

Package ‘sleepr’

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Title Analyse activity and sleep behaviour

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Version 0.0.0.9000

Description

Use behavioural variables to score activity and infer sleep from contiguous periods of immobility.

Depends R (>= 2.10)

Imports behavr,
data.table

Suggests testthat

License GPL-3

Encoding UTF-8

LazyData true

URL <https://github.com/rethomics/sleepr>

BugReports <https://github.com/rethomics/sleepr/issues>

RoxygenNote 6.0.1

Roxygen list(markdown = TRUE)

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bout_analysis	<i>Find "bouts" in categorical time series</i>
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Description

This function is used to find contiguous regions of unique value in a – potentially irregular/heterogeneous – univariate categorical time series.

Usage

```
bout_analysis(var, data)
```

Arguments

var name of the variable to use from data

data [data.table](#) containing behavioural variable from or one multiple animals. When it has a key, unique values, are assumed to represent unique individuals (e.g. in a [behavr](#) table). Otherwise, it analysis the data as coming from a single animal. data must have a column `t` representing time.

Value

an object of the same type as data (i.e. [data.table::data.table](#) or [behavr::behavr](#)). Each row is a specific bout characterised by three columns.

- `t` – its *onset*
- `duration` – its length
- `<var>` – a column with the same name as `var`. The value of `var` for this bout.

See Also

[todo](#)

Examples

```
#TODO
```

motion_detectors

Motion detector for Ethocope data

Description

Defines whether a *single animal* is moving according to:

Usage

```
max_velocity_detector(data, time_window_length,
  velocity_correction_coef = 0.003, masking_duration = 6)

max_velocity_detector_legacy(data, velocity_threshold = 0.006)

virtual_beam_cross_detector(data, time_window_length)
```

Arguments

data	data.table::data.table containing behavioural variables of <i>a single animal</i> (no id). It must have the columns xy_dist_log10x1000(for computing subpixel velocity), x(beam cross), t and has_interacted (whether a stimulus was delivered).
time_window_length	number of seconds to be used by the motion classifier. This corresponds to the sampling period of the output data.
velocity_correction_coef	an empirical coefficient to correct velocity with respect to variable framerate.
masking_duration	number of second during which any movement is ignored (velocity is set to 0) after a stimulus is delivered (aka interaction).
velocity_threshold	uncorrected velocity above which an animal is classified as ‘moving’ (for the legacy version).

Details

- Validated and corrected subpixel velocity ([max_velocity_detector](#)), the most rigorous
- Uncorrected subpixel velocity ([max_velocity_detector_legacy](#))
- Crossing a virtual beam in the middle of the region of interest ([virtual_beam_cross_detector](#))

[max_velocity_detector](#) is the default movement classification for real-time ethoscope experiments. It is benchmarked against human-generated ground truth.

These functions are *rarely called directly*, but typically used is in the context of [sleep_annotation](#).

Value

an object of the same type as data (i.e. [data.table::data.table](#) or [behavr::behavr](#)) with additional columns:

- moving Logical, TRUE iff. motion was detected.
- beam_crosses The number of beam crosses (when the animal crosses $x = 0.5$ – that is the midpoint of the region of interest) within the time window
- max_velocity The maximal velocity within the time window. The resulting data is sampled at a period equals to time_window_length.

See Also

TODO

- [sleep_annotation](#) – which requires a motion detector

sleep_annotation	<i>Score sleep behaviour from immobility</i>
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Description

This function first uses a motion classifier to decide whether an animal is moving during a given time window. Then, it defines sleep as contiguous immobility for a minimal duration.

Usage

```
sleep_annotation(data, time_window_length = 10, min_time_immobile = 300,
  motion_detector_FUN = max_velocity_detector, ...)

sleep_dam_annotation(data, time_window_length = 60, min_time_immobile = 300)
```

Arguments

data	data.table containing behavioural variable from or one multiple animals. When it has a key, unique values, are assumed to represent unique individuals (e.g. in a behavr table). Otherwise, it analysis the data as coming from a single animal. data must have a column t representing time.
time_window_length	number of seconds to be used by the motion classifier. This corresponds to the sampling period of the output data.
min_time_immobile	Minimal duration (in s) of a sleep bout. Immobility bouts longer or equal to this value are considered as sleep.
motion_detector_FUN	function used to classify movement
...	extra arguments to be passed to <code>motion_classifier_FUN</code> .

Details

The default `time_window_length` is 300 seconds also known as the "5 minute rule". `sleep_annotation` is typically used for ethoscope data, whilst `sleep_dam_annotation` only works on DAM2 data. These functions are *rarely used directly*, but rather passed as an argument to a data loading function, so that analysis can be performed on the go.

Value

a [behavr](#) table similar to data with additional variables/annotations (i.e. moving and asleep). The resulting data will only have one data point every `time_window_length` seconds.

See Also

- [motion_detectors](#) – options for the `motion_detector_FUN` argument
- [bout_analysis](#) – to further analyse sleep bouts in terms of onset and length

Examples

```
#todo
```

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