

Package: sftrack (via r-universe)

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Title Modern Classes for Tracking and Movement Data

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Depends R (>= 3.2.0)

Imports sf

Description Modern classes for tracking and movement data, building on 'sf' spatial infrastructure, and early theoretical work from Turchin (1998, ISBN: 9780878938476), and Calenge et al. (2009) <[doi:10.1016/j.ecoinf.2008.10.002](https://doi.org/10.1016/j.ecoinf.2008.10.002)>. Tracking data are series of locations with at least 2-dimensional spatial coordinates (x,y), a time index (t), and individual identification (id) of the object being monitored; movement data are made of trajectories, i.e. the line representation of the path, composed by steps (the straight-line segments connecting successive locations). 'sftrack' is designed to handle movement of both living organisms and inanimate objects.

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Encoding UTF-8

LazyData true

RoxygenNote 7.2.3

VignetteBuilder knitr

URL <https://mablab.org/sftrack/>, <https://github.com/mablab/sftrack>

BugReports <https://github.com/mablab/sftrack/issues>

Suggests testthat (>= 2.1.0), utils, ggplot2, adehabitatLT, knitr, geosphere, scales, covr, rmarkdown, lwgeom

Repository <https://mablab.r-universe.dev>

RemoteUrl <https://github.com/mablab/sftrack>

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active_group	<i>Access the active_group value</i>
--------------	--------------------------------------

Description

The active group is the combination of group names to group the data sets. The active_group acts essentially like a paste(names_of_groups, sep = ' ') grouping variable.

Usage

```
active_group(x)
```

Arguments

```
x                a c_grouping
```

Examples

```
#'
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
mb1 <- make_c_grouping(x = burstz, active_group = c("id", "month"))

# see the current active burst
active_group(mb1)

# change the active burst
active_group(mb1) <- "id"

# Using a full data set
my_track <- as_sftrack(raccoon,
  time = "timestamp",
  error = NA, coords = c("longitude", "latitude"),
  group = burstz
)

summary(my_track)

# change active group
active_group(my_track$sft_group) <- "id"

summary(my_track)
```

```
active_group<-      Set new active group
```

Description

Set new active group

Usage

```
active_group(x) <- value

## S3 replacement method for class 'sftrack'
active_group(x) <- value
```

```
## S3 replacement method for class 'sftraj'
active_group(x) <- value

## S3 replacement method for class 'c_grouping'
active_group(x) <- value
```

Arguments

x	sftrack/sftraj/c_grouping/s_group object
value	character vector of the grouping names to make active

as_sftrack	<i>Convert objects into sftrack objects.</i>
------------	--

Description

This function converts x,y,z data into an sftrack object with a sf_geometry column of sf_POINTS. Creates a 'grouping' column to group movement data and sets dedicated time and error columns.

Raw data inputted in two ways: vector or data.frame. 'Vector' inputs gives the argument as a vector where length = nrow(data). 'Data.frame' inputs gives the arguments as the column name of 'data' where the input can be found. Either input is allowed on any given argument.

Some options are global and required regardless

Usage

```
as_sftrack(data = data.frame(), ...)
```

```
## S3 method for class 'data.frame'
as_sftrack(
  data = data.frame(),
  ...,
  coords = c("x", "y"),
  group = "id",
  active_group = NA,
  time = "time",
  error = NA,
  crs = NA,
  zeroNA = FALSE,
  group_name = "sft_group",
  timestamp_name = "sft_timestamp",
  error_name = "sft_error",
  overwrite_names = FALSE
)
```

```
## S3 method for class 'sftraj'
as_sftrack(data, ...)
```

```

## S3 method for class 'ltraj'
as_sftrack(data, ...)

## S3 method for class 'sf'
as_sftrack(
  data,
  ...,
  coords,
  group,
  active_group = NA,
  time,
  error = NA,
  group_name = "sft_group",
  timestamp_name = "sft_timestamp",
  error_name = "sft_error",
  overwrite_names = FALSE
)

