# Package 'GGIR' 

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## Type Package <br> Title Raw Accelerometer Data Analysis

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Maintainer Vincent $T$ van Hees < $v$. vanhees@accelting. com>
Description A tool to process and analyse data collected with wearable raw acceleration sen-
sors as described in Migueles and colleagues (JMPB 2019), and van Hees and colleagues (JApplPhysiol 2014; PLoSONE 2015). The package has been developed and tested for binary data from 'GENEActiv' <https: //activinsights.com/>, binary (.gt3x) and .csvexport data from 'Actigraph' [https://theactigraph.com](https://theactigraph.com) devices, and binary (.cwa) and .csv-export data from 'Axivity' <https: //axivity.com>. These devices are currently widely used in research on human daily physical activity. Further, the package can handle accelerometer data file from any other sensor brand providing that the data is stored in csv format. Also the package allows for external function embedding.

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https://wadpac.github.io/GGIR/
BugReports https://github.com/wadpac/GGIR/issues
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Author Vincent T van Hees [aut, cre],
Jairo H Migueles [aut] ([https://orcid.org/0000-0003-0366-6935](https://orcid.org/0000-0003-0366-6935)),
Severine Sabia [ctb],
Matthew R Patterson [ctb],
Zhou Fang [ctb],

Joe Heywood [ctb],<br>Joan Capdevila Pujol [ctb],<br>Lena Kushleyeva [ctb],<br>Mathilde Chen [ctb],<br>Manasa Yerramalla [ctb],<br>Patrick Bos [ctb] ([https://orcid.org/0000-0002-6033-960X](https://orcid.org/0000-0002-6033-960X)),<br>Taren Sanders [ctb],<br>Chenxuan Zhao [ctb],<br>Medical Research Council UK [cph, fnd],<br>Accelting [cph, fnd],<br>French National Research Agency [cph, fnd]<br>Repository CRAN<br>Date/Publication 2024-07-02 15:50:05 UTC

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## Description

Disclaimer: If you are a new GGIR user then please see the GGIR github-pages for a narrative overview of GGIR.

This document is primarily aimed at documenting the functions and their input arguments.
Please note that there is google discussion group for this package (link below).
You can thank us for sharing the code in this package and for developing it as a generic purpose tool by citing the package name and by citing the supporting publications (e.g. Migueles et al. 2019) in your publications.

## Details

| Package: | GGIR |
| :--- | :--- |
| Type: | Package |
| Version: | $3.1-2$ |
| Date: | $2024-07-02$ |
| License: | Apache License $(==2.0)$ |
| Discussion group: | https://groups.google.com/forum/\#!forum/rpackageggir |

## Author(s)

- Vincent T van Hees < v.vanhees @accelting.com> main creator and developer
- Zhou Fang developed calibration algorithm used in function g.calibrate
- Joe Heywood helped develop the functionality to process specific recording days
- Severine Sabia, Mathilde Chen, and Manasa Yerramalla extensively tested and provided feedback on various functions
- Joan Capdevila Pujol helped to improve various functions
- Jairo H Migueles [jairohm@ugr.es](mailto:jairohm@ugr.es) helped to improve various functions
- Matthew R Patterson helped with enhancing the visual report.
- Lena Kushleyeva helped fix bug in sleep detection.
- Taren Sanders helped tidy up the parallel processing functionality


## References

- Migueles JH, Rowlands AV, et al. GGIR: A Research Community-Driven Open Source R Package for Generating Physical Activity and Sleep Outcomes From Multi-Day Raw Accelerometer Data. Journal for the Measurement of Physical Behaviour. 2(3) 2019. doi:10.1123/jmpb.20180063.
- van Hees VT, Gorzelniak L, Dean Leon EC, Eder M, Pias M, et al. (2013) Separating Movement and Gravity Components in an Acceleration Signal and Implications for the Assessment of Human Daily Physical Activity. PLoS ONE 8(4): e61691. doi:10.1371/journal.pone. 0061691
- van Hees VT, Fang Z, Langford J, Assah F, Mohammad A, da Silva IC, Trenell MI, White T, Wareham NJ, Brage S. Auto-calibration of accelerometer data for free-living physical activity assessment using local gravity and temperature: an evaluation on four continents. J Appl Physiol (1985). 2014 Aug 7
- van Hees VT, Sabia S, et al. (2015) A novel, open access method to assess sleep duration using a wrist-worn accelerometer, PLoS ONE, November 2015


## Examples

```
    ## Not run:
        #inspect file:
        I = g.inspectfile(datafile)
    #autocalibration:
    C = g.calibrate(datafile)
    #get meta-data:
    M = g.getmeta(datafile)
## End(Not run)
    data(data.getmeta)
    data(data.inspectfile)
    data(data.calibrate)
    #impute meta-data:
    IMP = g.impute(M = data.getmeta, I = data.inspectfile)
    #analyse and produce summary:
A = g.analyse(I = data.inspectfile, C = data.calibrate, M = data.getmeta, IMP, ID = "01wk0")
    #plot data
    g.plot(IMP, M = data.getmeta, I = data.inspectfile, durplot=4)
```

applyCosinorAnalyses Apply Cosinor Analyses to time series

## Description

Wrapper function around cosinorAnalyses that first prepares the time series before applying the cosinorAnlayses

