Package 'stepcount'

October 2, 2024

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Title Estimate Step Counts from 'Accelerometry' Data
Version 0.3.2
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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```
conda_create_walking_env
```

Create Conda Environment for Walking

Description

Create Conda Environment for Walking

Usage

```
conda_create_walking_env(envname = "stepcount", ...)
```

Arguments

```
envname environment name
```

... additional arguments to pass to reticulate::conda_create()

Value

Output of reticulate::conda_create

install_stepcount

Install the stepcount *Python Module*

Description

Install the stepcount Python Module

Usage

```
install_stepcount(packages = "stepcount", ...)
have_stepcount()
stepcount_check()
stepcount_version()
```

Arguments

```
packages
    packages to install. If stepcount is not included, it will be added. This package
    is known to work with stepcount==3.2.4
...
Additional arguments to pass to reticulate::py_install(), other than pip
    (pip = TRUE enforced)
```

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Value

Output of reticulate::py_install

Examples

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

sc_load_model

Load Stepcount Model

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()
```

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Value

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

sc_model_params

Run Stepcount Model on Data

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

mode1

type of the model: either random forest (rf) or Self-Supervised Learning model model_type (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 the file path to the model. If on disk, this can be re-used and not re-downloaded. model_path If NULL, will download to the temporary directory print diagnostic messages verbose keep_data should the data used in the prediction be in the output?

A model object loaded from sc_load_model, but as_python must be TRUE

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Value

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

Examples

```
file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")
if (stepcount_check()) {
 out = stepcount(file = file)
 st = out$step_times
}
## Not run:
 file = system.file("extdata/P30_wrist100.csv.gz", package = "stepcount")
 df = readr::read_csv(file)
 if (stepcount_check()) {
   out = stepcount(file = df)
   st = out$step_times
 if (requireNamespace("ggplot2", quietly = TRUE) &&
     requireNamespace("tidyr", quietly = TRUE) &&
     requireNamespace("dplyr", quietly = TRUE)) {
   dat = df[10000:12000,] \%>\%
     dplyr::select(-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))</pre>
   dat %>%
     ggplot2::ggplot(ggplot2::aes(x = time, y = value, colour = axis)) +
     ggplot2::geom_line() +
     ggplot2::geom_vline(data = st, ggplot2::aes(xintercept = time))
 }
## End(Not run)
```

sc_read

Read a Data Set for stepcount

Description

Read a Data Set for stepcount

Usage

```
sc_read(
  file,
```

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```
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 fre-

quency (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

Rename data for Stepcount

Description

Rename data for Stepcount

```
use_stepcount_condaenv
```

Usage

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

Arguments

data a data.frame of raw accelerometry

path path to the CSV output file

Value

A data.frame of renamed columns

```
use_stepcount_condaenv
```

Use Conda Environment for stepcount

Description

Use Conda Environment for stepcount

Usage

```
use_stepcount_condaenv(envname = "stepcount", ...)
conda_create_stepcount(envname = "stepcount", ..., python_version = "3.9")
unset_reticulate_python()
have_stepcount_condaenv()
```

Arguments

```
envname environment name for the conda environment
... additional arguments to pass to reticulate::use_condaenv() other than condaenv.

python_version version of Python to use for environment
```

Value

Nothing

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