```

Arguments

data	a data.frame of the movement data, if supplied all data.frame inputs, than is optional
...	extra information to be passed on to as_sftrack
coords	a character vector describing where the x,y,z coordinates are located in 'data' or a list with x,y,z (optional) vectors
group	a list of named vectors describing multiple grouping variables or a character vector naming the other grouping columns in 'data'.
active_group	a character vector of the burst names to be 'active' to group data by for analysis
time	a vector of time information, can be either POSIX or an integer or a character string naming the column in 'data' where the time information is located
error	(optional) a vector of error information for the movement data a character string naming the column in 'data' where the error information is located
crs	Coordinate reference system to be assigned; object of class 'crs'. Defaults to NA
zeroNA	logical whether to convert 0s in spatial data into NAs. Defaults to FALSE.
group_name	(optional) new column name for grouping data
timestamp_name	(optional) new column name for time data
error_name	(optional) new column name for error data
overwrite_names	T/F Whether to overwrite data if a group/time/error column name is supplied but already in data

Details

Convert objects into sftrack objects.

Examples

```

#'
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
# Input is a data.frame
my_track <- as_sftrack(raccoon,
  group = burstz, time = "timestamp",
  error = NA, coords = c("longitude", "latitude")
)

# Input is a ltraj
library("adehabitatLT")
ltraj_df <- as.ltraj(
  xy = raccoon[, c("longitude", "latitude")],
  date = as.POSIXct(raccoon$timestamp),
  id = raccoon$animal_id, typeII = TRUE,
  infolocs = raccoon[, 1:6]
)

my_sftrack <- as_sftrack(ltraj_df)
head(my_sftrack)

# Input is a sf object
library("sf")
df1 <- raccoon[!is.na(raccoon$latitude), ]
sf_df <- st_as_sf(df1, coords = c("longitude", "latitude"))

new_sftrack <- as_sftrack(sf_df, group = c(id = "animal_id"), time = "timestamp")
head(new_sftrack)

# Input is an sftraj object
my_traj <- as_sftraj(raccoon,
  time = "timestamp",
  error = NA, coords = c("longitude", "latitude"),
  group = burstz
)

new_track <- as_sftrack(my_traj)
head(new_track)
#####
# Builder

```

as_sftraj

Convert objects into sftrack objects.

Description

This function converts x,y,z data into an sftrack object with a sf_geometry column of sf_POINTS. Creates a 'grouping' column to group movement data and sets dedicated time and error columns.

Raw data inputted in two ways: vector or data.frame. 'Vector' inputs gives the argument as a vector where length = nrow(data). 'Data.frame' inputs gives the arguments as the column name of 'data' where the input can be found. Either input is allowed on any given argument.

Some options are global and required regardless

Usage

```
as_sftraj(data = data.frame(), ...)
```

```
## S3 method for class 'data.frame'
```

```
as_sftraj(  
  data = data.frame(),  
  ...,  
  coords = c("x", "y"),  
  group = "id",  
  active_group = NA,  
  time = "time",  
  error = NA,  
  crs = NA,  
  zeroNA = FALSE,  
  group_name = "sft_group",  
  timestamp_name = "sft_timestamp",  
  error_name = "sft_error",  
  overwrite_names = FALSE  
)
```

```
## S3 method for class 'sftrack'
```

```
as_sftraj(data, ...)
```

```
## S3 method for class 'sf'
```

```
as_sftraj(  
  data,  
  ...,  
  coords,  
  group,  
  active_group = NA,  
  time,  
  error = NA,  
  group_name = "sft_group",  
  timestamp_name = "sft_timestamp",  
  error_name = "sft_error",  
  overwrite_names = FALSE  
)
```

```
## S3 method for class 'ltraj'
```

```
as_sftraj(data, ...)
```

Arguments

data	a data.frame of the movement data, if supplied all data.frame inputs, than is optional
...	extra information to be passed on to as_sftrack
coords	a character vector describing where the x,y,z coordinates are located in 'data' or a list with x,y,z (optional) vectors
group	a list of named vectors describing multiple grouping variables or a character vector naming the other grouping columns in 'data'.
active_group	a character vector of the burst names to be 'active' to group data by for analysis
time	a vector of time information, can be either POSIX or an integer or a character string naming the column in 'data' where the time information is located
error	(optional) a vector of error information for the movement data a character string naming the column in 'data' where the error information is located
crs	Coordinate reference system to be assigned; object of class 'crs'. Defaults to NA
zeroNA	logical whether to convert 0s in spatial data into NAs. Defaults to FALSE.
group_name	(optional) new column name for grouping data
timestamp_name	(optional) new column name for time data
error_name	(optional) new column name for error data
overwrite_names	T/F Whether to overwrite data if a group/time/error column name is supplied but already in data

Details

Convert objects into sftrack objects.