## Usage

applyCosinorAnalyses(ts, qcheck, midnightsi, epochsizes)

## Arguments

ts Data.frame with timestamps and acceleration metric.
qcheck Vector of equal length as number of rows in ts with value 1 for invalid timestamps, 0 otherwise.
midnightsi Indices for midnights in the time series
epochsizes Epoch size for ts and qcheck respectively

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

```
create_test_acc_csv Creates csv data file for testing purposes
```


## Description

Creates file in the Actigraph csv data format with dummy data that can be used for testing. The file includes accelerometer data with bouts of higher acceleration, variations non-movement periods in a range of accelerometer positions to allow for testing the auto-calibration functionality.

## Usage

create_test_acc_csv(sf=3,Nmin=2000, storagelocation=c(),
start_time $=$ NULL, starts_at_midnight $=$ FALSE)

## Arguments

sf \begin{tabular}{l}
Sample frequency in Hertz, the default here is low to minimize file size <br>
Nmin <br>
storagelocation

 

Location where the test file named testfile.csv will be stored If no value is pro- <br>
vided then the function uses the current working directory
\end{tabular}

start_time $\quad$| Start time of the recording, in the hh:mm:ss format. |
| :--- |
| starts_at_midnight |

Boolean indicating whether the recording should start at midnight. Ignored if
start_time is specified.

## Value

The function does not produce any output values. Only the file is stored

## Examples

```
    ## Not run:
        create_test_acc_csv()
## End(Not run)
```

create_test_sleeplog_csv
Creates csv sleeplog file for testing purposes

## Description

Creates sleeplog file in the format as expected by g.part4 with dummy data (23:00 onset, 07:00 waking time for every night).

## Usage

```
create_test_sleeplog_csv(Nnights = 7, storagelocation = c(),
    advanced = FALSE, sep = ",")
```


## Arguments

Nnights $\quad$ Number of nights (minimum is 1 )
storagelocation
Location where the test file named testfile.csv will be stored If no value is provided then the function uses the current working directory
advanced Boolean to indicate whether to create an advanced sleeplog that also includes logs of nap times and nonwear
sep $\quad$ Character to indicate the column separator of the csv file.

## Value

The function does not produce any output values. Only the file is stored

## Examples

> \# Not run:

```
        create_test_sleeplog_csv()
```

    \#\# End(Not run)
    ```
data.calibrate Example output from g.calibrate
```


## Description

data.calibrate is example output from g.calibrate

## Usage

data(data.calibrate)

## Format

The format is: chr "data.calibrate"

## Source

The data was collected on one individual for testing purposes

## Examples

```
    data(data.calibrate)
```

    data.getmeta Example output from g.getmeta
    
## Description

data.getmeta is example output from g.getmeta

## Usage

data(data.getmeta)

## Format

The format is: chr "data.getmeta"

## Source

The data was collected on one individual for testing purposes

## Examples

```
data(data.getmeta)
```

data.inspectfile Example output from g.inspectfile

## Description

data.inspectfile is example output from g.inspectfile

## Usage

data(data.inspectfile)

## Format

The format is: chr "data.inspectfile"

## Source

The data was collected on one individual for testing purposes

## Examples

```
data(data.inspectfile)
```

data.metalong Metalong object as part of part 1 milestone data

## Description

data.metalong is example of the metalong data.frame stored g.part1

## Usage

data(data.metalong)

## Format

The format is: chr "data.metalong"

## Source

The data was collected on one individual for testing purposes

## Examples

data(data.metalong)
data.ts Time series data.frame stored by part 5

## Description

data.ts is example of the data.frame stored g.part5

## Usage

data(data.ts)

## Format

The format is: chr "data.ts"

## Source

The data was collected on one individual for testing purposes and matches the data in object data.metalong

## Examples

data(data.ts)

g.calibrate | function to estimate calibration error and make recommendation for |
| :--- |
| addressing it |

## Description

Function starts by identifying ten second windows of non-movement. Next, the average acceleration per axis per window is used to estimate calibration error (offset and scaling) per axis. The function provides recommended correction factors to address the calibration error and a summary of the callibration procedure.

## Usage

g.calibrate(datafile, params_rawdata $=c()$, params_general $=c()$,
params_cleaning $=c()$, inspectfileobject $=c()$, verbose $=$ TRUE, ...)

## Arguments

