

# Package ‘animaltracker’

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**Title** Animal Tracker

**Version** 0.1.0

**Description** Utilities for spatial-temporal analysis and visualization of animal (e.g. cattle) tracking data. The core feature is a 'shiny' web application for customized processing of GPS logs, including features for data augmentation (e.g. elevation lookup), data selection, export, plotting, and statistical summaries. A data validation application allows for side-by-side comparison via time series plots and extreme value detection described by J.P. van Brakel <<https://stackoverflow.com/questions/22583391/peak-signal-detection-in-realtime-timeseries-data/>>.

**Depends** R (>= 3.5.0)

**Imports** zoo (>= 1.8.6), forcats (>= 0.4.0), lubridate (>= 1.7.0), tibble (>= 2.1.0), shinyBS (>= 0.61), V8 (>= 2.0), shinyjs (>= 1.0), shiny (>= 1.2.0), shinyWidgets (>= 0.4.4), shinycssloaders (>= 0.2.0), shinythemes (>= 1.1.2), leaflet (>= 2.0.2), leaflet.extras (>= 1.0.0), dplyr (>= 0.7.5), ggplot2 (>= 3.1.0), scales (>= 1.0.0), tidyr (>= 0.8.2), sp (>= 1.3.1), rgdal (>= 1.3.6), raster (>= 2.7.15), elevatr (>= 0.2.0), geosphere (>= 1.5.7)

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<code>app_server</code>	<i>Defines logic for updating the app based on user interaction in the ui</i>
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---

**Description**

Defines logic for updating the app based on user interaction in the ui

**Usage**

```
app_server(input, output, session)
```

**Arguments**

<code>input</code>	see shiny app architecture
<code>output</code>	see shiny app architecture
<code>session</code>	see shiny app architecture

**Value**

server function for use in a shiny app

---

<code>app_ui</code>	<i>Defines a user interface for the 'shiny' app</i>
---------------------	---

---

**Description**

Defines a user interface for the 'shiny' app

**Usage**

```
app_ui()
```

**Value**

ui function for use in a 'shiny' app

---

boxplot\_altitude      *Generates a boxplot to visualize the distribution of altitude by GPS.*

---

**Description**

Generates a boxplot to visualize the distribution of altitude by GPS.

**Usage**

```
boxplot_altitude(rds_path)
```

**Arguments**

rds\_path      Path of .rds animal data file to read in

**Value**

overall boxplot of altitude by GPS

**Examples**

```
# Boxplot of altitude for demo data .rds  
boxplot_altitude(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

boxplot\_time\_unit      *Generates a boxplot to visualize the distribution of time between GPS measurements by GPS unit.*

---

**Description**

Generates a boxplot to visualize the distribution of time between GPS measurements by GPS unit.

**Usage**

```
boxplot_time_unit(rds_path)
```

**Arguments**

rds\_path      Path of .rds animal data file to read in

**Value**

distribution of time between GPS measurements by GPS unit, as a boxplot

**Examples**

```
# Boxplot of GPS measurement time differences for demo data .rds  
boxplot_time_unit(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

calc_bearing	<i>Helper function for cleaning Columbus P-1 datasets. Given lat and long coords in degree decimal, convert to radians and compute bearing.</i>
--------------	---

---

**Description**

Helper function for cleaning Columbus P-1 datasets. Given lat and long coords in degree decimal, convert to radians and compute bearing.

**Usage**

```
calc_bearing(lat1, lon1, lat2, lon2)
```

**Arguments**

lat1	latitude of starting point
lon1	longitude of starting point
lat2	latitude of ending point
lon2	longitude of ending point

**Value**

bearing computed from given coordinates

---

clean_batch_df	<i>Cleans a directory of animal data files</i>
----------------	--

---

**Description**

Cleans a directory of animal data files

**Usage**

```
clean_batch_df(data_info, filters = TRUE, tz_in = "UTC", tz_out = "UTC")
```

**Arguments**

data_info	list of animal data frames with information about the data, generated by store_batch
filters	filter bad data points, defaults to true
tz_in	input time zone, defaults to UTC
tz_out	output time zone, defaults to UTC

