

Package ‘RchivalTag’

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Description A set of functions to generate, access and analyze standard data products from archival tagging data.

Depends R (>= 3.0.1)

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R topics documented:

bin_TempTS	2
classify_DayTime	3
combine_histos	5
get_DayTimeLimits	6
get_thermalstrat	7
hist_tad	10
hist_tat	14
image_TempDepthProfiles	17
interpolate_TempDepthProfiles	19
merge_histos	21
plot_geopos	24
plot_TS	25

RchivalTag	28
read_histos	29
read_PDT	31
ts2histos	33
unmerge_histos	34

Index	36
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bin_TempTS	<i>bin depth-temperature time series data</i>
------------	-----------------------------------------------

Description

bins depth-temperature time series data to a user-defined resolution, returning the minimum, maximum and average temperature recorded at each depth interval (bin) per sampling day. The output is comparable to that of [read_PDT](#).

Why binning temperature data?

In case of archival tagging data, depth-temperature time series data at a given day may consist of multiple temperature profiles of different signatures, depending on the animal's behaviour. Slight differences in temperature profiles could impede further analyses (e.g. the estimation of the thermocline depth), if just the average profile is applied. To avoid such problems, it is useful to calculate the average temperature at given depth intervals (bins) and thus smooth temperature profiles of a given period.

In addition, temperature at depth profiles can be interpolated and then visualized using functions [interpolate_TempDepthProfiles](#) and [image_TempDepthProfiles](#), respectively. This facilitates the analysis of temporal changes of temperature profiles, for instance, in relation to animal behaviour (e.g. diving behaviour).

Usage

```
bin_TempTS(ts, res=8)
```

Arguments

ts	a data.frame with columns date, Depth and Temperature
res	the depth interval at which temperatures should be binned.

Value

A [data.frame](#) with the columns date, MeanTemp, MinTemp, MaxTemp, bin and MeanPDT (the latter being the average of the min and maximum water temperatures). Additional columns, used to distinguish tags, may include Serial, DeployID and Ptt, depending on their availability in the original ts-data.frame.

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

See Also

[read_PDT](#), [interpolate_TempDepthProfiles](#), [image_TempDepthProfiles](#)

Examples

```
# #### example 1) run on time series data:
## step I) read sample time series data file:
DepthTempTS <- read.table(system.file("example_files/104659-Series.csv",
                                     package="RchivalTag"),header = TRUE,sep=',')
DepthTempTS$date <- as.Date(DepthTempTS$Day,"%d-%b-%Y")
head(DepthTempTS)
#
#
# ## step Ib) bin temperature data on 10m depth bins
# ##           to increase later estimate accuracy (see Bauer et al. 2015):
# # DepthTempTS_binned <- bin_TempTS(DepthTempTS,res=10)
#
# ## step II) interpolate average temperature fields (MeanTemp) from binned data:
# m <- interpolate_TempDepthProfiles(DepthTempTS)
# # m <- interpolate_PDTs(DepthTempTS_binned)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
# ## step IV) plot interpolated profiles:
# image_TempDepthProfiles(m$station.1)
```

classify_DayTime

Classifying the time period of the day

Description

Classifying the time period of the day based on the timing of sunrise, sunset (and twilight events) or alternatively, geolocation estimates, as specified in [get_DayTimeLimits](#), that allow their internal estimation during the function call.

Usage

```
classify_DayTime(pos, twilight.set="ast")
```

Arguments

pos	A data.frame pos with the columns sunrise, sunset, dawn.ast,/dawn.naut and dawn.ast/dawn.naut in POSIXct-format. Note that the expected twilight vector (suffix "ast" for astronomical dawn and dusks; vs suffix "naut" for nautical twilight events) is defined by the function's second argument twilight.set (see description below).
twilight.set	character string, indicating the type of twilight used for the long daytime classification: "ast" (default) for astronomical and "naut" for nautical twilight events with sun angles of 18 vs 12 below the horizon, respectively. Corresponding (expected) dawn and dusk vector names are dawn.ast & dusk.ast vs dawn.naut & dusk.naut).

Value

The input [data.frame](#) pos extended by the time vectors daytime and daytime.long. In the former case, "Day" and "Night" periods are distinguished. In the latter case, "Day", "Night", "Dawn" and "Dusk".

Author(s)

Robert K. Bauer

References

Meeus, J. (1991) Astronomical Algorithms. Willmann-Bell, Inc.

See Also

[sunriset](#), [crepuscule](#), [get_DayTimeLimits](#)

Examples

```
#### example 1) estimate current times of dawn, sunrise, dusk and sunset in Mainz, Germany:
pos <- data.frame(Lat=8.2667, Lon=50)
pos$date.long <- strptime(Sys.Date(), "%Y-%m-%d")
get_DayTimeLimits(pos)

#### example 1b) classify current ime of the day in Mainz, Germany:
classify_DayTime(get_DayTimeLimits(pos))
```

combine_histos*combine lists of TAD/TAT frequency data*

Description

This function allows to combine separate lists of TAD/TAT frequency data from archival tags (i.e. by **Wildlife Computers**). The function requires ungrouped/unmerged TAD/TAT lists to avoid merging duplicate records (e.g. multiple TAD/TAT lists from the same individual). However, grouped/merged lists with TAD/TAT from multiple individuals and even duplicate records can be provided, as the function includes an internal call of [unmerge_histos](#) to meet this requirement.

Usage

```
combine_histos(hist_list1, hist_list2)
```

Arguments

```
hist_list1, hist_list2
```

Two list-of-lists to be combined, each containing TAD and TAT frequency data and the corresponding bin_breaks from one or several archival tags.

Value

A list-of-lists of ungrouped/unmerged TAD and TAT frequency data.

```
$ TAD:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
$ TAT:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
..$ ID2 : List of 2
...
```

Author(s)

Robert K. Bauer

See Also

[unmerge_histos](#), [merge_histos](#), [hist_tad](#), [hist_tat](#)

Examples

```
## example 1) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv",package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv",package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
hist_dat_merged <- merge_histos(hist_dat_combined,force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
unmerge_histos(hist_dat_merged)
```

get_DayTimeLimits	<i>Estimating the timing of sunrise, sunset, astronomical and nautical twilight events</i>
-------------------	--------------------------------------------------------------------------------------------

Description

Estimating the timing of sunrise, sunset, astronomical and nautical twilight events in POSIXct-format based on geolocations and a similar time vector. The function is a simplified call of the [sunriset](#) and [crepuscule](#) functions of the [maptools](#)-package that are based on algorithms provided by the National Oceanic & Atmospheric Administration (NOAA).

Usage

```
get_DayTimeLimits(pos)
```

Arguments

pos	a data.frame with the columns <code>date.long</code> (a time vector in POSIXct-format), <code>Lon</code> and <code>Lat</code> .
-----	-------------------------------------------------------------------------------------------------------------------------------------------------

Value

The input [data.frame](#) pos extended by the time vectors sunrise, sunset, dawn.naut, dawn.ast, dusk.naut and dusk.ast.

Author(s)

Robert K. Bauer

References

Meeus, J. (1991) Astronomical Algorithms. Willmann-Bell, Inc.

See Also

[sunriset](#), [crepuscule](#), [classify_DayTime](#)

Examples

```
#### example 1) estimate current times of dawn, sunrise, dusk and sunset in Mainz, Germany:
pos <- data.frame(Lat=8.2667, Lon=50)
pos$date.long <- strptime(Sys.Date(), "%Y-%m-%d")
get_DayTimeLimits(pos)

#### example 1b) classify current ime of the day in Mainz, Germany:
classify_DayTime(get_DayTimeLimits(pos))
```

get_thermalstrat	<i>estimate thermal stratification indices</i>
------------------	------------------------------------------------

Description

estimates thermal stratification indices, including thermocline depth, gradient, mixed later depth and stratification index from daily temperature at depth profiles, as illustrated by Bauer et al. (2015) for archival tagging data.