```
    datafile Name of accelerometer file
    params_rawdata See g.part1
    params_general See g.part1
    params_cleaning
        See g.part1
    inspectfileobject
```

                Output from the function g.inspectfile.
    verbose \(\quad\) Boolean \((\) default \(=T R U E)\). to indicate whether console message should be printed.
        Note that warnings and error are always printed and can be suppressed with sup-
        pressWarning() or suppressMessages().
    ... Any argument used in the previous version of g.calibrate, which will now be
    used to overrule the arguments specified with the parameter objects.
    
## Value

$\left.\begin{array}{ll}\text { scale } & \text { scaling correction values, e.g. c(1,1,1) } \\ \text { offset } & \text { offset correction values, e.g. c(0,0,0) } \\ \text { tempoffset } & \text { correction values related to temperature, e.g. c(0,0,0) } \\ \text { cal.error.start }\end{array} \quad \begin{array}{l}\text { absolute difference between Euclidean norm during all non-movement windows } \\ \text { and } 1 \mathrm{~g} \text { before autocalibration }\end{array}\right\}$

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com> Zhou Fang

## References

- van Hees VT, Fang Z, Langford J, Assah F, Mohammad A, da Silva IC, Trenell MI, White T, Wareham NJ, Brage S. Auto-calibration of accelerometer data for free-living physical activity assessment using local gravity and temperature: an evaluation on four continents. J Appl Physiol (1985). 2014 Aug 7


## Examples

        \#\# Not run:
        datafile = "C:/myfolder/testfile.bin"
        \#Apply autocalibration:
        C = g.calibrate(datafile)
        print (C\$scale)
        print (C\$offset)
    \#\# End(Not run)

```
g.getbout function to calculate bouts from vector of binary classes
```


## Description

To detect bouts of behaviour in time series. The function is used by g.analyse

## Usage

g.getbout(x, boutduration, boutcriter $=0.8$, ws3 $=5$ )

## Arguments

x
boutduration duration of bout in epochs
boutcriter Minimum percentage of boutduration for which the epoch values are expected to meet the threshold criterium
ws3 epoch length in seconds, only needed for bout.metric $=3$, because it needs to measure how many epochs equal 1 minute breaks

## Value

Vector with binary numbers indicator where bouts where detected

## Author(s)

Vincent T van Hees < v.vanhees @accelting.com> Jairo Hidalgo Migueles

## Examples

```
y = g.getbout(x=round(runif(1000, 0.4, 1)), boutduration = 120, boutcriter=0.9,
    ws3 = 5)
```

Function to extract meta-data (features) from data in accelerometer file

## Description

Reads a accelerometer file in blocks, extracts various features and stores average feature value per short or long epoch. Acceleration and angle metrics are stored at short epoch length. The non-wear indication score, the clipping score, temperature (if available), light (if available), and Euclidean norm are stored at long epoch length. The function has been designed and thoroughly tested with accelerometer files from GENEA and GENEActiv bin files. Further, the function should be able to cope with ActiGraph gt3x and csv files, Axivity cwa and csv files, Movisens bin files, and ad-hoc csv files read through the read.myacc.csv function.

## Usage

$$
\begin{aligned}
& \text { g.getmeta(datafile, params_metrics }=c(), \text { params_rawdata }=c(), \\
& \text { params_general }=c() \text {, params_cleaning }=c() \text {, daylimit }=\text { FALSE, } \\
& \text { offset }=c(0,0,0), \text { scale }=c(1,1,1) \text {, tempoffset }=c(0,0,0), \\
& \text { meantempcal }=c(), \text { myfun }=c() \text {, inspectfileobject }=c(), \\
& \text { verbose }=\text { TRUE, } \ldots)
\end{aligned}
$$

## Arguments



| verbose | Boolean $($ default $=$ TRUE $)$. to indicate whether console message should be printed <br> Note that warnings and error are always printed and can be suppressed with sup- <br> pressWarning () or suppressMessages(). |
| :--- | :--- |
| $\ldots$ | Any argument used in the previous version of g.getmeta, which will now be used <br> to overrule the arguments specified with the parameter objects. |

## Value

| metalong | dataframe with long epoch meta-data: EN, non-wear score, clipping score, tem- <br> perature |
| :--- | :--- |
| metashort | dataframe with short epoch meta-data: timestamp and metric |
| tooshort | indicator of whether file was too short for processing (TRUE or FALSE) |
| corrupt | indicator of whether file was considered corrupt (TRUE or FALSE) |

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## References

- van Hees VT, Gorzelniak L, Dean Leon EC, Eder M, Pias M, et al. (2013) Separating Movement and Gravity Components in an Acceleration Signal and Implications for the Assessment of Human Daily Physical Activity. PLoS ONE 8(4): e61691. doi:10.1371/journal.pone. 0061691
- Aittasalo M, Vaha-Ypya H, Vasankari T, Husu P, Jussila AM, and Sievanen H. Mean amplitude deviation calculated from raw acceleration data: a novel method for classifying the intensity of adolescents physical activity irrespective of accelerometer brand. BMC Sports Science, Medicine and Rehabilitation (2015).


## Examples