**Value**

clean df with all animal data files from the directory

---

clean_export_files	<i>Cleans all animal GPS datasets (in .csv format) in a chosen directory. Optionally exports the clean data as spreadsheets, a single .rds data file, or as a list of data frames</i>
--------------------	---

---

**Description**

Cleans all animal GPS datasets (in .csv format) in a chosen directory. Optionally exports the clean data as spreadsheets, a single .rds data file, or as a list of data frames

**Usage**

```
clean_export_files(
  data_dir,
  tz_in = "UTC",
  tz_out = "UTC",
  export = FALSE,
  cleaned_filename = NULL,
  cleaned_dir = NULL
)
```

**Arguments**

data_dir	directory of GPS tracking files (in csv)
tz_in	input time zone, defaults to UTC
tz_out	output time zone, defaults to UTC
export	logical, whether to export the clean data, defaults to False
cleaned_filename	full name of output file (ending in .rds) when export is True
cleaned_dir	directory to save the processed GPS datasets as spreadsheets (.csv) when export is True

**Value**

list of cleaned animal GPS datasets

**Examples**

```
# Clean all animal GPS .csv datasets in the demo directory

clean_export_files(system.file("extdata", "demo_nov19", package = "animaltracker"))
```

---

clean_location_data	<i>Cleans a raw animal GPS dataset, implementing a standardized procedure to remove impossible values</i>
---------------------	---

---

**Description**

Cleans a raw animal GPS dataset, implementing a standardized procedure to remove impossible values

**Usage**

```
clean_location_data(
  df,
  dtype,
  filters = TRUE,
  aniid = NA,
  gpsid = NA,
  maxrate = 84,
  maxcourse = 100,
  maxdist = 840,
  maxtime = 100,
  tz_in = "UTC",
  tz_out = "UTC"
)
```

**Arguments**

df	data frame in standardized format (e.g., from a raw spreadsheet)
dtype	data type, iGotU or Columbus P-1
filters	filter bad data points, defaults to true
aniid	identification code for the animal
gpsid	identification code for the GPS device
maxrate	maximum rate of travel (meters/minute) between consecutive points
maxcourse	maximum distance (meters) between consecutive points
maxdist	maximum geographic distance (meters) between consecutive points
maxtime	maximum time (minutes) between consecutive points
tz_in	input time zone, defaults to UTC
tz_out	output time zone, defaults to UTC

**Value**

df of clean animal GPS data

**Examples**

```
# Clean a data frame from csv

## Read igotU data
bannock_df <- read.csv(system.file("extdata", "demo_nov19/Bannock_2017_101_1149.csv",
package = "animaltracker"), skipNul=TRUE)

## Clean and filter
clean_location_data(bannock_df, dtype = "igotu", filters = TRUE, aniid = 1149,
gpsid = 101, maxrate = 84, maxdist = 840, maxtime = 100)

## Clean without filtering
clean_location_data(bannock_df, dtype = "igotu", filters = FALSE, aniid = 1149,
gpsid = 101, maxrate = 84, maxdist = 840, maxtime = 100)

# Clean a data frame from txt

## Read Columbus P-1 data
columbus_df <- read_columbus(system.file("extdata", "demo_columbus.TXT",
package = "animaltracker"))

## Clean and filter
clean_location_data(columbus_df, dtype = "columbus", filters = TRUE, aniid = 1149,
gpsid = 101, maxrate = 84, maxdist = 840, maxtime = 100)
```

---

clean_store_batch	<i>Cleans a directory of animal data files and stores them locally in rds format</i>
-------------------	--

---

**Description**

Cleans a directory of animal data files and stores them locally in rds format

**Usage**

```
clean_store_batch(
  data_info,
  filters = TRUE,
  zoom = 11,
  get_slope = TRUE,
  get_aspect = TRUE,
  min_lat = data_info$min_lat,
  max_lat = data_info$max_lat,
  min_long = data_info$min_long,
  max_long = data_info$max_long,
```

```

    tz_in = "UTC",
    tz_out = "UTC"
)