Examples

```
#
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
# Input is a data.frame
my_track <- as_sftraj(raccoon,
  group = burstz, time = "timestamp",
  error = NA, coords = c("longitude", "latitude")
)

# Input is a ltraj
library("adehabitatLT")
ltraj_df <- as.ltraj(
  xy = raccoon[, c("longitude", "latitude")],
  date = as.POSIXct(raccoon$timestamp),
  id = raccoon$animal_id, typeII = TRUE,
  infolocs = raccoon[, 1:6]
```



```

)

my_sftrack <- as_sftraj(ltraj_df)
head(my_sftrack)

# Input is a sf object
library("sf")
df1 <- raccoon[!is.na(raccoon$latitude), ]
sf_df <- st_as_sf(df1, coords = c("longitude", "latitude"))

new_sftrack <- as_sftrack(sf_df, group = c(id = "animal_id"), time = "timestamp")
head(new_sftrack)

# Input is an sftrack object
my_track <- as_sftrack(raccoon,
  time = "timestamp",
  error = NA, coords = c("longitude", "latitude"),
  group = burstz
)

new_traj <- as_sftraj(my_track)
head(new_traj)
#####

```

calc_sort_index

Calculate a new sort index for groups

Description

Calculate a new sort index for groups

Usage

```
calc_sort_index(x, active_group = NA)
```

Arguments

x	group or sftrack object
active_group	(optional), a new active group. If not included, defaults to the active group (if a c_grouping) or the group names

check_group_id	<i>Check there is a grouping id present</i>
----------------	---

Description

Check there is a grouping id present

Usage

```
check_group_id(x)
```

Arguments

x	a c_grouping
---	--------------

check_group_names	<i>Are group names equivalent for each s_group?</i>
-------------------	---

Description

Are group names equivalent for each s_group?

Usage

```
check_group_names(x)
```

Arguments

x	a c_grouping
---	--------------

check_names_exist	<i>Check if a set of column names are found in a data frame and return an error if not</i>
-------------------	--

Description

Check if a set of column names are found in a data frame and return an error if not

Usage

```
check_names_exist(data, names)
```

Arguments

data	a data.frame to check names against
names	the inputted column names

check_NA_coords	<i>Check if coordinates contain NAs in some columns but not others</i>
-----------------	--

Description

Check if coordinates contain NAs in some columns but not others

Usage

```
check_NA_coords(xyz)
```

Arguments

xyz	a data.frame of xy or xyz coordinates
-----	---------------------------------------

check_NA_group	<i>Check there are no NAs in burst</i>
----------------	--

Description

Check there are no NAs in burst

Usage

```
check_NA_group(x)
```

Arguments

x	a c_grouping
---	--------------

check_ordered	<i>Checks if grouping is ordered by time and then outputs the correct order</i>
---------------	---

Description

Checks if grouping is ordered by time and then outputs the correct order

Usage

```
check_ordered(group, time_data, return = TRUE)
```

Arguments

group	a c_grouping
time_data	a vector of time
return	T/F return the new order or just run check?

check_time	<i>Check if time is integer or posix</i>
------------	--

Description

Check if time is integer or posix

Usage

```
check_time(time)
```

Arguments

time	a vector of time
------	------------------

check_t_regular	<i>Check if time is regular for each burst and returns logical for each burst</i>
-----------------	---

Description

Check if time is regular for each burst and returns logical for each burst

Usage

```
check_t_regular(x)
```

Arguments

x	an sftack/sftraj object
---	-------------------------

dup_timestamp	<i>check that time is unique</i>
---------------	----------------------------------

Description

check that time is unique

Usage

```
dup_timestamp(x, time)
```

Arguments

x	An sftack/sftraj object or a multi_burst
time	vector of time, not required if given a sftack object.

fix_zero	<i>fix 0's to NAs in latitude and longitude</i>
----------	---

Description

fix 0's to NAs in latitude and longitude

Usage

```
fix_zero(xyz)
```

Arguments

xyz a data.frame of xy or xyz coordinates

Value

returns a data.frame with 0s replaced with NAs

geom_sftrack	<i>Function to plot sftrack objects in ggplot</i>
--------------	---