```
    ## Not run:
    datafile = "C:/myfolder/testfile.bin"
    #Extract meta-data:
    M = g.getmeta(datafile)
    #Inspect first couple of rows of long epoch length meta data:
    print(M$metalong[1:5,])
    #Inspect first couple of rows of short epoch length meta data:
    print(M$metalong[1:5,])
## End(Not run)
```


## Description

Removes all sample with a zero in each of the three axes, and then (as default) imputes time gaps by the last recorded value per axis normalised to $1{ }_{\text {_ }} \mathrm{g}_{-}$

## Usage

```
g.imputeTimegaps(x, sf, k = 0.25, impute = TRUE,
    PreviousLastValue = c(0,0,1),
    PreviousLastTime = NULL, epochsize = NULL)
```


## Arguments

x
sf Sample frequency in Hertz
k
impute Boolean to indicate whether the time gaps identified should be imputed
PreviousLastValue
Automatically identified last value in previous chunk of data read.
PreviousLastTime
Automatically identified last timestamp in previous chunk of data read.
epochsize Numeric vector of length two, with short and long epoch sizes.

## Value

List including: - x , data.frame based on input x with timegaps imputed (as default) or with recordings with 0 values in the three axes removed (if impute $=$ FALSE) - QClog, data.frame with information on the number of time gaps found and the total time imputed in minutes

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

g.inspectfile | function to inspect accelerometer file for brand, sample frequency and |
| :--- |
| header |

## Description

Inspects accelerometer file for key information, including: monitor brand, sample frequency and file header

## Usage

```
g.inspectfile(datafile, desiredtz = "", params_rawdata = c(),
```

                            configtz = c(), ...)
    
## Arguments

| datafile | name of data file |
| :--- | :--- |
| desiredtz | Desired timezone, see documentation g.getmeta |
| params_rawdata | See g.part1 |
| configtz | ... |

... Any argument used in the previous version of g.getmeta, which will now be used to overrule the arguments specified with the parameter objects.

## Value

| header | fileheader |
| :---: | :---: |
| monn | monitor name (genea, geneactive) |
| monc | monitor brand code ( $0-$ ad-hoc file format, $1=$ genea (non-commercial), $2=$ GENEActive, $3=$ actigraph, $4=$ Axivity (AX3, AX6), $5=$ Movisense, $6=$ Verisense) |
| dformn | data format name, e.g bin, csv, cwa, gt3x |
| dformc | data format code $(1=$.bin, $2=. \operatorname{csv}, 3=$. wav, $4=. c w a, 5=$ ad-hoc $. c s v, 6=$ .gt3x) |
| sf | samplefrequency in Hertz |
| filename | filename |

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## Description

Loads sleeplog from a csv input file and applies sanity checks before storing the output in a dataframe

## Usage

```
g.loadlog(loglocation=c(),coln1=c(),colid=c(),
    sleeplogsep=",", meta.sleep.folder = c(),
desiredtz="")
```


## Arguments

loglocation Location of the spreadsheet (csv) with sleep log information. See package vignette for explanation on expected format
coln1 Column number in the sleep log spreadsheet where the onset of the first night starts
colid Column number in the sleep $\log$ spreadsheet in which the participant ID code is stored $($ default $=1)$
sleeplogsep Value used as sep argument for reading sleeplog csv file, usually "," or ";". This argument has been deprecated.
meta.sleep.folder
Path to part3 milestone data, only specify if sleeplog is in advanced format.
desiredtz See g.part4

## Value

Data frame with sleeplog, which can be either in basic format or in advanced format. See GGIR package vignette for discussion of these two formats.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## Examples

```
## Not run:
    sleeplog = g.loadlog(loglocation="C:/mysleeplog.csv",coln1=2,
    colid=1)
## End(Not run)
```


## Description

Calls function g.getmeta and g.calibrate, and converts the output to .RData-format which will be the input for g.part2. Here, the function generates a folder structure to keep track of various output files. The reason why these g.part1 and g.part2 are not merged as one generic shell function is because g.part 1 takes much longer to and involves only minor decisions of interest to the movement scientist. Function g.part2 on the other hand is relatively fast and comes with all the decisions that directly impact on the variables that are of interest to the movement scientist. Therefore, the user may want to run g.part1 overnight or on a computing cluster, while g.part2 can then be the main playing ground for the movement scientist. Function GGIR provides the main shell that allows for operating g.part1 and g.part2.

## Usage