```

### Arguments

data_info	list of animal data frames with information about the data, generated by store_batch
filters	filter bad data points, defaults to true
zoom	level of zoom, defaults to 11
get_slope	logical, whether to compute slope (in degrees), defaults to true
get_aspect	logical, whether to compute aspect (in degrees), defaults to true
min_lat	minimum latitude for filtering, defaults to min in data_info
max_lat	maximum latitude for filtering, defaults to max in data_info
min_long	minimum longitude for filtering, defaults to min in data_info
max_long	maximum longitude for filtering, defaults to max in data_info
tz_in	input time zone, defaults to UTC
tz_out	output time zone, defaults to UTC

### Value

df of metadata for animal file directory

---

compare_flags	<i>Joins and reformats two animal data frames for the purpose of flag comparison</i>
---------------	--

---

### Description

Joins and reformats two animal data frames for the purpose of flag comparison

### Usage

```
compare_flags(correct, candidate, elev = TRUE, slope = TRUE)
```

### Arguments

correct	reference df
candidate	df to be compared to the reference
elev	logical, whether to include elevation, defaults to true
slope	logical, whether to include slope, defaults to true

### Value

joined and reformatted df

## Examples

```
# Join and reformat unfiltered demo data and filtered demo data

compare_flags(demo_unfiltered_elev, demo_filtered_elev)
```

---

compare\_summarise\_daily

*Compares two animal datasets and calculates daily summary statistics by GPS GPS, date, lat, long, course, distance, rate, elevation column names should match.*

---

## Description

Compares two animal datasets and calculates daily summary statistics by GPS GPS, date, lat, long, course, distance, rate, elevation column names should match.

## Usage

```
compare_summarise_daily(correct, candidate, export = TRUE, out = NULL)
```

## Arguments

correct	reference df
candidate	df to be compared to the reference
export	logical, whether to export summary to .csv, defaults to False
out	desired file name of .csv output summary when export is True

## Value

summary df

## Examples

```
# Compare and summarise unfiltered demo cows to filtered, grouped by both Date and GPS

compare_summarise_daily(demo_unfiltered_elev, demo_filtered_elev)
```

---

`compare_summarise_data`

*Compares two animal data frames and calculates summary statistics. GPS, date, lat, long, course, distance, rate, elevation column names should match.*

---

### Description

Compares two animal data frames and calculates summary statistics. GPS, date, lat, long, course, distance, rate, elevation column names should match.

### Usage

```
compare_summarise_data(  
  correct,  
  candidate,  
  export = FALSE,  
  gps_out = NULL,  
  date_out = NULL  
)
```

### Arguments

<code>correct</code>	reference df
<code>candidate</code>	df to be compared to the reference
<code>export</code>	logical, whether to export summaries to .csv, defaults to False
<code>gps_out</code>	desired file name of .csv output summary by GPS collar when export is True
<code>date_out</code>	desired file name of .csv output summary by date when export is True

### Value

list containing `gps_out` and `date_out` as dfs

### Examples

```
# Compare and summarise unfiltered demo cows to filtered  
compare_summarise_data(demo_unfiltered_elev, demo_filtered_elev)
```

---

deg_to_dec	<i>Helper function for cleaning Columbus P-1 datasets. Given lat or long coords in degrees and a direction, convert to decimal.</i>
------------	---

---

**Description**

Helper function for cleaning Columbus P-1 datasets. Given lat or long coords in degrees and a direction, convert to decimal.

**Usage**

```
deg_to_dec(x, direction)
```

**Arguments**

x	lat or long coords in degrees
direction	direction of lat/long

**Value**

converted x

---

demo	<i>Demo animal GPS data from cows</i>
------	---------------------------------------

---

**Description**

Demo animal GPS data from cows

**Usage**

```
demo
```

**Format**

A data frame with 2171 rows and 29 variables

---

demo_comparison	<i>Demo comparison of two animal datasets</i>
-----------------	---

---

**Description**

Demo comparison of two animal datasets

**Usage**

demo\_comparison

**Format**

A data frame with 2758 rows and 33 variables

---

demo_filtered	<i>Filtered demo animal GPS data from cows</i>
---------------	--

---

**Description**

Filtered demo animal GPS data from cows

**Usage**

demo\_filtered

**Format**

A data frame with 2187 rows and 26 variables

---

demo_filtered_elev	<i>Filtered demo animal GPS data from cows with elevation appended at zoom 1</i>
--------------------	--

