Package: boxfilter (via r-universe)

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Type Package
Title Filter Noisy Data
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Description Noise filter based on determining the proportion of neighboring points. A false point will be rejected if it has only few neighbors, but accepted if the proportion of neighbors in a rectangular frame is high. The size of the rectangular frame as well as the cut-off value, i.e. of a minimum proportion of neighbor-points, may be supplied or can be calculated automatically. Originally designed for the cleaning of heart rates, but suitable for filtering any slowly-changing physiological variable. For more information see Signer (2010)doi:10.1111/j.2041-210X.2009.00010.x.

License GPL (>= 3)

Encoding UTF-8

LazyData true

Imports ggplot2, gridExtra

Depends R (>= 3.50)

Suggests knitr, rmarkdown

VignetteBuilder knitr

Repository https://thomaspruf.r-universe.dev

RemoteUrl https://github.com/thomaspruf/boxfilter

RemoteRef HEAD

RemoteSha fbe9cf5da91776d6e165e1088f4698b4b0149dcd

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boxclip

Boxfilter Main Function

Description

Filters noise from data (e.g. heartrates) using x for x-axis data and y for y-axis, based on the proportion of neighbors of each point in a surrounding box of width and height (these may be determined automatically). It discards all data points that have less than a proportion of clipit neighbors.

Usage

Arguments

x	The x-axis of data, a datetime for example. Optional. If x=NULL x<-1:length(y) will be generated.
у	The y-axis of data, probably noisy. Required.
clipit	Optional. Y-values with less than a proportion of clipit neighbors will be discarded. If clipit is omitted it is set equal to the first trough in the neighbor proportion histogram.
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QI	Optional. An integer quality index for each data point.
width	Optional. The width of the box. If width is omitted it will be generated from $floor(length(x)*0.01)$.
height	Optional. The height of the box. If height is omitted it will be generated from floor(mean(y,na.rm=T)/4).
miny	The minimum y-value expected. Defaults to 10. Anything below miny is discarded.
plotit	Optional.If TRUE show a graph of the original and filtered data.
histo	Optional. If TRUE also show a histogram of the neighboring points.

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Details

Boxfilter mimics the human criterion of self-similarity. Data points with many neighbors are more trustworthy.

Value

x Original x-axis data y Original y-axis data

filtered Filtered data. Discarded data points are set to NA.

neighbors Proportion of neighbors of each point.

Note

To store only filtered data, use e.g.:

bc=boxclip (x,y) data=data.frame(x=bc\$x, hrf=bc\$filtered) data=na.omit(data) write.csv(data,file="myheartrates.csv")

Author(s)

Thomas Ruf (thomas.p.ruf@me.com)

See Also

clipview

```
data("wb_month")
data("ibex_hr")

x=wb_month$x
y=wb_month$hr

myclip=boxclip(x,y, histo=TRUE)
summary(myclip)

r=seq(1,28400,by=4)
myclip=boxclip(ibex_hr$Time[r],ibex_hr$Heartrate[r],0.65)
summary(myclip)
#store(myclip)

data("sleepduration")
Date=as.POSIXct(sleepduration$Date)
Duration=as.numeric(sleepduration$Bedtime)
boxclip(Date,Duration,miny=0)
```

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clipview	Get View Of Different clipit Values	
•	J JJ 1	

Description

Gives a view of the original data, a histogram of the neighbors, and results of four values of clipit. The cutoff-values clipit are determined by a sequence of length .-

Usage

```
clipview(x, y, clipit = NULL, width = NULL, height = NULL, miny=10)
```

Arguments

X	The x-axis of data, a datetime for example. Optional. If $x=NULL\ x<-1:length(y)$ will be generated.
У	The y-axis of data, probably noisy. Required.
clipit	This function requires a sequence of four items <1, e.g. clipit=seq $(0.1, 0.4, by=0.1)$
width	Optional. The width of the box. If width is omitted it will be generated from $floor(length(x)*0.01)$.
height	Optional. The height of the box. If height is omitted it will be generated from floor(mean(y,na.rm=T)/4).
miny	The minimum y-value expected. Anything below miny is discarded.
maxy	The minimum y-value expected. Anything abov miny is discarded.