```
g.part1(datadir = c(), metadatadir = c(), f0 = 1, f1 = c(),
    myfun = c(), params_metrics = c(), params_rawdata = c(),
    params_cleaning = c(), params_general = c(), verbose = TRUE, ...)
```


## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory where the output needs to be stored. Note that this function will attempt to create folders in this directory and uses those folder to keep output.
f0 File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., f1 $=0$ )
myfun External function object to be applied to raw data. See details applyExtFunction.
params_metrics See details in GGIR.
params_rawdata See details in GGIR.
params_cleaning
See details in GGIR.
params_general See details in GGIR.
verbose See details in GGIR.
... If you are working with a non-standard csv formatted files, g.part1 also takes any input arguments needed for function read.myacc.csv and argument rmc.noise from get_nw_clip_block_params. First test these argument with function read.myacc.csv directly. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_ objects as direct argument.

## Details

GGIR comes with many processing parameters, which have been thematically grouped in parameter objects (R list). By running print(load_params()) you can see the default values of all the parameter objects. When g.part 1 is used via GGIR you have the option to specifiy a configuration file, which will overrule the default parameter values. Further, as user you can set parameter values as input argument to both g.partl and GGIR. Directly specified argument overrule the configuration file and default values.

See the GGIR package vignette or the details section in GGIR for a more elaborate overview of parameter objects and their usage across GGIR.

## Value

The function provides no values, it only ensures that the output from other functions is stored in .RData(one file per accelerometer file) in folder structure

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

## References

- van Hees VT, Gorzelniak L, Dean Leon EC, Eder M, Pias M, et al. (2013) Separating Movement and Gravity Components in an Acceleration Signal and Implications for the Assessment of Human Daily Physical Activity. PLoS ONE 8(4): e61691. doi:10.1371/journal.pone. 0061691
- van Hees VT, Fang Z, Langford J, Assah F, Mohammad A, da Silva IC, Trenell MI, White T, Wareham NJ, Brage S. Auto-calibration of accelerometer data for free-living physical activity assessment using local gravity and temperature: an evaluation on four continents. J Appl Physiol (1985). 2014 Aug 7
- Aittasalo M, Vaha-Ypya H, Vasankari T, Husu P, Jussila AM, and Sievanen H. Mean amplitude deviation calculated from raw acceleration data: a novel method for classifying the intensity of adolescents physical activity irrespective of accelerometer brand. BMC Sports Science, Medicine and Rehabilitation (2015).


## Examples

```
    ## Not run:
        datafile = "C:/myfolder/mydata"
        outputdir = "C:/myresults"
        g.part1(datadir,outputdir)
## End(Not run)
```


## Description

Loads the output from g.part 1 and then applies g.impute and g.analyse, after which the output is converted to .RData-format which will be used by GGIR to generate reports. The variables in these reports are the same variables as described in g.analyse.

## Usage

$$
\begin{aligned}
& \text { g.part2(datadir }=c(), \text { metadatadir }=c(), f 0=c(), f 1=c(), \\
& \text { myfun }=c(), \text { params_cleaning }=c() \text {, params_247 }=c(), \\
& \text { params_phyact }=c() \text {, params_output }=c() \text {, params_general }=c() \text {, } \\
& \text { verbose }=\text { TRUE, } \ldots \text { ) }
\end{aligned}
$$

## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part1 and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default =1). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., f1 = 0)
myfun External function object to be applied to raw data. See details applyExtFunction.
params_cleaning
See details in GGIR.
params_247 See details in GGIR.
params_phyact See details in GGIR.
params_output See details in GGIR.
params_general See details in GGIR.
verbose See details in GGIR.
.. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_ objects as direct argument.

## Details

GGIR comes with many processing parameters, which have been thematically grouped in parameter objects (R list). By running print(load_params()) you can see the default values of all the parameter objects. When g.part 2 is used via GGIR you have the option to specifiy a configuration file, which will overrule the default parameter values. Further, as user you can set parameter values as input
argument to both g.part2 and GGIR. Directly specified argument overrule the configuration file and default values.
See the GGIR package vignette or the details section in GGIR for a more elaborate overview of parameter objects and their usage across GGIR.

## Value

The function provides no values, it only ensures that other functions are called and that their output is stored in the folder structure as created with g.part1.

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

## References

- van Hees VT, Gorzelniak L, Dean Leon EC, Eder M, Pias M, et al. (2013) Separating Movement and Gravity Components in an Acceleration Signal and Implications for the Assessment of Human Daily Physical Activity. PLoS ONE 8(4): e61691. doi:10.1371/journal.pone. 0061691
- van Hees VT, Fang Z, Langford J, Assah F, Mohammad A, da Silva IC, Trenell MI, White T, Wareham NJ, Brage S. Auto-calibration of accelerometer data for free-living physical activity assessment using local gravity and temperature: an evaluation on four continents. J Appl Physiol (1985). 2014 Aug 7


## Examples

