patterns in irrigated *Eucalyptus globulus* and *E. nitens* in relation to climate. *Trees* 14:102–111. doi:10.1007/PL00009752

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
sc.phase<-phase.sc(df=gf_nepa17, TreeNum=1, smoothing=12)
head(sc.phase[[1]],10)
head(sc.phase[[2]],10)
```

phase.zg	<i>Application of the zero-growth approach to calculate different phases, their duration and to plot them.</i>
----------	--

Description

This function analyses data using the zero-growth approach. Initially, it divides the data in two categories: 1) Tree water deficiency (TWD), i.e. the reversible shrinkage and expansion of the tree stem when the current reading is below the previous maximum and, 2) Increment (GRO), the irreversible expansion of the stem when the current reading is above the previous maximum. Then it calculates the TWD for each data point as the difference between the modelled "growth line" and the observed measurement. See Zweifel et. al.,(2016) for details.

The severity value of each TWD was introduced in version 0.1.4 of the package.

Usage

```
phase.zg(df, TreeNum)
```

Arguments

df	dataframe with first column containing date and time in the format yyyy-mm-dd HH:MM:SS. It should contain data with constant temporal resolution for best results.
TreeNum	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .

Value

A list of two dataframes. The first dataframe *ZG_cycle* contains the cyclic phases along with various statistics and the second dataframe *ZG_phase* with assigned phases for each data point. The contents of *ZG_cycle* are:

Columns	Description
DOY	Day of year for the corresponding phase.
Phase	TWD for tree water deficit and GRO for irreversible expansion.
start	Time when the corresponding phase starts.
end	Time when the corresponding phase ends.
Duration_h	Duration of the corresponding phase in hours.
Magnitude	Radial/circumferential change in millimeters.
rate	Rate of Radial/circumferential change in micrometers per hour.
Max.twd	Maximum TWD recorded for the corresponding TWD phase.
twd.severity	The severity of the individual TWD period (see description below).
Max.twd.time	Time of occurrence of maximum TWD value for each TWD phase.
Avg.twd	Average of TWD values for each TWD phase.
STD.twd	Standard deviation of TWD values for each TWD phase.

References

Zweifel R, Haeni M, Buchmann N, Eugster W (2016) Are trees able to grow in periods of stem shrinkage? *New Phytol* 211:839–849. doi:10.1111/nph.13995

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
zg.phase<-phase.zg(df=gf_nepa17[1:600,], TreeNum=1)
head(zg.phase[[1]],10)
head(zg.phase[[2]],10)
```

plot_mov.cor

Plotting moving correlation with climate.

Description

This function plots the moving correlation i.e., output of `mov.cor.dm`.

Usage

```
plot_mov.cor(mov.cor.output, sig.only = T, ci = 0.95, clim_vars = "all")
```

Arguments

`mov.cor.output` list the output of `mov.cor.dm` function.
`sig.only` logical TRUE to plot only significant correlation.
`ci` numeric confidence interval
`clim_vars` array of climate variables or "all" for all climate variables.

Value

A plot with correlation.

Examples

```

library(dendRoAnalyst)
data(gf_nepa17)
data(ktm_rain17)
out_corr<-mov.cor.dm(df=gf_nepa17, Clim=ktm_rain17, TreeNum=1, win_size=21)
plot_mov.cor(mov.cor.output=out_corr, sig.only=TRUE, ci=0.95)

```

plot_SC_output	<i>Plotting output of stem cycle approach (i.e., output of phase.sc function).</i>
----------------	--

Description

This function plots the stem cycle of dendrometer data.

Usage

```

plot_SC_output(
  SC_output,
  DOY,
  Year,
  cols = c("#fee8c8", "#fdbb84", "#e34a33"),
  phNames = c("Shrinkage", "Expansion", "Increment")
)

```

Arguments

`SC_output` list the output of `phase.sc` function.
`DOY` array with initial and final day for plotting. E.g. $c(a,b)$, where a = initial date and b = final date.
`Year` array for indicating year for plotting.
`cols` array of three elements: colour for each phases.
`phNames` array with three elements for three different phases. Default is **"Shrinkage", "Expansion" and "Increment"**.