Details

Note that a sequence is required for clipit here, while a scalar is required in boxclip().

Value

There is no return value. Six graphs are generated, original, histogram, and four for each value of clipit.

Author(s)

Thomas Ruf (thomas.p.ruf@me.com)

See Also

boxclip()

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Examples

```
data("wb_year")

r=seq(1,54179,by=5)

x=wb_year$x[r]
y=wb_year$y[r]

clipview(x,y,clipit=seq(0.2,0.5,0.1), miny=10)
```

ibex_hr

One year of heart rates of a capricorn free-living in the alps.

Description

Heart rates were obtained using acoustic loggers in the rumen.

Usage

```
data("ibex_hr")
```

Format

A data frame with 28454 observations on the following 2 variables.

Time datetime

Heartrate in bpm

References

Signer, C., Ruf, T., & Arnold, W. (2011). Hypometabolism and basking: The strategies of Alpine ibex to endure harsh over-wintering conditions. Functional Ecology, 25(3), 537-547.

```
data(ibex_hr)
```

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showdata

Boxfilter function

Description

Shows a graph of the data and its change over time. Called by boxclip()

Usage

```
showdata(x,y)
```

Arguments

The x-axis of data, a datetime for example.

y The y-axis of data, probably noisy.

Details

Asks the user whether to continue or not. Continue only if you want points to be deleted

Author(s)

Thomas Ruf (thomas.p.ruf@me.com)

Examples

```
set.seed(1234)
y=runif(1000,20,30)
ix=sample(1:1000,50)
y[ix]=runif(50,60,70)
showdata(1:1000,y)
```

store

Store data

Description

Stores original and filtered data.

Usage

```
store(object)
```

Arguments

object

Must be of class "boxclip", resulting from boxclip.

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Value

returns nothing

Author(s)

Thomas Ruf (thomas.p.ruf@me.com)

Examples

```
data ("wb_month.RData")
x=wb_month$x
y=wb_month$hr

myclip=boxclip(x,y,clipit=0.25,width=15)
summary(myclip)
store(myclip)
```

summary.boxclip

Summarize boxclip results

Description

Summary method for class boxclip.

Usage

```
## S3 method for class 'boxclip'
summary(object,...)
```

Arguments

object an object of class boxclip

... currently, no other arguments are required.

Value

summary.boxclip prints the following items:

clipit Cut-off value. Only points with a proportion >= clipit will be retained.

width Rectangle width in x-units. Filter criterion is the proportion of data points inside

the rectangle.

height Rectangle height in y-units. Filter criterion is the proportion of data points inside

the rectangle.

full Number of original data.

remaining Number of data remaining.

rest Percentage remaining.

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Author(s)

```
Thomas Ruf <thomas.p.ruf@me.com>
```

References

```
Signer (2010) <doi: 10.1111/j.2041-210X.2009.00010.x>
```

See Also

```
boxclip
```

Examples

```
data ("wb_month.RData")
x=wb_month$x
y=wb_month$hr

myclip=boxclip(x,y,clipit=0.25,width=15)
summary(myclip)
```

wb_month

One month of heart rates and their quality in a wild boar. Quality was assessed by Star-Oddi, Island.

Description

Heart rates were obtained from DST centi- HRT, Star-Oddi, Gardabaer, Iceland.

Usage

```
data("wb_month")
```

Format

A data frame with 3720 observations on the following 3 variables.

```
x x-axis, datetime
hr y-axis, heart rate
QI quality index (0-3) of the signal
```

```
data(wb_month)
```

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wb_year

One year of heart rates of a wild boar female.

Description

Heart rates were obtained from DST centi- HRT, Star-Oddi, Gardabaer, Iceland.

Usage

```
data("wb_year")
```

Format

A data frame with 3720 observations on the following 3 variables.

```
x x-axis, datetime
```

y y-axis, heart rate

References

Ruf, T., Vetter, S. G., Painer, J., Stalder, G., & Bieber, C. (2021). Atypical for northern ungulates, energy metabolism is lowest during summer in female wild boars (Sus scrofa). Scientific Reports, 11(1), 1-12.

```
data(wb_year)
```

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