```
    ## Not run:
    metadatadir = "C:/myresults/output_mystudy"
    g.part2(metadatadir)
## End(Not run)
```

| g.part 3 | Detection of sustained inactivity periods as needed for sleep detection <br> in g.part4. |
| :--- | :--- |

## Description

Function called by function GGIR. It estimates the sustained inactivity periods in each day, which are used as input for g.part4 which then labels them as nocturnal sleep or day time sustained inactivity periods. Typical users should work with function GGIR only.

## Usage

g.part3(metadatadir = c(), f0, f1, myfun = c(), params_sleep = c(), params_metrics = c(), params_output = c(), params_general = c(), verbose = TRUE, ...)

## Arguments

> metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.partl and GGIR will recognise what the value of metadatadir is.
> f0 File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
> f1 File index to finish with (defaults to number of files available, i.e., f1 $=0$ )
> myfun External function object to be applied to raw data. See details applyExtFunction.
> params_sleep See details in GGIR.
> params_metrics See details in GGIR.
> params_output See details in GGIR.
> params_general See details in GGIR.
> verbose See details in GGIR.
> ... To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_ objects as direct argument.

## Details

GGIR comes with many processing parameters, which have been thematically grouped in parameter objects (R list). By running print(load_params()) you can see the default values of all the parameter objects. When g.part 3 is used via GGIR you have the option to specifiy a configuration file, which will overrule the default parameter values. Further, as user you can set parameter values as input argument to both g.part3 and GGIR. Directly specified argument overrule the configuration file and default values.
See the GGIR package vignette or the details section in GGIR for a more elaborate overview of parameter objects and their usage across GGIR.

## Value

The function provides no values, it only ensures that other functions are called and that their output is stored in .RData files.

- night nightnumber
- definition definition of sustained inactivity. For example, T10A5 refers to 10 minute window and a 5 degree angle (see paper for further explaination).
- start. time. day timestamp when the day started
- nsib.periods number of sustained inactivity bouts
- tot.sib. dur.hrs total duration of all sustained inactivity bouts
- fraction.night. invalid fraction of the night for which accelerometer data was invalid, e.g. monitor not worn
- sib. period number of sustained inactivity period
- sib. onset. time onset time of sustained inactivity period
- sib.end.time end time of sustained inactivity period


## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

## References

- van Hees VT, Sabia S, et al. (2015) A novel, open access method to assess sleep duration using a wrist-worn accelerometer, PLoS ONE, November 2015
- van Hees VT, Sabia S, et al. (2018) Estimating sleep parameters using an accelerometer without sleep diary. Scientific Reports.


## Examples

```
    ## Not run:
    metadatadir = "C:/myfolder/meta" # assumes that there is a subfolder in
    # metadatadir named 'basic' containing the output from g.part1
    g.part3(metadatadir=metadatadir, anglethreshold=5,
    timethreshold=5, overwrite=FALSE)
## End(Not run)
```

g. part4

Labels detected sustained inactivity periods by g.part 3 as either part of the Sleep Period Time window or not

## Description

Combines output from g.part3 and guider information to estimate sleep variables. See vignette paragraph "Sleep and full day time-use analysis in GGIR" for an elaborate descript of the sleep detection.

## Usage

g.part4(datadir $=c()$, metadatadir $=c(), f 0=f 0, f 1=f 1$, params_sleep $=c()$, params_metrics = c(), params_cleaning = c(), params_output = c(), params_general = c(), verbose = TRUE, ...)

## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part1 and GGIR will recognise what the value of metadatadir is.
f0
File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., $\mathrm{f} 1=0$ )
params_sleep List of parameters used for sleep analysis (GGIR part 3, 4, and 5): see documentation g.part3.
params_metrics List of parameters used for metrics extraction (GGIR part 1): see documentation g.part1.
params_cleaning
See details in GGIR.
params_output See details in GGIR.
params_general See details in GGIR.
verbose See details in GGIR.
.. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_objects as direct argument.

## Value

The function does not produce values but generates an RData file in the milestone subfolder ms4.out which incudes a dataframe named nightsummary. This dataframe is used in g.report.part4 to create two reports one per night and one per person. See package vignette paragraph "Output part 4 " for description of all the variables.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## References

- van Hees VT, Sabia S, et al. (2018) AEstimating sleep parameters using an accelerometer without sleep diary, Scientific Reports.
- van Hees VT, Sabia S, et al. (2015) A novel, open access method to assess sleep duration using a wrist-worn accelerometer, PLoS ONE.


## Examples

```
    ## Not run:
        metadatadir = "C:/myfolder/meta" # assumes that there is a subfolder in
        # metadatadir named 'ms3.out' containing the output from g.part3
        g.part4(metadatadir=metadatadir)
## End(Not run)
```


## Description

Function to merge the output from g.part2 and g.part4 into one report enhanced with profiling of sleep and physical activity stratified across intensity levels and based on bouted periods as well as non-bouted periods.

## Usage

$$
\begin{aligned}
\text { g. part5(datadir }= & \mathrm{c}(), \text { metadatadir }=\mathrm{c}(), f 0=\mathrm{c}(), \mathrm{f} 1=\mathrm{c}(), \\
& \text { params_sleep }=\mathrm{c}(), \text { params_metrics }=\mathrm{c}(), \\
& \text { params_247 }=\mathrm{c}(), \text { params_phyact }=\mathrm{c}(), \\
& \text { params_cleaning }=\mathrm{c}(), \text { params_output }=\mathrm{c}(), \\
& \text { params_general }=\mathrm{c}(), \text { verbose }=\text { TRUE, } \ldots)
\end{aligned}
$$

## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part1 and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., f1 $=0$ )
params_sleep See details in GGIR.
params_metrics See details in GGIR.
params_247 See details in GGIR.
params_phyact See details in GGIR.
params_cleaning
See details in GGIR.
params_output See details in GGIR.
params_general See details in GGIR.
verbose See details in GGIR.
.. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_ objects as direct argument.

## Value

The function does not produce values but generates an RData file in the milestone subfolder ms5.out which incudes a dataframe named output. This dataframe is used in g.report.part5 to create two reports one per day and one per person. See package vignette paragraph "Output part 5 " for description of all the variables.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## Examples

