

Package ‘stepcount’

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Title Estimate Step Counts from 'Accelerometry' Data

Version 0.6.0

Description Interfaces the 'stepcount' Python module <<https://github.com/OxWearables/stepcount>> to estimate step counts and other activities from 'accelerometry' data.

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Depends R (>= 4.1.0)

Imports assertthat, curl, lubridate, magrittr, methods, readr, reticulate (>= 1.42.0), rlang

Suggests tidyr, dplyr, ggplot2, testthat (>= 3.0.0), spelling, callr

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have_stepcount	<i>Check the stepcount Python Module</i>
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Description

Check the stepcount Python Module

Usage

have_stepcount()

stepcount_check()

stepcount_version()

Value

A logical value indicating whether the stepcount Python module is available.

Examples

```
if (have_stepcount()) {  
    stepcount_version()  
}
```

py_require_stepcount	<i>Command for py_require for stepcount</i>
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Description

Command for py_require for stepcount

Usage

py_require_stepcount(...)

Arguments

... arguments to pass to `reticulate::py_require()`

Value

A logical value indicating whether the package is available.

py_stepcount	<i>Perform step count calculation in a separate Python environment</i>
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Description

Perform step count calculation in a separate Python environment

Usage

```
py_stepcount(  
  ...,  
  pyenv_function = function() {  
    stepcount::py_require_stepcount()  
  },  
  show = TRUE  
)
```

Arguments

...	arguments passed to stepcount()
pyenv_function	function that loads the stepcount Python package. By default, it uses py_require_stepcount . If this function has an args argument, the output of pyenv_function will be re-assigned to args.
show	Logical, whether to show the standard output on the screen while the child process is running, passed to callr::r()

Value

The output from [stepcount\(\)](#). A tibble with minute-level time, steps, and walking columns.

Examples

```
file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")  
if (stepcount_check()) {  
  df = readr::read_csv(file)  
  out = try({py_stepcount(df, sample_rate = 100)})  
}
```

sc_load_model	<i>Load Stepcount Model</i>
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Description

Load Stepcount Model

Usage

```
sc_load_model(
    model_type = c("ssl", "rf"),
    model_path = NULL,
    check_md5 = TRUE,
    force_download = FALSE,
    as_python = TRUE
)

sc_model_filename(model_type = c("ssl", "rf"))

sc_download_model(
    model_path,
    model_type = c("ssl", "rf"),
    check_md5 = TRUE,
    ...
)
```

Arguments

model_type	type of the model: either random forest (rf) or Self-Supervised Learning model (ssl)
model_path	the file path to the model. If on disk, this can be re-used and not re-downloaded. If NULL, will download to the temporary directory
check_md5	Do a MD5 checksum on the file
force_download	force a download of the model, even if the file exists
as_python	Keep model object as a python object
...	for sc_download_model, additional arguments to pass to curl::curl_download()

Value

A model from Python. sc_download_model returns a model file path.

sc_model_params	<i>Run Stepcount Model on Data</i>
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Description

Run Stepcount Model on Data

Usage

```
sc_model_params(model_type, pytorch_device)
```

```
stepcount(
    file,
    sample_rate = NULL,
    model_type = c("ssl", "rf"),
    model_path = NULL,
    pytorch_device = c("cpu", "cuda:0"),
    verbose = TRUE,
    keep_data = FALSE
)
```

```
stepcount_with_model(
    file,
    model_type = c("ssl", "rf"),
    model,
    sample_rate = NULL,
    pytorch_device = c("cpu", "cuda:0"),
    verbose = TRUE,
    keep_data = FALSE
)
```

Arguments

model_type	type of the model: either random forest (rf) or Self-Supervised Learning model (ssl)
pytorch_device	device to use for prediction for PyTorch.
file	accelerometry file to process, including CSV, CWA, GT3X, and GENEActiv bin files
sample_rate	the sample rate of the data. Set to NULL for stepcount to try to guess this
model_path	the file path to the model. If on disk, this can be re-used and not re-downloaded. If NULL, will download to the temporary directory
verbose	print diagnostic messages
keep_data	should the data used in the prediction be in the output?
model	A model object loaded from sc_load_model, but as_python must be TRUE

Value

A list of the results (`data.frame`), summary of the results, adjusted summary of the results, and information about the data.

Examples

```
library(magrittr)
file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")
if (stepcount_check()) {
  out = try({stepcount(file = file)})
  st = out$step_times
}

file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")
df = readr::read_csv(file)
if (stepcount_check()) {
  out = stepcount(file = df)
  st = out$step_times
}
if (requireNamespace("ggplot2", quietly = TRUE) &&
    requireNamespace("tidyr", quietly = TRUE) &&
    requireNamespace("dplyr", quietly = TRUE)) {
  dat = df[10000:12000,] |>
    dplyr::select(-dplyr::any_of("annotation")) |>
    tidyr::gather(axis, value, -time)
  st = st |>
    dplyr::mutate(time = lubridate::as_datetime(time)) |>
    dplyr::as_tibble()
  st = st |>
    dplyr::filter(time >= min(dat$time) & time <= max(dat$time))
  dat |>
    ggplot2::ggplot(ggplot2::aes(x = time, y = value, colour = axis)) +
    ggplot2::geom_line() +
    ggplot2::geom_vline(data = st, ggplot2::aes(xintercept = time))
}
```

sc_read

Read a Data Set for stepcount

Description

Read a Data Set for stepcount

Usage

```
sc_read(  
  file,  
  sample_rate = NULL,  
  resample_hz = "uniform",  
  verbose = TRUE,  
  keep_pandas = FALSE  
)
```

Arguments

file	path to the file for reading
sample_rate	the sample rate of the data. Set to NULL for stepcount to try to guess this
resample_hz	Target frequency (Hz) to resample the signal. If "uniform", use the implied frequency (use this option to fix any device sampling errors). Pass NULL to disable. Defaults to "uniform".
verbose	print diagnostic messages
keep_pandas	do not convert the data to a data.frame and keep as a pandas data.frame

Value

A list of the data and information about the data

Note

The data P30_wrist100 is from <https://ora.ox.ac.uk/objects/uuid:19d3cb34-e2b3-4177-91b6-1bad0e0163e7>, where we took the first 180,000 rows, the first 30 minutes of data from that participant as an example.

Examples

```
file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")  
if (stepcount_check()) {  
  out = sc_read(file)  
}  
  
## Not run:  
file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")  
if (stepcount_check()) {  
  out = sc_read(file, sample_rate = 100L)  
}  
  
## End(Not run)
```

sc_rename_data	<i>Rename data for Stepcount</i>
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Description

Rename data for Stepcount

Usage

```
sc_rename_data(data)
```

```
sc_write_csv(data, path = tempfile(fileext = ".csv"))
```

Arguments

data	a <code>data.frame</code> of raw accelerometry
path	path to the CSV output file

Value

A `data.frame` of renamed columns

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