

Package: anibehavr (via r-universe)

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Title Analyse Animal Behaviours

Version 0.1.0

Description What the package does (one paragraph).

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Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.1

Imports dplyr, forcats, janitor, magrittr, purrr, tidyr, tidyselect,
zoo, data.table, rlang, hms

Suggests ggplot2, testthat (>= 3.0.0)

Config/testthat/edition 3

URL <http://www.roald-arboel.com/anibehavr/>

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

RemoteUrl <https://github.com/roaldarbol/anibehavr>

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Contents

add_logticks	2
assign_video	3
classify_states	4
compress_observations	4
compress_observations2	5
correct_time	5
csv_clean	6
filter_classifications	6
filter_forward_backward	7
find_position	7
generete_state_numbers	8
identify_periods	8

inf_remove	9
inverse_ecdf	9
is_moving	10
join_timeseries	10
merge_videos	11
min_max_norm	11
mode_nonnumeric	12
read_align_all_data	12
read_align_data	13
read_switch_data	13
subset_around_event	14
sync_time	14

Index 15

add_logticks	<i>Add logticks</i>
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Description

Add logticks

Usage

```
add_logticks(
  base = 10,
  sides = "b1",
  scaled = TRUE,
  short = unit(0.1, "cm"),
  mid = unit(0.2, "cm"),
  long = unit(0.3, "cm"),
  colour = "black",
  size = 0.5,
  linetype = 1,
  alpha = 1,
  color = NULL,
  data = data.frame(x = NA),
  ...
)
```

Arguments

base	Base
sides	Sides
scaled	Scaled
short	Short
mid	Mid

long	Long
colour	Colour
size	Size
linetype	Linetype
alpha	Alpha
color	Color
data	Data
...	Other

Value

Added logticks

assign_video	<i>Assign video</i>
--------------	---------------------

Description

Assign video

Usage

```
assign_video(track_list, animal_ids, tracker = c("trex", "idtrackerai"))
```

Arguments

track_list	List from video files
animal_ids	Animal ids

Value

track_list

classify_states	<i>Classify states Across Time-scales</i>
-----------------	---

Description

Classify states Across Time-scales

Usage

```
classify_states(data, movement_var, window_widths, .keep = FALSE)
```

Arguments

data	Data frame
movement_var	Binary (1/0) variable to be used for the classification
window_widths	Window width for the filter
.keep	Keep both intermediate filter components (forward/backward)

Value

Data frame with classifications

compress_observations	<i>Compress observations</i>
-----------------------	------------------------------

Description

Get an average of your data where the output is n rows (approximately).

Arguments

data	Data
n_observations	Number of observations in output data frame
group	grouping variable

Value

Smaller data frame where values are means of N observations

`compress_observations2`*Compress observations2*

Description

Get an average of your data where the output is n rows (approximately).

Arguments

<code>data</code>	Data
<code>vars</code>	Variables to mean
<code>n_observations</code>	Number of observations in output data frame
<code>group</code>	grouping variable

Value

Smaller data frame where values are means of N observations

`correct_time`*Correct time*

Description

Correct time

Usage

```
correct_time(data)
```

Arguments

`data`

Value

A data frame where times are added.

`csv_clean`*csv_clean*

Description`csv_clean`**Usage**`csv_clean(df, tracker = c("trex", "idtrackerai"))`**Arguments**`df` Raw TRex csv file**Value**`df`

`filter_classifications`*Filter Classifications*

Description

Filter Classifications

Usage`filter_classifications(data, window_widths)`**Arguments**`data` Data frame`window_widths` Minimum duration of bout**Value**

Filtered data frame

filter_forward_backward
Forward-backward Filter

Description

Forward-backward Filter

Usage

```
filter_forward_backward(data, movement_var, window_width)
```

Arguments

data Data frame
group

find_position *Find position*

Description

Find position

Usage

```
find_position(df, exp_setup = c("wellplate", "tube"), animal_ids)
```

Arguments

df track_list of data frames.
exp_setup Experimental setup, eith "wellplate " or "tube".
animal_ids Vector of animal IDs, as strings.