```
    ## Not run:
        metadatadir = "C:/myfolder/meta"
        g.part5(metadatadir=metadatadir)
## End(Not run)
```

g.part5.analyseRest Analyse rest (internal function)

## Description

Analyses overlap self-reported napping, non-wear and sib. Internal function not intended for direct use by GGIR end-user.

## Usage

$$
\begin{aligned}
\text { g. part5.analyseRest(sibreport } & =\text { NULL, dsummary }=\text { NULL, } \\
& \text { ds_names }=\text { NULL, fi }=\text { NULL, } \\
& \text { di }=\text { NULL, time }=\text { NULL, tz }=\text { NULL, } \\
& \text { possible_nap_dur }=0, \\
& \text { possible_nap_edge_acc }=\text { Inf) }
\end{aligned}
$$

## Arguments

sibreport sibreport data.frame produced by g.sibreport
dsummary matrix created internally by g.part5
ds_names character vector with variable names corresponding to dsummary created internally by g.part5
fi Numeric scalar to indicate variable index, created internally by g.part5
di Numeric scalar to indicate recording index, created internally by g.part5
time Daytime section of time column from the ts object, created internally by g.part5,
tz Timezone database name
possible_nap_dur
Minimum sib duration to be considered. For self-reported naps/nonwear there is a minimum duration of 1 epoch.
possible_nap_edge_acc
Maximum acceleration before or after the SIB for it to be considered.

## Value

List with updated objects dsummary, ds_names, fi, and di

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>
g.part6 Perform temporal pattern analyses

## Description

This function aims to facilitate time-pattern analysis building on the labelled time series derived in GGIR part 5

## Usage

$$
\begin{aligned}
\text { g. part6(datadir }= & c(), \text { metadatadir }=c(), f 0=c(), f 1=c(), \\
& \text { params_general }=c(), \text { params_phyact }=c(), \text { params_247 }=c(), \\
& \text { verbose }=\text { TRUE }, \ldots)
\end{aligned}
$$

## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.partl and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default =1). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., $\mathrm{f} 1=0$ )
params_general See details in GGIR.
params_phyact See details in GGIR.
params_247 See details in GGIR.
verbose See details in GGIR.
.. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the params_objects as direct argument.

## Value

The function does not produce values but generates an RData file in the milestone subfolder ms6.out which incudes ... (TO BE COMPLETED). This dataframe is used in g.report.part6 to create reports. See package vignette paragraph (TO BE COMPLETED) for description of all the variables.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## Examples

```
    ## Not run:
        metadatadir = "C:/myfolder/meta"
        g.part6(metadatadir=metadatadir)
## End(Not run)
```

g.plot5

Generate user-friendly visual report. The first part of the report summarizes important daily metrics in bar plot format. The second part of the report shows the raw data and annotations in 24 -hr periods. Angle-z is shown with sleep annotations during the SPT (sleep period time) window. ENMO is shown with daytime inactivity and PA (physical activity) annotations in the lower section of each 24-hr plot. The PA annotations are based on a 10 minute bout metric and 80 of a 10 minute bout of MVPA. Vigorous PA is a short window of time above threshold.vig that is part of a bout of MVPA. Light PA is a short window of time above threshold.lig that is part of a bout of light PA.

## Description

Function called by GGIR to generate report. Not intended for direct use by user

## Usage

```
g.plot5(metadatadir = c(), dofirstpage = FALSE, viewingwindow = 1,
f0 = c(), f1 = c(), overwrite = FALSE, metric="ENMO",desiredtz = "",
threshold.lig = 30, threshold.mod = 100, threshold.vig = 400,
visualreport_without_invalid = TRUE, includedaycrit = 0.66, includenightcrit = 0.66,
    verbose = TRUE)
```


## Arguments

metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part1 and GGIR will recognise what the value of metadatadir is.
dofirstpage Boolean to indicate whether a first page with historgrams summarizing the whole measurement should be added
viewingwindow
See GGIR
f0 File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., f1 $=0$ )
overwrite See GGIR
metric Which one of the metrics do you want to consider to describe behaviour. The metric of interest need to be calculated in M (see g.part1)

| desiredtz | See GGIR |  |
| :--- | :--- | :---: |
| threshold.lig | See GGIR |  |
| threshold.mod | See GGIR |  |
| threshold.vig | See GGIR |  |
| visualreport_without_invalid |  |  |
| See GGIR |  |  |
| includenightcrit |  |  |
| See GGIR |  |  |
| includedaycrit | See GGIR |  |
| verbose | See GGIR |  |

## Value

No values, this function only generates a plot

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com> Matthew R Patterson <mpatterson@ shimmersensing.com>

## Examples