Description

This function can be added to ggplot() to plot an sftrack and sftraj Function does not yet work with ggplot grammar so you must put data= in this function

Usage

```
geom_sftrack(mapping, data, ...)

## S3 method for class 'sftrack'
geom_sftrack(mapping = ggplot2::aes(), data = NULL, ...)

## S3 method for class 'sftraj'
geom_sftrack(mapping = ggplot2::aes(), data = NULL, ..., step_mode = FALSE)
```

Arguments

mapping mapping aesthetics for ggplot.
 data the sftraj or sftrack object.
 ... arguments to passed to ggplot
 step_mode TRUE/FALSE, whether to plot in step_mode, See details

Details

step mode refers to considering the trajectory as individual 'steps', in the case of plot this means it will plot each line & point individually. This approach is much slower to plot when $n(\text{steps}) > 10,000$. The alternative method is to merge the steps into a multilinestring of continuous lines. This is much faster to plot.

Examples

```
#'
require("ggplot2")
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- c(id = "animal_id")

# sftraj will as well for the most part, however as its a more complex
# structure to speed up plotting.
my_sftraj <- as_sftraj(raccoon,
  time = "timestamp",
  coords = c("longitude", "latitude"),
  group = burstz
)

ggplot() +
  geom_sftrack(data = my_sftraj)
```

grouping-class

A class to group movement data

Description

This class describes grouping variables for movement data. The grouping object is composed of a list with named vectors. One of which must be 'id', this is the id of subject being monitored (commonly animal id in movement data) Can be any number of groups after that.

Usage

```
make_s_group(x)

make_c_grouping(x = NULL, active_group = NULL)

## S3 method for class 's_group'
c(...)

## S3 method for class 'c_grouping'
c(..., recursive = FALSE)
```

Arguments

x	a list containing named grouping variables, one item must be named 'id'. ex: list(id = 1, month = 'may'). For a c_grouping: A list of s_groups or a list of equal length named vectors which will be combined to create a c_grouping. ex: list(x = 1st_vector, y = 2nd_vector)
active_group	a vector of the names of the groups to be considered 'active'.
...	objects to be pasted together into a c_grouping
recursive	ignored

Details

A grouping is a list of possible categories to group the data. The 'active group' of these is the current grouping variables to be considered for analysis. The 'active group' can be any combination of the categories in a burst, and can change with the use of 'active_group()'.

An 's_group' is a single row group. It is a 1xn dimensional list with any length(n) > 1. Atleast one of the groups must be named 'id' which is the subjects id.

A 'c_grouping' is a collection of 's_groups's, it is a data.frame with dimensions of 1xnrow(data). One c_grouping has one 'active group' which describes the set of names in each s_group to group the data. When you change the 'active group', calculations and plots change accordingly to the new grouping levels.

You can create bursts with make_s_group and make_c_grouping.

Examples

```
# Make a single group
#'
make_s_group(x = list(id = "CJ11", month = 3, height = 10))

# Make a c_grouping
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
mb1 <- make_c_grouping(x = burstz, active_group = c("id", "month"))
str(mb1)

# Make a multi_burst from many ind_bursts
a <- make_s_group(list(id = 1, year = 2020))
b <- make_s_group(list(id = 1, year = 2020))
c <- make_s_group(list(id = 2, year = 2020))

c(a, b, c)
```

group_labels	<i>Shows grouping labels created from the s_group and the c_grouping</i>
--------------	--

Description

Shows grouping labels created from the s_group and the c_grouping

Usage

```
group_labels(x)

## S3 method for class 'sftrack'
group_labels(x)

## S3 method for class 'sftraj'
group_labels(x)

## S3 method for class 'c_grouping'
group_labels(x)
```

Arguments

x a sftrack or grouping object

group_names	<i>Display the levels of the sort index</i>
-------------	---

Description

Display the levels of the sort index

Usage

```
group_names(x)

## S3 method for class 'c_grouping'
group_names(x)

## S3 method for class 'sftraj'
group_names(x)

## S3 method for class 'sftrack'
group_names(x)
```

Arguments

x sftrack/sftraj/c_grouping/s_group object

is_linestring	<i>Is a trajectory geometry a linestring or a point</i>
---------------	---

Description

A step is a movement from one point to the next, with an sftraj object this manifests as a linestring. If, however, one of these two points is missing, the sftraj is created as a geometry collection of two points, the beginning and the end point, where one of the steps is NA. This function checks a trajectory geometry if its a linestring and returns a vector of T/F.