---

**Description**

Filtered demo animal GPS data from cows with elevation appended at zoom 1

**Usage**

demo\_filtered\_elev

**Format**

A data frame with 2187 rows and 29 variables

---

demo_info	<i>Raw demo animal GPS data from cows with information</i>
-----------	--

---

**Description**

Raw demo animal GPS data from cows with information

**Usage**

demo\_info

**Format**

A list with 10 elements

---

demo_meta	<i>Metadata for demo animal GPS data from cows</i>
-----------	--

---

**Description**

Metadata for demo animal GPS data from cows

**Usage**

demo\_meta

**Format**

A data frame with 6 rows and 11 variables

---

demo_unfiltered	<i>Unfiltered demo animal GPS data from cows</i>
-----------------	--

---

**Description**

Unfiltered demo animal GPS data from cows

**Usage**

demo\_unfiltered

**Format**

A data frame with 2288 rows and 32 variables

---

demo_unfiltered_elev	<i>Unfiltered demo animal GPS data from cows with elevation appended at zoom 1</i>
----------------------	--

---

**Description**

Unfiltered demo animal GPS data from cows with elevation appended at zoom 1

**Usage**

```
demo_unfiltered_elev
```

**Format**

A data frame with 2288 rows and 35 variables

---

detect_peak_modz	<i>Alternative implementation of the robust peak detection algorithm by van Brakel 2016</i>
------------------	---

---

**Description**

Alternative implementation of the robust peak detection algorithm by van Brakel 2016

**Usage**

```
detect_peak_modz(df_comparison, lag = 5, max_score = 3.5)
```

**Arguments**

df_comparison	output of compare_flags
lag	width of interval to compute rolling median and MAD, defaults to 5
max_score	modified z-score cutoff to classify observations as outliers, defaults to 3.5

**Value**

df with classifications

**Examples**

```
# Join and reformat unfiltered demo data and filtered demo data  
  
detect_peak_modz(demo_comparison, lag = 5, max_score = 3.5)
```

---

dev\_add\_to\_gitignore    *Add big files to a .gitignore file*

---

**Description**

Add big files to a .gitignore file

**Usage**

```
dev_add_to_gitignore(data_dir)
```

**Arguments**

data\_dir            directory of animal data files

**Value**

None

---

get\_data\_from\_meta    *Get animal data set from specified meta. If date range is invalid, automatically returns all animal data specified by meta\_df.*

---

**Description**

Get animal data set from specified meta. If date range is invalid, automatically returns all animal data specified by meta\_df.

**Usage**

```
get_data_from_meta(meta_df, min_date, max_date)
```

**Arguments**

meta\_df            data frame of specified meta  
 min\_date          minimum date specified by user  
 max\_date          maximum date specified by user

**Value**

df of animal data from specified meta

---

get_file_meta	<i>Generate metadata for a directory of animal data files</i>
---------------	---

---

**Description**

Generate metadata for a directory of animal data files

**Usage**

```
get_file_meta(data_dir)
```

**Arguments**

data_dir	directory of animal data files
----------	--------------------------------

**Value**

list of data info as a list of animal IDs and GPS units

**Examples**

```
# Get metadata for demo directory

get_file_meta(system.file("extdata", "demo_nov19", package = "animaltracker"))
```

---

get_meta	<i>Generate metadata for an animal data frame - filename, site, date min/max, animals, min/max lat/longitude, storage location</i>
----------	--

---

**Description**

Generate metadata for an animal data frame - filename, site, date min/max, animals, min/max lat/longitude, storage location

**Usage**

```
get_meta(df, file_id, file_name, site, ani_id, storage_loc)
```

**Arguments**

df	clean animal data frame
file_id	ID number of .csv source of animal data frame
file_name	.csv source of animal data frame
site	physical source of animal data
ani_id	ID of animal found in data frame
storage_loc	.rds storage location of animal data frame

**Value**

df of metadata for animal data frame

---

histogram\_animal\_elevation

*Generate a histogram of the distribution of modeled elevation - measured altitude*

---

**Description**

Generate a histogram of the distribution of modeled elevation - measured altitude

**Usage**

```
histogram_animal_elevation(datapts)
```

**Arguments**

datapts            GPS data with measured Altitude and computed Elevation data

**Value**

histogram of the distribution of modeled elevation - measured altitude

**Examples**

```
# Histogram of elevation - altitude for the demo data
histogram_animal_elevation(demo)
```

---

histogram\_time

*Generates a histogram to visualize the distribution of time between GPS measurements.*

---

**Description**

Generates a histogram to visualize the distribution of time between GPS measurements.