Usage

```
get_thermalstrat(x, dz=20, strat_lim=100, na.rm=FALSE,
                 show_info=TRUE, Depth_res, all_info=FALSE)
```

Arguments

x	A list generated by interpolate_TempDepthProfiles , containing interpolated temperature at depth profiles and their corresponding date and depth vectors: \$ Data_Source.ID_key:List of 3 ..\$ Temperature_matrix: num ..\$ Depth : num ..\$ Date :Date ..\$ sm :data.frame\$ Date :chr\$ nrecs :int\$ Depths :chr
dz	size of the moving window in meters between which temperature values should be compared for the estimation of the thermocline gradient and depth (by default 20).
na.rm	whether interpolated temperature at depth profiles with missing values should be treated (default is FALSE).
strat_lim	up to which depth (in meters) temperature values should be considered for the estimation of the stratification index (by default 100).
Depth_res	numeric value, defining the depth resolution at which the temperature data should be interpolated.
show_info	whether the process of the function run should be indicated (by default TRUE).
all_info	whether the summary information of the input file should be generated in the output (by default FALSE).

Value

a [data.frame](#) composed of

Date a date vector (see input argument x)

maxDepth_interp the maximum depth (in meters) of a daily temperature at depth profile to which its interpolation is limited)

tgrad the maximum temperature gradient of all possible moving windows of size dz)

tcline the thermocline depth, defined as the average depth (of the depth range) of the moving window(s) with the maximum temperature gradient tgrad)

dz size of the moving window in meters between which temperature values were compared

mld mixed layer depth, defined as the average depth of the first moving window that meets maximum temperature gradient criterium)

mld_0.5 mixed layer depth, defined as as the depth at which $T = SST - 0.5$ degrees, the temperature criterion of Monterey and Levitus (1997).

mld_0.8 mixed layer depth, defined as the depth at which $T = SST - 0.8$ degrees, the temperature criterion of Kara et al. (2000, 2003).

strat_index stratification index, defined as the standard deviation of all interpolated temperature values up to the depth defined by the argument strat_lim

... optional columns to be taken from the sm data.frame of the input list (in case that all_info=TRUE)
 nrecs number of records of the non-interpolated daily temperature at depth profiles
 Depths unique depth records of the non-interpolated daily temperature at depth profiles,
 seperated by `;`

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

Kara, A. B., P. A. Rochford, and H. E. Hurlburt (2000). An optimal definition for ocean mixed layer depth. Journal of Geophysical Research, 105:16803-16821, doi: [10.1029/2000JC900072](https://doi.org/10.1029/2000JC900072)

Kara, A. B., P. A. Rochford, and H. E. Hurlburt (2003) Mixed layer depth variability over the global ocean. Journal of Geophysical Research, 108:3079, doi: [10.1029/2000JC000736](https://doi.org/10.1029/2000JC000736)

Monterey, G., and S. Levitus (1997) Seasonal variability of mixed layer depth for the world ocean. NOAA Atlas NESDIS 14, U. S. Govt. Printing Office.

See Also

[interpolate_TempDepthProfiles](#)

Examples

```
#### example 1) run on PDT file:
## step I) read sample PDT data file:
setwd(system.file("example_files",package="RchivalTag"))
PDT <- read_PDT("104659-PDTs.csv")
head(PDT)
#
# ## step II) interpolate average temperature fields (MeanPDT) from PDT file:
# m <- interpolate_PDTs(PDT)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
#
# #### example 2) run on time series data:
## step I) read sample time series data file:
DepthTempTS <- read.table(system.file("example_files/104659-Series.csv",
                                     package="RchivalTag"),header = TRUE,sep=',')
#
# DepthTempTS$date <- as.Date(DepthTempTS$Day,"%d-%b-%Y")
# head(DepthTempTS)
#
```

```
#
# ## step Ib) bin temperature data on 10m depth bins
# ##           to increase later estimate accuracy (see Bauer et al. 2015):
# # DepthTempTS_binned <- bin_TempTS(DepthTempTS,res=10)
#
# ## step II) interpolate average temperature fields (MeanTemp) from binned data:
# m <- interpolate_TempDepthProfiles(DepthTempTS)
# # m <- interpolate_PDTs(DepthTempTS_binned)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
```

hist_tad

Time-at-Depth histogram

Description

generates daily or back-to-back (e.g. Day-vs-Night-) Time-at-Depth histograms from binned depth or depth time series data

Usage

```
hist_tad(df,
  bin_breaks=NULL, bin_prefix="Bin",
  select_id, select_from='Ptt', aggregate_by='Ptt',
  date,
  main, xlab='Time at Depth (%)', ylab="Depth (m)", labeling=TRUE,
  xlim=c(0, 100), adaptive.xlim=FALSE,
  split_by=NULL, split_levels, xlab2=split_levels,
  ylab.side=2, ylab.line, ylab.font=1,
  xlab.side=3, xlab.line=2.5, xlab.font=1,
  xlab2.side=1, xlab2.line=1, xlab2.font=2,
  main.side=2, main.line=3.8, main.font=1,
  col=c("darkgrey", "white"),
  do_mid.ticks=TRUE, yaxis.pos=0,
  mars,
  plot_sd=TRUE, plot_nrec=TRUE, plot_ntags=TRUE,
  cex=1.2, cex.main=cex, cex.lab=cex, cex.inf=cex,
  return.sm=FALSE,
  subplot=FALSE, inside=FALSE, Type="TAD")
```

Arguments

<code>df</code>	dataframe that either contains depth time series data (as a vector "Depth") or several vectors of Time-at-Depth frequencies. In the latter case, column names composed of a common <code>bin_prefix</code> (default is "Bin.") hold the pre-binned Time-at-Depth frequencies whose depth limits are defined in <code>bin_breaks</code> .
<code>bin_breaks, bin_prefix</code>	<code>bin_breaks</code> is a numeric vector of depth bin breaks for the histogram data. In case of binned data (e.g. from standard wildlife computer histogram files), column names with a <code>bin_prefix</code> are expected to contain the preprocessed data (by default: Bin1, Bin2, Bin3, etc.). Alternatively, depth time series data will be directly converted using function <code>ts2histos</code> .
<code>select_id, select_from</code>	these arguments allow to take a direct subset of the input dataframe. <code>select_from</code> defines the vector whereas <code>select_id</code> defines the identification key(s) that should be selected.
<code>aggregate_by</code>	character vector defining the columns by which the tagging data should be aggregated. Should contain columns that identify tags (e.g. Serial, Ptt, DeployID) the date and/or day time period (to separate records from night, day, dawn and dusk see classify_DayTime). Default values are: date, Day and Ptt.
<code>date</code>	An optional vector to select depth data of a specified date/-range.
<code>main, xlab, ylab, labeling</code>	The titles for the plot, x- and y-axes to be plotted if <code>labeling</code> is set TRUE (default).
<code>xlim, adaptive.xlim</code>	a vector defining the limits (x1,x2) of the x-axis, by default <code>c(0,100)</code> . However, if <code>adaptive.xlim</code> is set TRUE, these limits will be overwritten, and the maximum value (<code>xlim[2]</code>) will be chosen from the histogram data.
<code>split_by</code>	Name of the logical vector by which TaD data should be splitted (e.g. <code>day.time</code>).
<code>split_levels, xlab2</code>	Character vector defining the name and order of the levels of the <code>split_by</code> vector (e.g. <code>c("Night", "Day")</code>) for <code>split_by</code> vector ' <code>day.time</code> '. The same groups are plotted as a second x-axis label if not defined otherwise (<code>xlab2=split_levels</code>).
<code>ylab.side, ylab.line, ylab.font</code>	side, line and font of second y-axis label.
<code>xlab.side, xlab.line, xlab.font</code>	side, line and font of first x-axis label.
<code>xlab2.side, xlab2.line, xlab2.font</code>	side, line and font of second x-axis labels.
<code>main.side, main.line, main.font</code>	side, line and font of plot title.
<code>col</code>	colours to be used for the TaD-histogram, by default 'grey' and 'white' (corresponding to the values of <code>split_by/split_levels</code>).
<code>do_mid.ticks</code>	whether centered tick-labels, indicating the depth range of histogram cells, shall be plotted (by default FALSE). Alternatively, tick labels will be indicated at the breakpoints of the histogram cells.