Usage

```
is_linestring(x)
```

Arguments

x	an sftraj object
---	------------------

make_step_geom	<i>Calculate step geometries given a set of groupings, time, and geometries</i>
----------------	---

Description

This calculates step geometries as individual line segments based on the active_group

Usage

```
make_step_geom(group, time_data, geometry)
```

Arguments

group	a c_grouping object
time_data	time vector
geometry	the geometry data from either sf or sf_track. Must be an sf geometry class

Examples

```
#'
library("sf")
geom <- st_as_sf(data.frame(
  x = c(1, 2, 2, 5),
  y = c(0, 1, 5, 7)
), coords = c("x", "y"))

burst <- list(id = rep(1, 4))
```

```

time <- 1:4

cg <- make_c_grouping(burst)

make_step_geom(
  group = cg,
  geometry = geom$geometry,
  time_data = time
)

```

merge_traj	<i>Merge connected lines and create an sf object</i>
------------	--

Description

This function returns a sf object grouped by each burst with a geometry column of multilinestrings for each grouping

Usage

```
merge_traj(x)
```

Arguments

x an sftraj object

new_sftrack	<i>Define an sftack</i>
-------------	-------------------------

Description

Define an sftack

Usage

```
new_sftrack(data, group_col, sf_col, time_col, error_col = NA)
```

Arguments

data	data.frame with multi_burst column, geometry column, time_col (integer/POSIXct), and error column (optional)
group_col	column name of grouping info in 'data'
sf_col	column name of geometry info in 'data'
time_col	column name of time info in 'data'
error_col	column name of error info in 'data'

new_sftraj	<i>Define an sftraj</i>
------------	-------------------------

Description

Define an sftraj

Usage

```
new_sftraj(data, group_col, sf_col, time_col, error_col = NA)
```

Arguments

data	data.frame with multi_burst column, geometry column, time_col (integer/POSIXct), and error column (optional)
group_col	column name of multi_burst in 'data'
sf_col	column name of geometry in 'data'
time_col	column name of time in 'data'
error_col	column name of error in 'data'

plot_sftrack	<i>Methods for plotting sftrack/sftraj</i>
--------------	--

Description

Methods for plotting sftrack/sftraj

Methods for plotting sftrack/sftraj

Usage

```
## S3 method for class 'sftrack'
plot(x, y, key.pos, key.width, ...)

## S3 method for class 'sftraj'
plot(x, y, key.pos, key.width, ..., step_mode)
```

Arguments

x	'sftrack' or 'sftraj' object
y	ignored
key.pos	Integer; side to plot a color key: 1 bottom, 2 left, 3 top, 4 right; set to NULL to omit key, or -1 to select automatically (defaults to 4; see plot_sf for more details).

key.width	Amount of space reserved for the key, including labels (see <code>plot_sf</code> for more details.)
...	Further arguments passed to <code>'plot.sf'</code> . Among others, arguments for the key are set differently in <code>'sftrack'</code> to allow for longer labels by default (but can be nevertheless adjusted).
step_mode	Logical; whether to plot in step mode, see details, defaults to TRUE, unless there are more than 10,000 steps.

Details

Step mode refers to considering the trajectory as individual 'steps', in the case of plot this means it will plot each line & point individually. This approach is much slower to plot with large objects, and is thus turned off when `n(steps)>10,000`. The alternative, much faster method is to merge the steps into a multiline string as continuous lines.

Examples

```
## Prepare an 'sftrack' object:
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- c(id = "animal_id")
my_sftrack <- as_sftrack(raccoon,
  time = "timestamp",
  coords = c("longitude", "latitude"),
  group = burstz
)

## Plotting with sftrack is just like sf. `...` will accept most
## arguments as 'plot.sf':
plot(my_sftrack, axes = TRUE, lwd = 5, cex = 5, bgc = "gray50")

## sftraj will as well for the most part; however it is a more complex
## structure that combines points and steps (in step mode):
my_sftraj <- as_sftraj(raccoon,
  time = "timestamp",
  coords = c("longitude", "latitude"),
  group = burstz
)
plot(my_sftraj, lwd = 5, cex = 5, bgc = "gray50", graticule = TRUE)
```