**Usage**

```
histogram_time(rds_path)
```

**Arguments**

rds\_path            Path of .rds cow data file to read in

**Value**

distribution of time between GPS measurements, as a histogram

**Examples**

```
# Histogram of GPS measurement time differences for demo data .rds
histogram_time(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

histogram_time_unit	<i>Generates a histogram to visualize the distribution of time between GPS measurements by GPS unit.</i>
---------------------	--

---

**Description**

Generates a histogram to visualize the distribution of time between GPS measurements by GPS unit.

**Usage**

```
histogram_time_unit(rds_path)
```

**Arguments**

rds\_path            Path of .rds animal data file to read in

**Value**

distribution of time between GPS measurements by GPS unit, as a histogram

**Examples**

```
# Histogram of GPS measurement time differences by GPS unit for demo data .rds
histogram_time_unit(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

join_summaries	<i>Joins two animal data frame summaries by a column and appends differences</i>
----------------	--

---

### Description

Joins two animal data frame summaries by a column and appends differences

### Usage

```
join_summaries(correct_summary, candidate_summary, by_str, daily = FALSE)
```

### Arguments

correct_summary	summary df of reference dataset, returned by summarise_anidf
candidate_summary	summary df of dataset to be compared to reference, returned by summarise_anidf
by_str	column to join by as a string, null if daily=TRUE
daily	whether to group by both GPS and Date for daily summary, defaults to False

### Value

df of joined summaries with differences

### Examples

```
# Join date summaries of unfiltered and filtered demo data
## Summarise unfiltered demo by date
unfiltered_summary <- summarise_anidf(demo_unfiltered_elev, Date, Latitude, Longitude,
Distance, Course, Rate, Elevation, daily=FALSE)

## Summarise filtered demo by date
filtered_summary <- summarise_anidf(demo_filtered_elev, Date, Latitude, Longitude,
Distance, Course, Rate, Elevation, daily=FALSE)

## Join
join_summaries(unfiltered_summary, filtered_summary, "Date", daily=FALSE)
```

---

line_compare	<i>Compares moving averages of a variable for two datasets over time, grouped by GPS GPS, Date, and col columns should match</i>
--------------	--

---

**Description**

Compares moving averages of a variable for two datasets over time, grouped by GPS GPS, Date, and col columns should match

**Usage**

```
line_compare(correct, candidate, col, export = FALSE, out = NULL)
```

**Arguments**

correct	reference df
candidate	df to be compared to the reference
col	variable to plot the moving average for
export	logical, whether to export plot, defaults to False
out	.png file name to save plot when export is True

**Value**

faceted line plot of moving averages over time grouped by GPS

**Examples**

```
# Faceted line plot comparing moving averages over time
# grouped by GPS for unfiltered and filtered demo data
## Set distance as the y axis
line_compare(demo_unfiltered, demo_filtered, Distance)
```

---

lookup_elevation_aws	<i>Add elevation data from public AWS terrain tiles to long/lat coordinates of animal gps data</i>
----------------------	--

---

**Description**

Add elevation data from public AWS terrain tiles to long/lat coordinates of animal gps data

**Usage**

```
lookup_elevation_aws(anidf, zoom = 11, get_slope = TRUE, get_aspect = TRUE)
```

**Arguments**

anidf	animal tracking dataframe
zoom	level of zoom, defaults to 11
get_slope	logical, whether to compute slope (in degrees), defaults to true
get_aspect	logical, whether to compute aspect (in degrees), defaults to true

**Value**

original data frame, with Elevation column appended

---

lookup\_elevation\_file *Add elevation data from terrain tiles to long/lat coordinates of animal gps data*