<code>yaxis.pos</code>	x-axis coordinate at which the y-axis should be plotted (by default <code>xlim[1]</code> , and thus 0).
<code>mars</code>	a numerical vector of the form <code>c(bottom, left, top, right)</code> , describing the number of margin lines to be specified on the each side of the plot. The default is <code>c(2.1, 4.1, 6.1, 2.1)</code> . In case that <code>do_mid.ticks</code> is TRUE margins are: <code>c(2.1, 8, 6.1, 2.1)</code> .
<code>plot_sd, plot_nrec , plot_ntags</code>	whether standard deviation bars, the number of records and tags shall be plotted (default is TRUE) inside the TaD/TaT histogram.
<code>cex, cex.main, cex.lab, cex.inf</code>	font size of the title (<code>cex.main</code>), x- and y-axes labels (<code>cex.lab</code>), and other labels, like the number of records (<code>cex.inf</code>).
<code>return.sm</code>	whether summary information of the TaD histograms, including the number of records per summary period, the relative frequencies per bin and corresponding standard deviation, should be plotted (default is TRUE).
<code>subplot, inside</code>	whether the TaD histogram is a subplot or an inner plot of a figure (default is FALSE). If subplot or inside are set TRUE, graphic margins will not be set by <code>hist_tad</code> . In case that inside is TRUE, no axis-labels and titles will be plotted.
Type	The Type of data to be plotted (TAD: Time-at-Depth histograms; TAT: Time-at-Temperature histograms)

Details

Time-at-Temperature (Tat) and Time-at-Depth (TaD) frequencies are a standard data product of archival tags (incl. tag models TDR-Mk9, PAT-Mk10 and miniPAT by [Wildlife Computers](#)) that allow to assess habitat preferences of tagged animals (see function [read_histos](#)). It can be likewise generated from transmitted or recovered time series data sets using function [ts2histos](#).

However, different depth and temperature bin breaks are often used during different deployment programs, which makes a later comparative analysis of TaT and TaD data difficult. For such cases, the function [combine_histos](#) and [merge_histos](#) can be applied to merge TaT and TaD frequencies based on common bin breaks of different tags.

The purpose of this function is the visualization of Time-at-Depth (TaD) histograms, whereas [hist_tad](#) is the related function for Time-at-Temperature (TaT) data.

Author(s)

Robert K. Bauer

See Also

[ts2histos](#), [combine_histos](#), [merge_histos](#), [hist_tat](#)

Examples

```
ts_file <- system.file("example_files/104659-Series.csv", package="RchivalTag")
ts_df <- read.table(ts_file, header = TRUE, sep = ",")
```

```

head(ts_df)

tad_breaks <- c(0, 2, 5, 10, 20, 50, 100, 200, 300, 400, 600, 2000)
tat_breaks <- c(10,12,15,17,18,19,20,21,22,23,24,27)

## example 1a) convert only DepthTS data to daily TAD frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks, do_mid.ticks = FALSE)

## convert 1b) only TemperatureTS data to daily TAT frequencies:
tat <- ts2histos(ts_df, tat_breaks = tat_breaks)
hist_tat(ts_df, bin_breaks = tat_breaks, do_mid.ticks = FALSE)
hist_tat(tat$TAT$merged, do_mid.ticks = FALSE)

## convert 1c) DepthTS & TemperatureTS data to daily TAD & TAT frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks, tat_breaks = tat_breaks)

## example 2) rebin daily TAD frequencies:
tad <- ts2histos(ts_df, tad_breaks = tad_breaks)
tad2 <- rebin_histos(hist_list = tad, tad_breaks = tad_breaks[c(1:3,6:12)])
par(mfrow=c(2,2))
hist_tad(tad, do_mid.ticks = FALSE) ## example for multiple individuals
hist_tad(tad$TAD$merged, do_mid.ticks = FALSE)
hist_tad(tad$TAD$merged, bin_breaks = tad_breaks[c(1:3,6:12)]) ## from inside hist_tad

## example 3) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv",package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv",package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
hist_dat_merged <- merge_histos(hist_dat_combined,force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
unmerge_histos(hist_dat_merged)

```

hist_tat	<i>Time-at-Temperature histogram</i>
----------	--------------------------------------

Description

generates daily or back-to-back (e.g. Day-vs-Night-) Time-at-Temperature histograms from binned Temperature or Temperature time series data

Usage

```
hist_tat(df,
         bin_breaks=NULL, bin_prefix="Bin",
         main, xlab="Time at Temperature (%)",
         ylab=expression(paste("Temperature (",degree,"C)")), labeling=TRUE,
         Type="TAT", ...)
```

Arguments

df	dataframe that either contains Temperature time series data (as a vector "Temperature") or several vectors of Time-at-Temperature frequencies. In the latter case, vector names are composed of a common bin_prefix (default is "tad."), followed by the upper Temperature limit (bin break).
bin_breaks, bin_prefix	bin_breaks is a numeric vector of depth bin breaks for the histogram data. In case of binned data (e.g. from standard wildlife computer histogram files), column names with a bin_prefix are expected to contain the preprocessed data (by default: Bin1, Bin2, Bin3, etc.). Alternatively, depth time series data will be directly converted using function ts2histos .
main, xlab, ylab, labeling	The titles for the plot, x- and y-axes to be plotted if labeling is set TRUE (default).
Type	The Type of data to be plotted (TAD: Time-at-Depth histograms; TAT: Time-at-Temperature histograms)
...	additional arguments to be passed:
select_id, select_from	these arguments allow to take a direct subset of the input dataframe. select_from defines the vector whereas select_id defines the identification key(s) that should be selected.
aggregate_by	character vector defining the columns by which the tagging data should be aggregated. Should contain columns that identify tags (e.g. Serial, Ptt, DeployID) the date and/or day time period (to separate records from night, day, dawn and dusk see classify_DayTime). Default values are: date, Day and Ptt.
date	An optional vector to select depth data of a specified date/-range.

- xlim, adaptive.xlim** a vector defining the limits (x1,x2) of the x-axis, by default `c(0,100)`. However, if `adaptive.xlim` is set `TRUE`, these limits will be overwritten, and the maximum value (`xlim[2]`) will be chosen from the histogram data.
- split_by** Name of the logical vector by which TaD data should be splitted (e.g. `day.time`).
- split_levels, xlab2** Character vector defining the name and order of the levels of the `split_by` vector (e.g. `c("Night", "Day")` for `split_by` vector `'day.time'`. The same groups are plotted as a second x-axis label if not defined otherwise (`xlab2=split_levels`).
- ylab.side, ylab.line, ylab.font** side, line and font of second y-axis label.
- xlab.side, xlab.line, xlab.font** side, line and font of first x-axis label.
- xlab2.side, xlab2.line, xlab2.font** side, line and font of second x-axis labels.
- main.side, main.line, main.font** side, line and font of plot title.
- col** colours to be used for the TaD-histogram, by default `'grey'` and `'white'` (corresponding to the values of `split_by/split_levels`).
- do_mid.ticks** whether centered tick-labels, indicating the depth range of histogram cells, shall be plotted (by default `FALSE`). Alternatively, tick labels will be indicated at the breakpoints of the histogram cells.
- yaxis.pos** x-axis coordinate at which the y-axis should be plotted (by default `xlim[1]`, and thus 0).
- mars** a numerical vector of the form `c(bottom, left, top, right)`, describing the number of margin lines to be specified on the each side of the plot. The default is `c(2.1, 4.1, 6.1, 2.1)`. In case that `do_mid.ticks` is `TRUE` margins are: `c(2.1, 8, 6.1, 2.1)`.
- plot_sd, plot_nrec, plot_ntags** whether standard deviation bars, the number of records and tags shall be plotted (default is `TRUE`) inside the TaD/TaT histogram.
- cex, cex.main, cex.lab, cex.inf** font size of the title (`cex.main`), x- and y-axes labels (`cex.lab`), and other labels, like the number of records (`cex.inf`).
- return.sm** whether summary information of the TaD histograms, including the number of records per summary period, the relative frequencies per bin and corresponding standard deviation, should be plotted (default is `TRUE`).
- subplot, inside** whether the TaD histogram is a subplot or an inner plot of a figure (default is `FALSE`). If `subplot` or `inside` are set `TRUE`, graphic margins will not be set by `hist_tat`. In case that `inside` is `TRUE`, no axis-labels and titles will be plotted.