Print_sftrack_objects *Print methods for sftrack*

Description

Print methods for sftrack

Usage

```
## S3 method for class 'sftrack'  
print(x, n_row, n_col, ...)
```

Arguments

x	sftraj object
n_row	Integer of number of rows to display. Defaults to global option default if non supplied
n_col	Integer of number of columns to display + required sftrack columns (burst, geometry, time, and error). Defaults to global option default if non supplied
...	other arguments passed onto print

Print_sftraj_objects *Print methods for sftraj*

Description

Print methods for sftraj

Usage

```
## S3 method for class 'sftraj'  
print(x, n_row, n_col, ...)
```

Arguments

x	sftraj object
n_row	Integer of number of rows to display. Defaults to global option default if non supplied
n_col	Integer of number of columns to display + required sftrack columns (burst, geometry, time, and error). Defaults to global option default if non supplied
...	other arguments passed onto print

 raccoon

Movements of two raccoons in an urban park in Florida

Description

A dataset of two raccoons collared with GPS collars for one month in January 2019 in Tree Tops Park, Broward County, Florida, US.

Usage

```
raccoon
```

Format

A data frame with 445 rows and 10 variables:

animal_id ID of individual. TTP: tree tops park, i.e the tagging site.

timestamp The date and time of gps fix in UTC

latitude Latitude in degrees

longitude Longitude in degrees

height Altitude in meters based on satellite positios

hdop Horizontal precision

vdop Vertical precision

fix The number of satellite fixes

 step_metrics

Calculates step metrics including distance, dt, dx, and dy.

Description

Calculates step metrics including distance, dt, dx, and dy.

Usage

```
step_metrics(sftraj)
```

Arguments

sftraj an sftrack/sftraj object. sftrack objects will be converted to sftraj internally for calculation.

Examples

```

#'
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
# Input is a data.frame
my_sftraj <- as_sftraj(raccoon,
  group = burstz, time = "timestamp",
  error = NA, coords = c("longitude", "latitude")
)

step_metrics(my_sftraj)[1:10, ]

```

step_recalc

recalculate step geometry

Description

Step geometries in sftraj objects are linestrings going from t1 to t2 of a 'step'. As these are stored at the row level they are not dynamic to changes in t2. `step_recalc` allows you to recalculate these geometries if your data.frame has changed because of subsetting or filtering.

Usage

```
step_recalc(x, return = FALSE)
```

Arguments

x	an sftraj object.
return	return <code>step_geometry</code> instead of replacing sftraj object with new step geometry. Defaults to FALSE

summary_sftrack

Summarize sftrack objects

Description

Summarize sftrack objects

Usage

```
summary_sftrack(x)
```

Arguments

x	an sftrack object
---	-------------------

traj_geom	<i>Return a list of sf_POINTS or a data.frame from a sftraj object</i>
-----------	--

Description

Return a list of sf_POINTS or a data.frame from a sftraj object

Usage

```
pts_traj(traj, sfc = FALSE)

coord_traj(traj)
```

Arguments

traj	a trajectory geometry from sf_traj
sfc	TRUE/FALSE should the return by an sfc or a list of points. Defaults to FALSE

Examples

```
#'
data("raccoon")
raccoon$timestamp <- as.POSIXct(raccoon$timestamp, "EST")
burstz <- list(id = raccoon$animal_id, month = as.POSIXlt(raccoon$timestamp)$mon)
# Input is a data.frame
my_traj <- as_sftraj(raccoon,
  time = "timestamp",
  error = NA, coords = c("longitude", "latitude"),
  group = burstz
)
print(my_traj, 5, 10)

# extract a list of points
pts_traj(my_traj)[1:10]

# or a data.frame of points
coord_traj(my_traj)[1:10]
```

which_duplicated	<i>Which grouping/time stamp combos are duplicated.</i>
------------------	---

Description

This function returns a data.frame of which rows are duplicated and their time stamps.

Usage

```
which_duplicated(data = data.frame(), group, time)
```

Arguments

<code>data</code>	a data.frame containing burst or time data (if necessary)
<code>group</code>	a list where each entry is a vector of groupings where length == nrow(data)/nrow(time). Or a character vector describing the column name they are located in data
<code>time</code>	a vector of as.POSIXct time, or a character of the column name where it can be found in data

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