

# Package ‘Sojourn’

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**Type** Package

**Title** Apply Sojourn Methods for Processing ActiGraph Accelerometer Data

**Version** 1.2.1

**Depends** R (>= 3.1.0)

**Description** Provides a simple way for utilizing Sojourn methods for accelerometer processing, as detailed in Lyden K, Keadle S, Staudenmayer J, & Freedson P (2014) <[doi:10.1249/MSS.0b013e3182a42a2d](https://doi.org/10.1249/MSS.0b013e3182a42a2d)>, Ellingson LD, Schwabacher IJ, Kim Y, Welk GJ, & Cook DB (2016) <[doi:10.1249/MSS.0000000000000915](https://doi.org/10.1249/MSS.0000000000000915)>, and Hibbing PR, Ellingson LD, Dixon PM, & Welk GJ (2018) <[doi:10.1249/MSS.0000000000001486](https://doi.org/10.1249/MSS.0000000000001486)>.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**Imports** dplyr (>= 0.7), magrittr (>= 1.5), lubridate (>= 1.7.4), nnet (>= 7.3), PAutilities (>= 0.2.0), rlang (>= 0.2), stats, svDialogs (>= 1.0), utils, zoo (>= 1.8)

**RoxygenNote** 7.1.2

**URL** <https://github.com/paulhibbing/Sojourn>

**BugReports** <https://github.com/paulhibbing/Sojourn/issues>

**Suggests** data.table, testthat

**NeedsCompilation** no

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## Contents

apply_youth_sojourn	2
compute.bouts.info	3
enhance_actigraph	4
example_data	4
get_youth_sojourns	5
input_demographic	6
read_AP	7
SIP_ag	7
SIP_ap	8
Sojourn	9
sojourn_3x_SIP	9
soj_1x_original	10
soj_3x_original	11
<b>Index</b>	<b>12</b>

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apply_youth_sojourn	<i>Apply the youth Sojourn method</i>
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## Description

Function for using the youth Sojourn method developed by [Hibbing et al. \(2018\)](#)

## Usage

```
apply_youth_sojourn(
  AG,
  vm = c("Vector.Magnitude", "ENMO"),
  Site = c("Hip", "Wrist"),
  demo_interactive = FALSE,
  verbose = FALSE,
  ...
)
```

## Arguments

AG	a data frame of monitor and demographic data
vm	the variable to use for processing, either "Vector.Magnitude" (for activity counts) or "ENMO" (for raw acceleration)
Site	the wear location of the monitor, either "Hip" or "Wrist"
demo_interactive	logical. Input demographics interactively if missing variables are identified during format checking?
verbose	logical. Print processing updates to the console?
...	Further arguments passed to <a href="#">youth_name_test</a>

**Value**

The original data frame, plus additional predictions made by the Sojourn method

**Note**

The functions `AGread::read_AG_counts` and `AGread::read_AG_raw` are recommended for assembling the monitor-specific portion of the AG data frame.

**Examples**

```
data(example_data, package = "Sojourn")

results_youth_soj <- apply_youth_sojourn(
  AG = example_data,
  vm = "Vector.Magnitude",
  Site = "Hip"
)
utils::head(results_youth_soj)
```

---

`compute.bouts.info`      *Summarize outcomes from data processed using the Sojourn method*

---

**Description**

A function to summarize predictions made by the original Sojourn method of [Lyden et al. \(2014\)](#).

**Usage**

```
compute.bouts.info(est.mets, units = c("secs", "mins"))
```

**Arguments**

<code>est.mets</code>	numeric vector of predicted metabolic equivalents
<code>units</code>	time units associated with each row of data

**Value**

a data frame summarizing the predictions made by the Sojourn method.

**Examples**

```
data(example_data, package = "Sojourn")
example_data <- soj_3x_original(
  example_data$axis1,
  example_data$axis2,
  example_data$axis3,
  example_data$Vector.Magnitude
```

```
)  
compute.bouts.info(example_data$METs)
```

---

enhance\_actigraph      *Combine ActiGraph and activPAL data*

---

### Description

Merge data streams for separate monitors in the **SIP method**

### Usage

```
enhance_actigraph(ag, ap, verbose = FALSE)
```

### Arguments

ag	ActiGraph data
ap	activPAL data
verbose	logical. Print information to console?