Details

Time-at-Temperature (Tat) and Time-at-Depth (TaD) frequencies are a standard data product of archival tags (incl. tag models TDR-Mk9, PAT-Mk10 and miniPAT by [Wildlife Computers](#)) that allow to assess habitat preferences of tagged animals (see function [read_histos](#)). It can be likewise generated from transmitted or recovered time series data sets using function [ts2histos](#).

However, different depth and temperature bin breaks are often used during different deployment programs, which makes a later comparative analysis of TaT and TaT data difficult. For such cases,

the function [combine_histos](#) and [merge_histos](#) can be applied to merge TaT and TaD frequencies based on common bin breaks of different tags.

The purpose of this function is the visualization of Time-at-Temperature (TaT) histograms, whereas [hist_tad](#) is the related function for Time-at-Depth (TaD) data.

Author(s)

Robert K. Bauer

See Also

[ts2histos](#), [combine_histos](#), [merge_histos](#), [hist_tad](#)

Examples

```
ts_file <- system.file("example_files/104659-Series.csv",package="RchivalTag")
ts_df <- read.table(ts_file, header = TRUE, sep = ",")
head(ts_df)

tad_breaks <- c(0, 2, 5, 10, 20, 50, 100, 200, 300, 400, 600, 2000)
tat_breaks <- c(10,12,15,17,18,19,20,21,22,23,24,27)

## example 1a) convert only DepthTS data to daily TAD frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks, do_mid.ticks = FALSE)

## convert 1b) only TemperatureTS data to daily TAT frequencies:
tat <- ts2histos(ts_df, tat_breaks = tat_breaks)
hist_tat(ts_df, bin_breaks = tat_breaks, do_mid.ticks = FALSE)
hist_tat(tat$TAT$merged, do_mid.ticks = FALSE)

## convert 1c) DepthTS & TemperatureTS data to daily TAD & TAT frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks, tat_breaks = tat_breaks)

## example 2) rebin daily TAD frequencies:
tad <- ts2histos(ts_df, tad_breaks = tad_breaks)
tad2 <- rebin_histos(hist_list = tad, tad_breaks = tad_breaks[c(1:3,6:12)])
par(mfrow=c(2,2))
hist_tad(tad, do_mid.ticks = FALSE) ## example for multiple individuals
hist_tad(tad$TAD$merged, do_mid.ticks = FALSE)
hist_tad(tad$TAD$merged, bin_breaks = tad_breaks[c(1:3,6:12)]) ## from inside hist_tad

## example 3) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv",package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv",package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
```



```

hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
hist_dat_merged <- merge_histos(hist_dat_combined,force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
unmerge_histos(hist_dat_merged)

```

image_TempDepthProfiles

plots interpolated daily temperature at depth profiles

Description

plots interpolated daily temperature at depth profiles, thus facilitating the analysis of temporal changes of temperature profiles, for instance, in relation to animal behaviour (e.g. diving behaviour). See Bauer et al. (2015) for further examples.

Usage

```

image_TempDepthProfiles(x, main=NULL, xlab='Date', ylab="Depth (m)",
                        cb.xlab=expression(paste("Temperature (",degree,"C)")),
                        cex.cb.xlab=1, cex.cb.ticks=1,
                        xlim, ylim, zlim, pal="jet", only.months, month.line, mars, ...)

```

Arguments

x	A list , generated by interpolate_TempDepthProfiles or interpolate_PDTs , containing interpolated temperature at depth profiles and their corresponding date and interpolated depth values as well as a summary table with the original depth values and their number per day: \$ Temperature_matrix: num \$ Depth : num \$ Date :Date \$ sm :data.frame
main, xlab, ylab	the title, x- and y-axis labels to be plotted.
cb.xlab	character string indicating the x-axis label of the colorbar.

cex.cb.xlab, cex.cb.ticks
cex.cb.xlab: font size of the x-axis label of the colorbar (by default 1). *cex.cb.ticks*: font size of the x-axis tick labels of the colorbar (by default 1).

xlim, ylim, zlim
the x, y and z limits of the plot.

pal
color map to be plotted (default is 'jet'). See [cmap](#) for available color maps.

only.months, month.line
whether only mid-months shall be plotted as tick labels of the x-axis (by default FALSE for time ranges of less than 3 months (93 days)). In case, that only.months is set TRUE, month.line defines the line where the month labels shall be plotted.

mars
a numerical vector defining the plot margins c(bottom, left, top, right) (by default c(5,4,4,9)).

...
additional arguments to be passed to [set.colorbarp](#)

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

See Also

[read_PDT](#), [bin_TempTS](#), [get_thermalstrat](#), [image_TempDepthProfiles](#)

Examples

```
#### example 1) run on PDT file:
## step I) read sample PDT data file:
setwd(system.file("example_files", package="RchivalTag"))
PDT <- read_PDT("104659-PDTs.csv")
head(PDT)
#
# ## step II) interpolate average temperature fields (MeanPDT) from PDT file:
# m <- interpolate_PDTs(PDT)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
# ## step IV) plot interpolated profiles:
# image_TempDepthProfiles(m$station.1)
#
#
```

```

##### example 2) run on time series data:
## step I) read sample time series data file:
DepthTempTS <- read.table(system.file("example_files/104659-Series.csv",
                                     package="RchivalTag"),header = TRUE,sep=',')
#
DepthTempTS$date <- as.Date(DepthTempTS$Day,"%d-%b-%Y")
# head(DepthTempTS)
#
#
## step Ib) bin temperature data on 10m depth bins
##
# to increase later estimate accuracy (see Bauer et al. 2015):
DepthTempTS_binned <- bin_TempTS(DepthTempTS,res=10)
#
## step II) interpolate average temperature fields (MeanTemp) from binned data:
m <- interpolate_TempDepthProfiles(DepthTempTS)
# m <- interpolate_PDTs(DepthTempTS_binned)
# str(m)
# m$sm
#
## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
## step IV) plot interpolated profiles:
# image_TempDepthProfiles(m$station.1)

```

interpolate_TempDepthProfiles

interpolate daily temperature at depth profiles

Description

interpolates depth-temperature data and returns daily average temperature at depth profiles on a user-specified resolution (Depth_res).

Results are returned as a list containing the interpolated Temperature-matrix, and the corresponding date and depth values. Thus interpolated temperature at depth profiles can be visualized using function [image_TempDepthProfiles](#) and facilitates the analysis of temporal changes of temperature profiles, for instance, in relation to animal behaviour (e.g. diving behaviour).