Value

A single tibble

```
generete_state_numbers
```

Generate bout numbers for binary variable

Description

Generate bout numbers for binary variable

Usage

```
generete_state_numbers(data, var, .keep = FALSE)
```

Arguments

data	Data frame
var	Variable name (in quotes)
.keep	Keep both intermediate filter components (forward/backward)

Value

Data frame with classifications

```
identify_periods
```

IDENTIFY PERIODS OF INTEREST

Description

IDENTIFY PERIODS OF INTEREST

Usage

```
identify_periods(data, var, newvar, threshold = "minmax", as_factor = FALSE)
```

Arguments

data	Data frame
var	Variable to be identified
newvar	New variable name
threshold	Method for thresholding. Either "minmax", "mediansd" or a number.
as_factor	Whether an output should be returned as factor

Value

Thresholded data frame

inf_remove	<i>Title</i>
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Description

Title

Usage

```
inf_remove(df)
```

Arguments

df	Data frame
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Value

A data frame without Inf values

inverse_ecdf	<i>inverse_ecdf</i>
--------------	---------------------

Description

inverse_ecdf

Usage

```
inverse_ecdf(data, col, min_length, rest_state = FALSE)
```

Arguments

data	A data frame (e.g. a tibble)
col	Variable to make the density function for.
min_length	Minimal starting duration in frames
rest_state	Whether to make the ecdf for resting or active periods. TRUE or FALSE (default)

Value

Inverse ECDF

is_moving	<i>is_moving</i>
-----------	------------------

Description

is_moving

Usage

```
is_moving(df, type, n_frames = 50)
```

Arguments

df	Data frame.
type	Use mean or median.
n_frames	Number of frames used to generate rolling average.

Value

Data frame with a value for whether the animal is moving

join_timeseries	<i>Ensure that grouping levels match across data frames - add an error/warning if not.</i>
-----------------	--

Description

Ensure that grouping levels match across data frames - add an error/warning if not.

Usage

```
join_timeseries(
  .x,
  y,
  by = NULL,
  group = NULL,
  method = NA,
  copy = FALSE,
  suffix = c("", ".y"),
  keep = NULL
)
```

merge_videos	<i>Merge videos</i>
--------------	---------------------

Description

Merges data sets for consecutive videos for each animal.

Usage

```
merge_videos(df_list)
```

Arguments

df_list	List of data frames
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Value

Merged data frame from list

min_max_norm	<i>Min-max normalisation</i>
--------------	------------------------------

Description

Min-max normalisation

Usage

```
min_max_norm(x)
```

Arguments

x

Value

Normalisation

mode_nonnumeric	<i>Non-numeric mode</i>
-----------------	-------------------------

Description

Non-numeric mode

Usage

mode_nonnumeric(x)

Arguments

x	Vector to find the mode over
---	------------------------------

Value

Mode

read_align_all_data	<i>Title</i>
---------------------	--------------

Description

Title

Usage

read_align_all_data(filenamees, bin_seconds)

Arguments

filenamees	Filenamees
bin_seconds	Bin duration in seconds

Value

Aligned data frame

read_align_data	<i>Title</i>
-----------------	--------------

Description

Title

Usage

```
read_align_data(data, bin_seconds, time, animal_id)
```

Arguments

bin_seconds	Bin duration in seconds
filename	Filenames

Value

Aligned data frame

read_switch_data	<i>Title</i>
------------------	--------------

Description

Title

Usage

```
read_switch_data(filenamees, bin_seconds, time_radius = FALSE)
```

Arguments

filenamees	Filenames
bin_seconds	Bin duration in seconds
time_radius	Radius around change

Value

Switch data

subset_around_event *Subset Around Event*

Description

Subset Around Event

Usage

```
subset_around_event(data, var, from, time_var, time_radius = FALSE)
```

Arguments

data	Data frame
var	Variable to center around
from	Reference point (e.g. "Day"/"Night")
time_var	Which variable represents time
time_radius	Time around point

Value

Subset of data frame around point in time

sync_time *Sync time*

Description

Sync time

Usage

```
sync_time(.data, data2, x, y)
```

Arguments

.data	Data frame or tibble
data2	The second data frame
x	Parameter used to merge (often time)
y	Variable to be synced along with x

Value

Tibble

Index

[add_logticks](#), 2
[assign_video](#), 3

[classify_states](#), 4
[compress_observations](#), 4
[compress_observations2](#), 5
[correct_time](#), 5
[csv_clean](#), 6

[filter_classifications](#), 6
[filter_forward_backward](#), 7
[find_position](#), 7

[generete_state_numbers](#), 8

[identify_periods](#), 8
[inf_remove](#), 9
[inverse_ecdf](#), 9
[is_moving](#), 10

[join_timeseries](#), 10

[merge_videos](#), 11
[min_max_norm](#), 11
[mode_nonnumeric](#), 12

[read_align_all_data](#), 12
[read_align_data](#), 13
[read_switch_data](#), 13

[subset_around_event](#), 14
[sync_time](#), 14