### Value

data frame of combined ActiGraph and activPAL data

### Examples

```
data(SIP_ag, package = "Sojourn")  
data(SIP_ap, package = "Sojourn")  
combined_data <- enhance_actigraph(SIP_ag, SIP_ap)  
utils::head(combined_data)
```

---

example\_data      *Sample data for exploring original and youth Sojourn methods*

---

### Description

Sample data for exploring original and youth Sojourn methods

### Usage

```
example_data
```

**Format**

A data frame with 79989 rows and 11 variables:

**id** An example ID for the data set

**Sex** An example sex for the data set

**Age** An example age (in years) for the data set

**BMI** An example body mass index (in kg/m<sup>2</sup>) for the data set

**Timestamp** POSIX-formatted variable giving the timestamp for each observation

**axis1** Activity counts from the first axis

**axis2** Activity counts from the second axis

**axis3** Activity counts from the third axis

**Vector.Magnitude** Vector magnitude of activity counts ( $\sqrt{\text{sum}(\text{axis1}^2, \text{axis2}^2, \text{axis3}^2)}$ )

**steps** Predicted steps taken

**incline** Inclinometer status (0 = off, 1 = lying, 2 = sitting, 3 = standing)

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get_youth_sojourns	<i>Label Sojourns in a data stream according to the youth-specific algorithm</i>
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**Description**

Identify Sojourns using the algorithm of [Hibbing et al. \(2018\)](#)

**Usage**

```
get_youth_sojourns(
  vm,
  short = 30,
  Output = c("Counts", "Raw"),
  Site = c("Hip", "Wrist"),
  epoch = 1,
  difference = 15,
  threshold = 100,
  verbose = FALSE
)
```

**Arguments**

vm	vector of triaxial accelerometer values, either the vector magnitude for activity counts, or the Euclidian Norm Minus One for raw acceleration
short	numeric scalar. Shortest allowable duration for a Sojourn. Should be detected automatically from the internally-stored grid search values for the attachment site (hip or wrist) and data type (counts or raw)

Output	the data type (counts or raw)
Site	the attachment site (hip or wrist)
epoch	the epoch length, in seconds
difference	the difference parameter
threshold	the threshold parameter
verbose	logical. Print processing updates to the console?

**Value**

A data frame (with nrow equal to length(vm)) that gives sojourn labels and durations

**Examples**

```
data(example_data, package = "Sojourn")
get_youth_sojourns(example_data$Vector.Magnitude,
  Output = "Counts", Site = "Hip")
```

---

input\_demographic      *Interactively input demographic information*

---

**Description**

Interactively input demographic information

**Usage**

```
input_demographic(...)
```

**Arguments**

...                      Further arguments passed to svDialogs functions

**Value**

A data frame containing an ID, attachment site of the monitor (hip or wrist), and the participant's sex, age, and BMI

**Examples**

```
if (interactive()) {
  input_demographic()
}
```

---

read_AP	<i>Read an activPAL events file</i>
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**Description**

Read an activPAL events file

**Usage**

```
read_AP(filename, tz = "UTC")
```

**Arguments**

filename	character. Path to the file
tz	character. The timezone to use

**Value**

Data frame reflecting the data contained in filename.

**Note**

There must be a corresponding `.def` file located in the same directory as filename

**Examples**

```
ap_file <- system.file(  
  "extdata/sampled_data_Events.csv",  
  package = "Sojourn"  
)  
if (isTRUE(requireNamespace("data.table"))) {  
  ap_data <- read_AP(ap_file)  
  utils::head(ap_data)  
}
```

---

SIP_ag	<i>ActiGraph sample data for exploring Sojourns Including Posture (SIP) method.</i>
--------	---

---

**Description**

ActiGraph sample data for exploring Sojourns Including Posture (SIP) method.

**Usage**

```
SIP_ag
```

**Format**

A data frame with 12257 rows and 5 variables:

**counts** Activity counts from the first axis

**axis2** Activity counts from the second axis

**axis3** Activity counts from the third axis

**vm** Vector magnitude of activity counts

**Time** POSIX-formatted variable giving the timestamp for each observation

---

SIP_ap	<i>activPAL sample data for exploring Sojourns Including Posture (SIP) method</i>
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---

**Description**

activPAL sample data for exploring Sojourns Including Posture (SIP) method

**Usage**

SIP\_ap

**Format**

A data frame with 12257 rows and 5 variables:

**Time** POSIX-formatted variable giving the timestamp for each observation

**DataCount** Integer value giving the index of the sample from which the row of data is drawn

**Interval** Duration (in seconds) of the interval from one data point to the next

**ActivityCode** Integer giving the posture activity classification: 0 is sedentary, 1 is standing, and 2 is stepping

**CumulativeStepCount** Integer giving the cumulative step count from the start of the file to the current data point.

**ActivityScore** Numeric giving MET-hours

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Sojourn	<i>Sojourn: Apply Sojourn Methods for Processing ActiGraph Accelerometer Data</i>
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### Description

The Sojourn package provides a convenient way to apply the family of Sojourn methods published in previous works including Lyden K, Keadle S, Staudenmayer J, & Freedson P (2014) <doi:10.1249/MSS.0b013e3182a42a2d>, Ellingson LD, Schwabacher IJ, Kim Y, Welk GJ, & Cook DB (2016) <doi:10.1249/MSS.0000000000000915>, and Hibbing PR, Ellingson LD, Dixon PM, & Welk GJ (2018) <doi:10.1249/MSS.0000000000001486>.

### Details

It is meant for use with data from ActiGraph monitors and (in the case of Sojourns Including Posture, by Ellingson et al. (2016)) activPAL monitors. File reading is not included in the functionality of the Sojourn package. For help with that preliminary step, users are directed to the packages AGread (for ActiGraph files) and activpalProcessing (for activPAL files).

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sojourn_3x_SIP	<i>Triaxial Sojourn method for the SIP method</i> <a href="https://pubmed.ncbi.nlm.nih.gov/27015380/SIP_method">Rhrefhttps://pubmed.ncbi.nlm.nih.gov/27015380/SIP method</a>
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### Description

Triaxial Sojourn method for the **SIP method**

### Usage

```
sojourn_3x_SIP(ag, short = 30)
```

### Arguments

ag	combined ActiGraph and activPAL data on which to identify transitions and make predictions
short	the minimum duration of a qualifying Sojourn

### Value

A data frame of processed data using the **SIP method**

### Examples

```
data(SIP_ag, package = "Sojourn")
data(SIP_ap, package = "Sojourn")
data <- Sojourn::enhance_actigraph(SIP_ag, SIP_ap)
utils::head(sojourn_3x_SIP(data))
```

---

soj_1x_original	<i>Invoke the original uni-axial Sojourn method</i>
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---

### Description

Calls the uni-axial Sojourn method from [Lyden et al. \(2014\)](#).

### Usage

```
soj_1x_original(  
  counts,  
  perc.cut = 0.05,  
  perc.cut.2 = 0.12,  
  perc.cut.3 = 0.55,  
  too.short = 10,  
  sit.cut = 90,  
  long.soj = 120  
)
```

### Arguments

counts	numeric vector of vertical axis counts
perc.cut	decision tree threshold 1
perc.cut.2	decision tree threshold 2
perc.cut.3	decision tree threshold 3
too.short	minimum length of one Sojourn
sit.cut	cut-point for determining sitting
long.soj	maximum length of one Sojourn

### Value

a data frame of processed data

### Examples

```
data(example_data, package = "Sojourn")  
results_1x <- soj_1x_original(example_data$axis1)  
utils::head(results_1x)
```

---

soj_3x_original	<i>Invoke the original triaxial Sojourn method</i>
-----------------	--

---

**Description**

Calls the triaxial Sojourn method from [Lyden et al. \(2014\)](#).

**Usage**

```
soj_3x_original(  
  counts,  
  counts.2,  
  counts.3,  
  vect.mag,  
  short = 30,  
  verbose = FALSE  
)
```

**Arguments**

counts	numeric vector of activity counts from the first axis
counts.2	numeric vector of activity counts from the second axis
counts.3	numeric vector of activity counts from the third axis
vect.mag	vector magnitude of the activity counts
short	minimum length of one Sojourn
verbose	logical. Print updates to console?

**Value**

a data frame of processed data

**Examples**

```
data(example_data, package = "Sojourn")  
results_3x <- soj_3x_original(  
  example_data$axis1,  
  example_data$axis2,  
  example_data$axis3,  
  example_data$Vector.Magnitude  
)  
  
utils::head(results_3x)
```

# Index

## \* datasets

example\_data, 4

SIP\_ag, 7

SIP\_ap, 8

apply\_youth\_sojourn, 2

compute.bouts.info, 3

enhance\_actigraph, 4

example\_data, 4

get\_youth\_sojourns, 5

input\_demographic, 6

read\_AP, 7

SIP\_ag, 7

SIP\_ap, 8

soj\_1x\_original, 10

soj\_3x\_original, 11

Sojourn, 9

sojourn\_3x\_SIP, 9

youth\_name\_test, 2