Usage

```

interpolate_TempDepthProfiles(ts, Temp_field="Temperature", ID_key="Serial",
                              Depth_res=.5, show_info=TRUE, Data_Source='station')

interpolate_PDTs(ts, Temp_field="MeanPDT", ID_key="Serial", #return_as_matrix=FALSE,
                 Depth_res=.5, show_info=TRUE, Data_Source='station')

```

Arguments

ts, Temp_field, ID_key	ts is a data.frame with temperature at depth data. Required columns are Depth for the depth data and a column containing temperature data, whose name is defined by Temp_field. ID_key specifies the name of an optional column on which sampling stations or tags can be distinguished (by default Serial).
Depth_res	numeric value, defining the depth resolution at which the temperature data should be interpolated.
show_info	whether the sampling dates and ids of stations or tags, as defined by the columns date and ID_key, should be indicated during the interpolation process.
Data_Source	a character string, defining the data source (by default station).

Value

A list containing the interpolated temperature at depth profiles and their corresponding date and interpolated depth values as well as a summary table with the original depth values and their number per day:

```
$ Data_Source.ID_key:List of 4
..$ Temperature_matrix: num
..$ Depth : num
..$ Date :Date
..$ sm :data.frame
```

Please see the examples for further understanding.

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

See Also

[read_PDT](#), [bin_TempTS](#), [get_thermalstrat](#), [image_TempDepthProfiles](#)

Examples

```
#### example 1) run on PDT file:
## step I) read sample PDT data file:
setwd(system.file("example_files",package="RchivalTag"))
PDT <- read_PDT("104659-PDTs.csv")
head(PDT)
#
# ## step II) interpolate average temperature fields (MeanPDT) from PDT file:
```

```

# m <- interpolate_PDTs(PDT)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
# ## step IV) plot interpolated profiles:
# image_TempDepthProfiles(m$station.1)
#
#
# ##### example 2) run on time series data:
# ## step I) read sample time series data file:
# DepthTempTS <- read.table(system.file("example_files/104659-Series.csv",
#                                     package="RchivalTag"),header = TRUE,sep=',')
# DepthTempTS$date <- as.Date(DepthTempTS$Day,"%d-%b-%Y")
# head(DepthTempTS)
#
#
# ## step Ib) bin temperature data on 10m depth bins
# ##           to increase later estimate accuracy (see Bauer et al. 2015):
# DepthTempTS_binned <- bin_TempTS(DepthTempTS,res=10)
#
# ## step II) interpolate average temperature fields (MeanTemp) from binned data:
# m <- interpolate_TempDepthProfiles(DepthTempTS)
# m <- interpolate_PDTs(DepthTempTS_binned)
# str(m)
# m$sm
#
# ## step III) calculate thermal stratification indicators per day (and tag):
# get_thermalstrat(m, all_info = TRUE)
# get_thermalstrat(m, all_info = FALSE)
#
# ## step IV) plot interpolated profiles:
# image_TempDepthProfiles(m$station.1)

```

merge_histos

merge and/or rebin TAD/TAT-frequency data

Description

The joint analysis of archival tagging data from different tagging programs is often hampered by differences in the tags' setups, e.g. by the user-specified temporal resolution of time series data or the definition of summary data products. The latter particularly concerns different selected bin breaks of Time-at-Depth (TAD) and Time-at-Temperature (TAT) frequency data from archival tags by **Wildlife Computers**.

The purpose of this function is to allow:

- 1) a grouping of TAD and TAT data from multiple tags based on similar bin breaks (For this, run the function with default statements, i.e. `force_merge` is `FALSE`),
- 2) merging (rebinning) of TAD and TAT data from multiple tags based on the bin breaks that all tags have in common (To do so, run the function with `force_merge` set `TRUE`).
- 3) merging (rebinning) of TAD and TAT data from multiple tags based on new user-specified `tad_breaks` and/or `tat_breaks`. In this case, the `force_merge`-statements `TRUE` and `FALSE` will omit or separately group tags that do not share all user-specified bin breaks, respectively.

To combine of TAD/TAT data of several `hist_lists`, see [combine_histos](#).

To visualize Time-at-Temperature (TaT) and Time-at-Depth (TaD) data, please see [hist_tat](#) and [hist_tad](#), respectively.

Usage

```
merge_histos(hist_list, tad_breaks=NULL, tat_breaks=NULL, force_merge=FALSE)
rebin_histos(hist_list, tad_breaks=NULL, tat_breaks=NULL, force_merge=FALSE)
```

Arguments

<code>hist_list</code>	A list-of-lists containing the TAD and TAT frequency data and the corresponding <code>bin_breaks</code> from one or several tags.
<code>tad_breaks</code>	<p>a numeric vector defining the <code>bin_breaks</code> for the merging (rebinning) of the TAD frequency data.</p> <p>In case that the additional argument <code>force_merge</code> is set <code>TRUE</code>, only tags whose original TAD bin breaks included all of the user-specified <code>tad_breaks</code> will be merged in a single group ('merged') based on the new bin breaks, while other tags will be omitted in the output. By contrast, if <code>force_merge</code> is set <code>FALSE</code>, tags that do not contain all specified <code>tad_breaks</code> will be merged in separate groups (<code>group2</code>, <code>group3</code>, etc.), based on similar <code>bin_breaks</code>.</p>
<code>tat_breaks</code>	<p>a numeric vector defining the <code>bin_breaks</code> for the merging (rebinning) of the TAT frequency data.</p> <p>In case that the additional argument <code>force_merge</code> is set <code>TRUE</code>, only tags whose original TAT bin breaks included all of the user-specified <code>tat_breaks</code> will be merged in a single group ('merged') based on the new bin breaks, while other tags will be omitted in the output. By contrast, if <code>force_merge</code> is set <code>FALSE</code>, tags that do not contain all specified <code>tat_breaks</code> will be merged in separate groups (<code>group2</code>, <code>group3</code>, etc.), based on similar <code>bin_breaks</code>.</p>
<code>force_merge</code>	<p>If <code>FALSE</code> (default), groups of tags with similar TAD and TAT-<code>bin_breaks</code> will be combined (no merging on new bin breaks) and identifier labels renamed as <code>group1</code>, <code>group2</code>, etc.</p> <p>If set <code>TRUE</code>, TAD and TAT frequency data will be merged on user-specified <code>tad/tat_breaks</code> or, if those arguments are missing, on the <code>bin_breaks</code> that</p>

all tags have in common. In both latter cases, identifier labels will be renamed "merged".

Value

A list-of-lists of grouped or merged TAD and TAT frequency data.

```
$ TAD:List
..$ group1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
$ TAT:List
..$ group1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
..$ group2 : List of 2
...
```

Author(s)

Robert K. Bauer

See Also

[unmerge_histos](#), [combine_histos](#), [hist_tad](#)

Examples

```
## example 1) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv",package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv",package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
hist_dat_merged <- merge_histos(hist_dat_combined,force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
```

```
unmerge_histos(hist_dat_merged)
```

plot_geopos

plot geolocation estimates derived from archival tagging data

Description

In case that geolocations are provided by csv-files or data frames, line and scatter plots are implemented. In case of ncdf-files, generated by the [Wildlife Computers-data portal](#), are selected, surface probability maps are illustrated. The latter procedure is based on the R-code given in the [location processing user guide](#) by [Wildlife Computers](#).

Usage

```
plot_geopos(file, pos, xlim, ylim,
             prob_lim=.75, pal="jet", alpha=70, type="p", pch=19, add=FALSE, ...)
```

Arguments

file	path and file name of .csv or .nc-file.
pos	alternative data.frame containing horizontal position records (allowed column names are 'Most.Likely.Longitude', 'Longitude' or 'Lon' and 'Most.Likely.Latitude', 'Latitude' or 'Lat', respectively).
xlim, ylim	Numeric vector, defining the limits of the x and y-axes.
prob_lim	in case that a netcdf-file (.nc) is selected, the value defines the limit of the probability surfaces in % (By default 75%). Otherwise ignored.
pal	color map to be plotted in case of polygon (.nc-files) or scatter plots (default is the 'jet'-colormap). See cmap for pre-installed color maps. Note that tracking data with constant time steps is being assumed in the color assignment. To verify this, a data.frame containing the colors at each time steps will be returned for polygon and scatter plots.
alpha	transparency of polygons and dots to be plotted in percent (By default 70%).
type	character string giving the type of plot desired. The following values are possible, for details (By default "p" for points, but "l" for lines is also implemented).
pch	dot-type to be plotted if 'points' have been selected (By default '19' for solid dots).
add	whether the a the plot should be added to an existent figure (default is FALSE)
...	additional arguments to be passed to plot .