Value

A plot with different phases.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
sc.phase<-phase.sc(df=gf_nepa17, TreeNum=1, smoothing=12)
plot_SC_output(SC_output=sc.phase,D0Y=c(50,60), Year=2017)
```

plot_ZG_output	<i>Plotting output of zero-growth approach (i.e., output of phase.zg function).</i>
----------------	---

Description

This function plots the GRO and TWD of dendrometer data for a defined time period.

Usage

```
plot_ZG_output(ZG_output, D0Y, Year)
```

Arguments

ZG_output	list the output of phase.zg function.
D0Y	array with initial and final day for plotting. E.g. $c(a,b)$, where a = initial date and b = final date.
Year	array for indicating year for plotting.

Value

A plot with GRO and TWD in two different subplots.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
zg.phase<-phase.zg(df=gf_nepa17, TreeNum=1)
plot_ZG_output(ZG_output=zg.phase,D0Y=c(50,51), Year=2017)
```

read.dendrometer	<i>Reading dendrometer data.</i>
------------------	----------------------------------

Description

This function reads dendrometer data from .csv or .txt or .xlsx files. This function automatically recognizes the date time format of the first column of the data frame and changes it to "yyyy-mm-dd HH:MM:SS format".

Usage

```
read.dendrometer(file, sep = NULL, dec = NULL)
```

Arguments

file	string file name or path of the file.
sep	string the separator of the files. Only if they are different than the standard separators such as tab for .txt file and comma for .csv file.
dec	the character used in the file for decimal points.

Value

A dataframe with the dendrometer data:

twd.maxima	<i>Locating the maxima of TWD periods</i>
------------	---

Description

This function detects the TWD phases, including their beginning (TWDb), using the phase.zg function. Then it calculates the number, time of occurrence (Tm) and value of every local maximum within each TWD phase. In addition it calculates the time difference between 'TWDb' and each 'Tm' within each TWD phase.

Usage

```
twd.maxima(df, TreeNum, smoothing = 5)
```

Arguments

df	data frame with first column containing date and time in the format yyyy-mm-dd HH:MM:SS. It should contain data with constant temporal resolution for best results.
TreeNum	numerical value indicating the tree to be analysed. E.g. '1' refers to the first dendrometer data column in <i>df</i> .
smoothing	numerical value from 1 to 12 which indicates the length of the smoothing spline, i.e. 1 = 1 hour and 12 = 12 hours. Default is 5.

Value

A data frame with statistics of maxima in each TWD phase.

Examples

```
library(dendRoAnalyst)
data(gf_nepa17)
df1=gf_nepa17[2500:3500,]
twd_max<-twd.maxima(df=df1, TreeNum=2)
head(twd_max,10)
```

Index

* datasets

- gf_nepa17, [8](#)
- ktm_rain17, [10](#)
- nepa, [12](#)
- nepa17, [13](#)
- nepa2, [13](#)

clim.twd, [2](#)

daily.data, [3](#)

dendro.resample, [3](#), [4](#)

dendro.truncate, [5](#)

dm.detrend, [6](#)

dm.fit.gompertz, [7](#)

dm.na.interpolation, [7](#)

gf_nepa17, [8](#)

i.jump.locator, [9](#), [9](#)

jump.locator, [9](#)

ktm_rain17, [10](#)

mean_detrended.dm, [11](#)

mgcv, [6](#)

mov.cor.dm, [11](#)

na.interp, [7](#)

na.spline, [7](#)

nepa, [12](#)

nepa17, [13](#)

nepa2, [13](#)

network.interpolation, [14](#)

phase.sc, [15](#)

phase.zg, [16](#)

plot_mov.cor, [17](#)

plot_SC_output, [18](#)

plot_ZG_output, [19](#)

read.dendrometer, [20](#)

smooth.Pspline, [15](#)

twd.maxima, [20](#)