---

**Description**

Add elevation data from terrain tiles to long/lat coordinates of animal gps data

**Usage**

```
lookup_elevation_file(
  elev,
  anidf,
  zoom = 11,
  get_slope = TRUE,
  get_aspect = TRUE
)
```

**Arguments**

elev	elevation data as raster
anidf	animal tracking dataframe
zoom	level of zoom, defaults to 11
get_slope	logical, whether to compute slope (in degrees), defaults to true
get_aspect	logical, whether to compute aspect (in degrees), defaults to true

**Value**

original data frame, with terrain column(s) appended

---

process_elevation	<i>Process and optionally export modeled elevation data from existing animal data file</i>
-------------------	--

---

**Description**

Process and optionally export modeled elevation data from existing animal data file

**Usage**

```
process_elevation(
  zoom = 11,
  get_slope = TRUE,
  get_aspect = TRUE,
  in_path,
  export = FALSE,
  out_path = NULL
)
```

**Arguments**

zoom	level of zoom, defaults to 11
get_slope	logical, whether to compute slope (in degrees), defaults to True
get_aspect	logical, whether to compute aspect (in degrees), defaults to True
in_path	animal tracking data file to model elevation from
export	logical, whether to export data with elevation, defaults to False
out_path	.rds file path for processed data when export is True

**Value**

list of data frames with gps data augmented by elevation

---

qqplot_time	<i>Generates a QQ plot to show the distribution of time between GPS measurements.</i>
-------------	---

---

**Description**

Generates a QQ plot to show the distribution of time between GPS measurements.

**Usage**

```
qqplot_time(rds_path)
```

**Arguments**

rds\_path            Path of .rds animal data file to read in

**Value**

quantile-quantile plot to show distribution of time between GPS measurements

**Examples**

```
# QQ plot of GPS measurment time differences for demo data .rds
qqplot_time(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

quantile_time	<i>Determines the GPS measurement time value difference values roughly corresponding to quantiles with .05 intervals.</i>
---------------	---

---

**Description**

Determines the GPS measurement time value difference values roughly corresponding to quantiles with .05 intervals.

**Usage**

```
quantile_time(rds_path)
```

**Arguments**

rds\_path            Path of .rds animal data file to read in

**Value**

approximate time difference values corresponding to quantiles (.05 intervals)

**Examples**

```
# Read in .rds of demo data and calculate time difference quantiles
quantile_time(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

read_columbus	<i>Read and process a Columbus P-1 data file containing NMEA records into a data frame</i>
---------------	--

---

**Description**

Read and process a Columbus P-1 data file containing NMEA records into a data frame

**Usage**

```
read_columbus(filename)
```

**Arguments**

filename            path of Columbus P-1 data file

**Value**

NMEA records in RMC and GGA formats as a data frame

**Examples**

```
read_columbus(system.file("extdata", "demo_columbus.TXT", package = "animaltracker"))
```

---

read_gps	<i>Reads a GPS dataset of unknown format at location filename</i>
----------	---

---

**Description**

Reads a GPS dataset of unknown format at location filename

**Usage**

```
read_gps(filename)
```

**Arguments**

filename            location of the GPS dataset

**Value**

list containing the dataset as a df and the format

---

read\_zip\_to\_rasters     *Read an archive of altitude mask files and convert the first file into a raster object*

---

**Description**

Read an archive of altitude mask files and convert the first file into a raster object

**Usage**

```
read_zip_to_rasters(filename, exdir = "inst/extdata/elev")
```

**Arguments**

filename	path of altitude mask file archive
exdir	path to extract files

**Value**

the first altitude mask file as a raster object

---

run\_shiny\_animaltracker

*Run the animaltracker 'shiny' app by calling this function. Depending on the size of input files, it may be advisable to increase the maximum request size.*

---

**Description**

Run the animaltracker 'shiny' app by calling this function. Depending on the size of input files, it may be advisable to increase the maximum request size.