Author(s)

Robert K. Bauer

See Also

[plotmap](#), [plot_DepthTS](#), [hist_tat](#), [hist_tad](#)

Examples

```
## example 1a) line plot from csv-file:
csv_file <- system.file("example_files/15P1019-104659-1-GPE3.csv",package="RchivalTag")
plot_geopos(csv_file, type='l', add=FALSE) ## show tracks as line plot

## example 1b) scatter plot from csv-file on existing landmask:
require('oceanmap')
plotmap('lion') ## use keyword to derive area limits
plot_geopos(csv_file, add=TRUE) ## show tracks as scatter plot

## example 2) probability surfaces of horizontal tracks from nc-file:
## this can take some time as it includes time consuming data processing
nc_file <- system.file("example_files/15P1019-104659-1-GPE3.nc",package="RchivalTag")
plot_geopos(nc_file)
```

plot_TS

plot time series data

Description

plotting functions for time series data (e.g. depth or temperature time series data from archival tags) with user specified xtick intervals.

Usage

```
plot_DepthTS(df, y="Depth", xlim, ylim, xticks_interval,
             ylab=y, xlab="Time (UTC)", main, main.line=1, plot_info=TRUE,
             ID, ID_label="Serial",
             plot_DayTimePeriods=TRUE, twilight.set="ast",
             type="l", las=1, xaxs="i", yaxs="i", cex=1,
             plot_box=TRUE, bty="l", Return=FALSE, ...)

plot_TS(df, y="Depth", xlim, ylim, xticks_interval,
        ylab=y, xlab="Time (UTC)", main, main.line=1, plot_info=TRUE,
        ID, ID_label="Serial",
        plot_DayTimePeriods=TRUE, twilight.set="ast",
        type="l", las=1, xaxs="i", yaxs="i", cex=1,
        plot_box=TRUE, bty="l", Return=FALSE, ...)

empty.plot_TS(xlim, ylim, xticks_interval, ylab="", xlab="Time (UTC)",
              las=1, xaxs="i", yaxs="i", cex=1,
              plot_box=TRUE, bty="l", ...)
```

Arguments

df	data.frame holding the time series data to be plotted, including the x-vector 'date.long' (in POSIXct-format and UTC), and the numeric y-vector whose label is defined by y.
y	character label of time series vector to be plotted (by default 'Depth').
xlim	the x limits (x1, x2) of the plot (by default range(df\$date.long), but needs to be specified in empty.plot_TS).
ylim	the y limits of the plot (by default range(df[[y]]), but needs to be specified in empty.plot_TS).
xticks_interval	time step of the x-axis ticklabels in (full) hours. By default 3 hours for xlim differences <= 1 day, and 6 hours for differences > 1 day.
ylab, xlab	the y- and x-axis labels.
main, main.line	main title (by default "Tag ID") for the plot and its line (see mtext for reference).
plot_info	whether the plot title and axes labels should be shown (by default TRUE).
ID, ID_label	Tag ID and its label (column name; by default "Serial") to be selected (e.g. if input data frame holds tagging data from several tags).
type	what type of plot should be drawn. Possible types are: <ul style="list-style-type: none"> • "p" for points, • "l" for lines (default), • "b" for both, • "c" for the lines part alone of "b", • "o" for both 'overlapped', • "n" for nothing (similar to empty.plot_TS-function call)
las	numeric in 0,1,2,3; the style of axis labels. 0: always parallel 1: always horizontal (default) 3: always perpendicular 4: always vertical
xaxs, yaxs	The style of axis interval calculation to be used for the x-and y-axes. Possible values are "r" and "i" (default). The styles are generally controlled by the range of data or xlim, if given. Style "r" (regular) first extends the data range by 4 percent at each end and then finds an axis with pretty labels that fits within the extended range. Style "i" (internal) just finds an axis with pretty labels that fits within the original data range.
cex	The standard font size (by default 1). Attention, the (internal) standard font size of axis labels cex.axis is 0.9*cex and the plot title cex.main= 1.2*cex.

`plot_TimePeriods`, `twilight.set`
 whether day-time periods ('Night', 'Dawn', 'Day', 'Dusk') should be plotted as shaded areas. In case that `plot_TimePeriods` is set TRUE, the limits of each time period are required (columns `sunrise`, `sunset`, `dawn.ast/dawn.naut` and `dawn.ast/dawn.naut` in POSIXct-format. In case of the twilight events, the additional argument `twilight.set` defines the suffix of the twilight-set to be selected ("ast" for astronomical dawn and dusks vs "naut" for nautical twilight events). If any of the day-time columns, described above, is missing, it/they will be calculated based on geolocation estimates (required columns `Lon` and `Lat`) through an internal call of function `get_TimeLimits`.

`plot_box`, `bty` whether a box of box-type `bty` should be plotted (by default TRUE. `bty` is one of "o" (the default), "l", "7", "c", "u", or "j" the resulting box resembles the corresponding upper case letter.

Return whether edited time series data set should be returned (by default FALSE).

... additional arguments to be passed to [plot](#).

Author(s)

Robert K. Bauer

See Also

[hist_tad](#), [hist_tat](#)

Examples

```
### load sample depth and temperature time series data from miniPAT:
ts_file <- system.file("example_files/104659-Series.csv", package="RchivalTag")
ts_df <- read.table(ts_file, header = TRUE, sep = ",")
head(ts_df)
ts_df$date.long <- as.POSIXct(strptime(paste(ts_df$Day, ts_df$Time),
                                         "%d-%b-%Y %H:%M:%S", tz = "UTC"))
ts_df$date <- as.Date(ts_df$date.long)

### select subsets (dates to plot)
plot_DepthTS(ts_df, plot_TimePeriods = FALSE, xlim = unique(ts_df$date)[2:3])
xlim <- c("2016-08-10 6:10:00", "2016-08-11 17:40:00")
plot_DepthTS(ts_df, plot_TimePeriods = FALSE, xlim = xlim)

### check xtick time step:
plot_DepthTS(ts_df, plot_TimePeriods = FALSE, xlim = "2016-08-10")
plot_DepthTS(ts_df, plot_TimePeriods = FALSE, xlim = "2016-08-10", xticks_interval = 2)

### add daytime periods during plot-function call and return extended data set
ts_df$Lon <- 5; ts_df$Lat <- 43
plot_DepthTS(ts_df, plot_TimePeriods = TRUE, xlim = unique(ts_df$date)[2:3])
ts_df2 <- plot_DepthTS(ts_df, plot_TimePeriods = TRUE, Return = TRUE)
names(ts_df2)
```

```

names(ts_df2)

### add daytime periods before function call
ts_df_extended <- get_DayTimeLimits(ts_df)
plot_DepthTS(ts_df_extended, plot_DayTimePeriods = TRUE)
plot_DepthTS(ts_df_extended, plot_DayTimePeriods = TRUE, twilight.set = "naut")

### introduce data transmission gaps that are then filled internally
### as well as daytime periods based on interpolated Lon & Lat positions
ts_df_cutted <- ts_df[-c(200:400, 1800:2200), ]
plot_DepthTS(ts_df_cutted, plot_DayTimePeriods = FALSE)
plot_DepthTS(ts_df_cutted, plot_DayTimePeriods = TRUE)

### example for empty.plotTS and adding time series data as line:
empty.plot_TS(xlim="2016-08-10",ylim=c(100,0))
lines(ts_df$date.long, ts_df$Depth)

### alternative:
plot_DepthTS(ts_df, xlim=c("2016-08-10","2016-08-12"), plot_DayTimePeriods = TRUE, type='n')
lines(ts_df$date.long, ts_df$Depth)

```

RchivalTag

RchivalTag - Analyzing Archival Tagging Data

Description

RchivalTag provides a set of functions to analyze and visualize (aquatic) archival tagging data, including:

- "(Depth) time series data" (See [empty.plot_TS](#), [plot_TS](#) & [plot_DepthTS](#))
- "Time-at-Depth (TaD) and Time-at-Temperature (TaT) frequencies" (See [ts2histos](#), [merge_histos](#), [hist_tad](#) & [hist_tat](#))
- "Depth Temperature profiles (time series data)" (See [interpolate_TempDepthProfiles](#), [get_thermalstrat](#) & [image_TempDepthProfiles](#))
- "PDT (PAT-style Depth Temperature profiles) data" (See [read_PDT](#), [interpolate_TempDepthProfiles](#), [get_thermalstrat](#) & [image_TempDepthProfiles](#))
- "visualization of geolocation estimates" (See: [plot_geopos](#))

Details

TaD-/TaT-histogram data

- The package allows to read and calculate standard summary data products (TaD-/TaT-profiles, see above) from recovered or transmitted time series data sets as well as to merge and visualize such summary data products from different tag setups/tagging programs. For more information on these data products, please see: Wildlife Computers (2016).

Depth time series data

- data visualization, optionally highlighting daytime differences (dawn, day, dusk, night).

Depth-temperature time series data

- data visualization and examination of the thermal stratification of the water column (i.e. thermocline depth, gradient and stratification index), based on previously interpolated. The paper by Bauer et al. (2015) is highly recommended in this context.

Compatibility

So far, the package is mainly adapted for archival tagging data from **Wildlife Computers**, but can also be applied to data from other tag manufacturers (e.g. see [ts2histos](#) in order to calculate TaD & TaT-frequencies from time series data). Function examples are based on the transmitted data sets of a miniPAT-tag from the BLUEMED-project <http://bluemed-project.com/>, funded by the French National Research Agency (ANR; <http://www.agence-nationale-recherche.fr>).

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

Wildlife Computers (2016) MiniPAT-User-Guide, 4 April 2016, 26 pp. <http://wildlifecomputers.com/wp-content/uploads/manuals/MiniPAT-User-Guide.pdf>

read_histos

reads a TAD/TAT-histogram file from archival tags

Description

reads or posttreats a manually loaded standard histogram data file, containing Time-at-Depth (TAD) and Time-at-Temperature (TAT) frequency data, from archival tags by **Wildlife Computers**.

Usage

```
read_histos(hist_file)
```

Arguments

hist_file	character string indicating the name of a standard Wildlife Computers file to read or the data.frame of a manually loaded histogram data file. The Date-vector of the file is expected to be or the format "%H:%M:%S %d-%b-%Y, tz='UTC'". The combination of the columns DeployID, Ptt and Serial is assumed to provide an unique key to distinguish data from individual tags.
-----------	--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------

Details

This function reads or posttreats a manually loaded standard Wildlife Computers histogram file including Time-at-Depth (TAD) and Time-at-Temperature (TAT) frequency data. In the post-treatment, the histogram data is split in lists of TAD and TAT per individual (see below). Thus processed data from several histogram files (or similarly processed time series data) can be combined using the function [combine_histos](#). Merging of histogram data from several tags, based on similar or user-specified TAD and TAT-bin_breaks, can be done by applying function [merge_histos](#). To generate TAD/TAT histogram data from depth and temperature time series data, see [ts2histos](#).

Value

A list-of-lists containing the loaded histogram data. Lists of TAD and TAT data are distinguished at the first nesting level. Further sublists contain the bin_breaks and [data.frames](#) of the histogram data per tag (ID). Tag IDs are constructed based on the columns DeployID, Ptt and Serial keys (e.g. DeployID.101_Ptt.102525).

```
$ TAD:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
$ TAT:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
..$ ID2 : List of 2
...
```

Author(s)

Robert K. Bauer

See Also

[ts2histos](#), [combine_histos](#), [merge_histos](#), [hist_tad](#), [hist_tat](#)

Examples

```
## example 1) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv", package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv", package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
```

```

hist_dat_merged <- merge_histos(hist_dat_combined, force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
unmerge_histos(hist_dat_merged)

```

read_PDT

read PDT data from archival one or multiple tags

Description

reads PDT data (PAT-style Depth Temperature profiles) from archival tags by **Wildlife Computers**). The PDT file can contain data from one or multiple tags.

What are PDTs?

PDT data provides minimum and maximum water temperatures during a user-programmed interval (usually 24h) at 8 to 16 depths. The sampled depths are thereby rounded (binned) to multiples of 8 and include the minimum and maximum depth bins as well as the 6 to 14 most frequent depth bins at which the tagged animal was located. The total number of depth bins (8 or 16) also depends on the tagged animals' behaviour. If the animal was in waters deeper than 400 m during the summary data period, the range of temperature at 16 depth bins will be reported, otherwise 8.

Why using PDT data?

Despite its low resolution, PDT data can give accurate information on the in-situ thermal stratification of the water column (e.g. thermocline depth, stratification index, ocean heat content) experienced by the tagged animal, as illustrated by Bauer et al. (2015). Accordingly, PDT data can provide precious insights into the relations between animal behaviour and environmental conditions. See the example section below on how to obtain thermal stratification indicators of the water column from PDT data.

For instance, daily PDT data can be interpolated and then visualized using functions [interpolate_PDTs](#) and [image_TempDepthProfiles](#), respectively. This facilitates the analysis of temporal changes of temperature profiles, for instance, in relation to animal behaviour (e.g. diving behaviour).

Usage

```
read_PDT(pdt_file, sep=",")
```

Arguments

pdt_file	character string indicating the name of a standard PDT-file. The Date-vector of the file is expected to be or the format "%H:%M:%S %d-%b-%Y, tz='UTC'".
----------	---------------------------------------------------------------------------------------------------------------------------------------------------------

sep the field separator character. Values on each line of the file are separated by this character (default is ',').

Value

A [data.frame](#) with the columns:
 "pdt_file", "DeployID", "Ptt", "NumBins", "Depth", "MinTemp", "MaxTemp", "date.long", "date",
 "MeanPDT"

Attention: Column "MeanPDT" is not measured but calculated as the average of "MinTemp" and "MaxTemp" values.

Author(s)

Robert K. Bauer

References

Bauer, R., F. Forget and JM. Fromentin (2015) Optimizing PAT data transmission: assessing the accuracy of temperature summary data to estimate environmental conditions. Fisheries Oceanography, 24(6): 533-539, doi: [10.1111/fog.12127](https://doi.org/10.1111/fog.12127)

See Also

[bin_TempTS](#), [interpolate_PDTs](#), [image_TempDepthProfiles](#)

Examples

```
## step I) read sample PDT data file:
setwd(system.file("example_files", package="RchivalTag"))
PDT <- read_PDT("104659-PDTs.csv")
head(PDT)

## step II) interpolate average temperature fields (MeanPDT) from PDT file:
m <- interpolate_PDTs(PDT)
str(m)
m$sm

## step III) calculate thermal stratification indicators per day (and tag):
strat <- get_thermalstrat(m, all_info = TRUE)
strat <- get_thermalstrat(m, all_info = FALSE)

## step IV) plot interpolated profiles:
image_TempDepthProfiles(m$station.1)
```

ts2histos	<i>convert depth and temperature time series data to discrete Time-at-Depth and Time-at-Temperature data (histogram data)</i>
-----------	-------------------------------------------------------------------------------------------------------------------------------

Description

convert depth and temperature time series data to discrete Time-at-Depth (TaD) and Time-at-Temperature (TaT) data (histogram data) at user-defined breakpoints

Usage

```
ts2histos(ts_df, tad_breaks=NULL, tat_breaks=NULL, split_by=NULL, aggregate_by="Ptt")
```

Arguments

ts_df	dataframe of depth time series data. Obligatory columns are the numeric vector "Depth", "date" (of class Date) and "Serial". split.by defines an optional vector to consider (e.g. day.period).
tad_breaks, tat_breaks	a numeric vector, defining the depth and/or temperature breakpoints of the histogram cells.
split_by	Name of the logical vector by which TaD/TaT data shall be splitted (e.g. day.time).
aggregate_by	character vector defining the columns by which the tagging data should be aggregated. Should contain columns that identify tags (e.g. Serial, Ptt, DeployID) the date and/or day time period (to separate records from night, day, dawn and dusk see classify_DayTime). Default values are: date, Day and Ptt.