**Usage**

```
run_shiny_animaltracker(browser = TRUE, showcase = FALSE)
```

**Arguments**

browser	logical, whether to launch the app in your default browser (defaults to TRUE)
showcase	logical, whether to launch the app in 'showcase' mode (defaults to FALSE)

**Value**

None

---

run_validation_app	<i>Run the 'shiny' validation app. Depending on the size of input files, it may be advisable to increase the maximum request size.</i>
--------------------	--

---

**Description**

Run the 'shiny' validation app. Depending on the size of input files, it may be advisable to increase the maximum request size.

**Usage**

```
run_validation_app()
```

**Value**

None

---

save_meta	<i>Save metadata to a data frame and return it</i>
-----------	--

---

**Description**

Save metadata to a data frame and return it

**Usage**

```
save_meta(meta_df, file_meta)
```

**Arguments**

meta_df	the data frame to store metadata in
file_meta	meta for a .csv file generated by get_meta

**Value**

df of metadata

---

store_batch_list	<i>Generates basic metadata about a directory of animal data files and stores the files as data frames as a list with the meta</i>
------------------	--

---

**Description**

Generates basic metadata about a directory of animal data files and stores the files as data frames as a list with the meta

**Usage**

```
store_batch_list(data_dir)
```

**Arguments**

data_dir	location of animal data files, in list format
----------	---

**Value**

a list of animal data frames with information about the data

---

summarise_anidf	<i>Calculates summary statistics for an animal data frame</i>
-----------------	---

---

**Description**

Calculates summary statistics for an animal data frame

**Usage**

```
summarise_anidf(anidf, by, lat, long, dist, course, rate, elev, daily = FALSE)
```

**Arguments**

anidf	the animal data frame
by	column to group by, null if daily=TRUE
lat	latitude column
long	longitude column
dist	distance column
course	course column
rate	rate column
elev	elevation column
daily	whether to group by both GPS and Date for daily summary, defaults to False

**Value**

df of summary statistics for the animal data frame

**Examples**

```
# Summary of demo data by date
summarise_anidf(demo, Date, Latitude, Longitude, Distance, Course, Rate, Elevation, daily = FALSE)
```

---

summarise_col	<i>Get summary statistics for a single column in an animal data frame</i>
---------------	---

---

**Description**

Get summary statistics for a single column in an animal data frame

**Usage**

```
summarise_col(df, col)
```

**Arguments**

df	animal data frame
col	column to get summary stats for, as a string

**Value**

data frame of summary stats for col

**Examples**

```
# Get summary statistics for Distance column of demo data
summarise_col(demo, Distance)
```

---

summarise_unit	<i>Summarise a number of animal datasets by GPS unit</i>
----------------	--

---

**Description**

Summarise a number of animal datasets by GPS unit

**Usage**

```
summarise_unit(rds_path)
```

**Arguments**

rds\_path            Path of .rds cow data file to read in

**Value**

summary statistics for animals by GPS unit

**Examples**

```
# Read in .rds of demo data and summarise by GPS unit

summarise_unit(system.file("extdata", "demo_nov19.rds", package = "animaltracker"))
```

---

violin_compare	<i>Compares summary statistics from two datasets as side-by-side violin plots</i>
----------------	---

---

**Description**

Compares summary statistics from two datasets as side-by-side violin plots

**Usage**

```
violin_compare(df_summary, by, col_name, export = FALSE, out = NULL)
```

**Arguments**

df\_summary        data frame of summary statistics from both datasets to be compared  
 by                GPS or Date  
 col\_name         variable in df\_summary to be used for the y-axis, as a string  
 export            logical, whether to export plot, defaults to False  
 out               .png file name to save plot when export is True

**Value**

side-by-side violin plots

**Examples**

```
# Violin plot comparing unfiltered and filtered demo data summaries by date for a single variable
## Summarise unfiltered demo
unfiltered_summary <- summarise_anidf(demo_unfiltered_elev, Date, Latitude, Longitude,
Distance, Course, Rate, Elevation, daily=FALSE)

## Summarise filtered demo
filtered_summary <- summarise_anidf(demo_filtered_elev, Date, Latitude, Longitude,
Distance, Course, Rate, Elevation, daily=FALSE)

## Join
summary <- join_summaries(unfiltered_summary, filtered_summary, "Date", daily=FALSE)

## Violin plot

violin_compare(summary, Date, "meanElev")
```

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