Details

Time-at-Depth and Time-at-Temperature frequencies (histograms) are a standard data product of archival tags (incl. tag models TDR-Mk9, PAT-Mk10 and miniPAT by [Wildlife Computers](#)) that allow to assess habitat preferences of tagged animals. It can be likewise generated from transmitted or recovered time series data sets, which is the purpose of this function.

However, different depth and temperature bin breaks are often used during different deployment programs, which makes a later comparative analysis of TaT and TaD data difficult. For such cases, the functions [combine_histos](#) and [merge_histos](#) can be applied to merge TaT and TaD frequencies based on common bin breaks of different tags.

To visualize Time-at-Temperature (TaT) and Time-at-Depth (TaD) data, please see [hist_tat](#) and [hist_tad](#), respectively.

Author(s)

Robert K. Bauer

See Also

[read_histos](#), [hist_tad](#), [merge_histos](#)

Examples

```
### load sample depth and temperature time series data from miniPAT:
ts_file <- system.file("example_files/104659-Series.csv",package="RchivalTag")
ts_df <- read.table(ts_file, header = TRUE, sep = ",")
head(ts_df)

tad_breaks <- c(0, 2, 5, 10, 20, 50, 100, 200, 300, 400, 600, 2000)
tat_breaks <- c(10,12,15,17,18,19,20,21,22,23,24,27)

## example 1a) convert only DepthTS data to daily TAD frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks)
hist_tad(ts_df, bin_breaks = tad_breaks, do_mid.ticks = FALSE)

## convert 1b) only TemperatureTS data to daily TAT frequencies:
tat <- ts2histos(ts_df, tat_breaks = tat_breaks)
hist_tat(ts_df, bin_breaks = tat_breaks, do_mid.ticks = FALSE)
hist_tat(tat$TAT$merged, do_mid.ticks = FALSE)

## convert 1c) DepthTS & TemperatureTS data to daily TAD & TAT frequencies:
ts2histos(ts_df, tad_breaks = tad_breaks, tat_breaks = tat_breaks)

## example 2) rebin daily TAD frequencies:
tad <- ts2histos(ts_df, tad_breaks = tad_breaks)
tad2 <- rebin_histos(hist_list = tad, tad_breaks = tad_breaks[c(1:3,6:12)])
par(mfrow=c(2,2))
hist_tad(tad, do_mid.ticks = FALSE) ## example for multiple individuals
hist_tad(tad$TAD$merged, do_mid.ticks = FALSE)
hist_tad(tad$TAD$merged, bin_breaks = tad_breaks[c(1:3,6:12)]) ## from inside hist_tad
```

unmerge_histos	<i>unmerge previously grouped or merged lists of TAD/TAT frequency data</i>
----------------	-----------------------------------------------------------------------------

Description

This function unmerges previously grouped or merged lists of TAD/TAT frequency data, and thus allows to add TAD/TAT lists from new tags (see [combine_histos](#)).

Usage

```
unmerge_histos(hist_list)
```

Arguments

hist_list A previously grouped or merged list-of-lists to be unmerged (seperated by tags).

Value

A list-of-lists of ungrouped/unmerged TAD and TAT frequency data.

```
$ TAD:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
$ TAT:List
..$ ID1 : List of 2
.. ..$ bin_breaks: num
.. ..$ df : data.frame
..$ ID2 : List of 2
...
```

Author(s)

Robert K. Bauer

See Also

[combine_histos](#), [merge_histos](#), [hist_tad](#)

Examples

```
## example 1) read, merge and plot TAD frequency data from several files:
## part I - read histogram data from two files:
hist_dat_1 <- read_histos(system.file("example_files/104659-Histos.csv",package="RchivalTag"))
hist_dat_2 <- read_histos(system.file("example_files/104659b-Histos.csv",package="RchivalTag"))
## note the second list is based on the same data (tag), but on different bin_breaks

## part II - combine TAD/TAT frequency data from separate files in one list:
hist_dat_combined <- combine_histos(hist_dat_1, hist_dat_2)
par(mfrow=c(2,1))
hist_tad(hist_dat_combined)
hist_tat(hist_dat_combined)

## part III - force merge TAD/TAT frequency data from separate files
# in one list, by applying common bin_breaks:
hist_dat_merged <- merge_histos(hist_dat_combined,force_merge = TRUE)
hist_tad(hist_dat_merged)
hist_tat(hist_dat_merged)

## part IV - plot merged data:
hist_tad(hist_dat_merged) # of all tags
unique(hist_dat_merged$TAD$merged$df$DeployID) ## list unique tags in merged list
hist_tad(hist_dat_merged, select_id = "15P1019b", select_from = 'DeployID') # of one tag

## part V - unmerge data:
unmerge_histos(hist_dat_merged)
```

Index

bin_TempTS, [2](#), [18](#), [20](#), [32](#)

classify_DayTime, [3](#), [7](#), [11](#), [14](#), [33](#)

cmap, [18](#), [24](#)

combine_histos, [5](#), [12](#), [16](#), [22](#), [23](#), [30](#), [33–35](#)

crepuscule, [4](#), [6](#), [7](#)

data.frame, [2](#), [4](#), [6–8](#), [20](#), [24](#), [26](#), [29](#), [30](#), [32](#)

Date, [33](#)

empty.plot_TS, [28](#)

empty.plot_TS(plot_TS), [25](#)

get_DayTimeLimits, [3](#), [4](#), [6](#)

get_thermalstrat, [7](#), [18](#), [20](#), [28](#)

hist_tad, [5](#), [10](#), [12](#), [16](#), [22–24](#), [27](#), [28](#), [30](#), [33](#), [35](#)

hist_tat, [5](#), [12](#), [14](#), [22](#), [24](#), [27](#), [28](#), [30](#), [33](#)

image_TempDepthProfiles, [2](#), [3](#), [17](#), [18–20](#), [28](#), [31](#), [32](#)

interpolate_PDTs, [17](#), [31](#), [32](#)

interpolate_PDTs
(interpolate_TempDepthProfiles),
[19](#)

interpolate_TempDepthProfiles, [2](#), [3](#), [8](#), [9](#),
[17](#), [19](#), [28](#)

merge_histos, [5](#), [12](#), [16](#), [21](#), [28](#), [30](#), [33](#), [35](#)

mtext, [26](#)

plot, [24](#), [27](#)

plot_DepthTS, [24](#), [28](#)

plot_DepthTS(plot_TS), [25](#)

plot_geopos, [24](#), [28](#)

plot_TS, [25](#), [28](#)

plotmap, [24](#)

RchivalTag, [28](#)

read_histos, [12](#), [15](#), [29](#), [33](#)

read_PDT, [2](#), [3](#), [18](#), [20](#), [28](#), [31](#)

rebin_histos(merge_histos), [21](#)

set.colorbarp, [18](#)

sunriset, [4](#), [6](#), [7](#)

ts2histos, [11](#), [12](#), [14–16](#), [28–30](#), [33](#)

unmerge_histos, [5](#), [23](#), [34](#)