```
## Not run:
        # generate plots for the first 10 files:
        g.plot5(metadatadir="C:/output_mystudy/meta/basic",dofirstpage=TRUE,
        viewingwindow = 1,f0=1,f1=10,overwrite=FALSE,desiredtz = "Europe/London",
        threshold.lig,threshold.mod,threshold.vig)
## End(Not run)
```

    g. report. part2
    Generate report from milestone data produced by g.part 2
    
## Description

Creates report from milestone data produced by g.part2. Not intended for direct use by package user

## Usage

g. report. part2(metadatadir $=c(), f 0=c(), f 1=c()$, maxdur $=0$, store.long = FALSE, params_output, verbose = TRUE)

## Arguments

metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.partl and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., $\mathrm{f} 1=0$ )
maxdur see g.part2
store.long Booelean to indicate whether output should stored in long format in addition to default wide format. Automatically turned to TRUE if using day segmentation with qwindow.
params_output Parameters object, see GGIR
verbose See details in GGIR.

## Value

Function does not produce data, but only writes reports in csv format and visual reports in pdf format

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

```
g.report.part4 Generate report from milestone data produced by g.part4
```


## Description

Creates report from milestone data produced by g.part4. Not intended for direct use by package user

## Usage

```
g.report.part4(datadir = c(), metadatadir = c(), loglocation = c(), f0 = c(),
    f1 = c(), data_cleaning_file = c(),
    sleepwindowType = "SPT", params_output, verbose = TRUE)
```


## Arguments

datadir Directory where the accelerometer files are stored, e.g. "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part1 and GGIR will recognise what the value of metadatadir is.
loglocation see g.part4

```
f0
File index to start with (default \(=1\) ). Index refers to the filenames sorted in alphabetical order File index to finish with (defaults to number of files available, i.e., \(\mathrm{f} 1=0\) )
data_cleaning_file
    see GGIR
sleepwindowType
    see GGIR
params_output Parameters object, see GGIR
verbose See details in GGIR.
```


## Value

Function does not produce data, but only writes reports in csv format and a visual report in pdf.
The following files are stored in the root of the results folder: part4_nightsummary_sleep_cleaned.csv part4_summary_sleep_cleaned.csv
The following files are stored in the folder results/QC: part4_nightsummary_sleep_full.csv part4_summary_sleep_full.csv
If a sleeplog is used *_full.csv as stored in the QC folder includes estimates for all nights in the data, and *_cleaned.csv in the results folder includes estimates for all nights in the data excluding the nights that did not had a sleeplog entry or had no valid accelerometer data.

If a sleep log is not used then * _cleaned.csv includes the nights that are in *_full.csv excluding the nights with insufficient data.

If you have a study where the sleeplog was available for a subset of the participants, but you want to include all individuals in your analysis, then use the *_full.csv output and clean the night level data yourself by excluding rows with cleaningcode $>1$ which are the cases where no or invalid accelerometer data was present.

The above means that for studies with missing sleeplog entries for some individuals and some nights using the *_full.csv output and excluding rows (nights) with cleaningcode $>1$ will lead to the same as * _cleaned.csv plus sleep estimates for the nights with missing sleeplog, providing that there was enough accelerometer data for those nights.

In other words, *_cleaned.csv is perfect if you only want to rely on nights with a sleeplog or if you do not use a sleeplog at all. For all other scenarios We advise using the *_full.csv report and to clean it yourself.

See package vignette sections "Sleep analysis" and "Output part 4" for a more elaborative description of the sleep analysis and reporting.

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

## Description

Creates report from milestone data produced by g.part5. Not intended for direct use by package user

## Usage

```
g.report.part5(metadatadir = c(), f0 = c(), f1 = c(), loglocation = c(),
    params_cleaning = NULL,
    LUX_day_segments = c(), params_output,
    verbose = TRUE)
```


## Arguments

metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.partl and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default =1). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., $\mathrm{f} 1=0$ )
loglocation see g.part4
params_cleaning
See details in GGIR.
LUX_day_segments
see g.part5
params_output Parameters object, see GGIR
verbose See details in GGIR.

## Value

Function does not produce data, but only writes reports in csv format
The following files are stored in the root of the results folder: part5_daysummary_* part5_personsummary_*
The following files are stored in the folder results/QC: part5_daysummary_full_*
See package vignette paragraph "Waking-waking or 24 hour time-use analysis" and "Output part 5" for a more elaborative description of the full day time-use and analysis and reporting.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>
g.report.part5_dictionary

Generate data dictionary for reports from milestone data produced by g.part5

## Description

Creates a data dictionary with the definitions of the outcomes exported in the reports from milestone data produced by g.part5. Not intended for direct use by package user.

## Usage

g.report.part5_dictionary(metadatadir, params_output)

## Arguments

metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.partl and GGIR will recognise what the value of metadatadir is.
params_output Parameters object, see GGIR

## Value

Function does not produce data, but only writes data dictionaries for the reports in csv format
The following files are stored in the root of the results folder: part5_dictionary_daysummary_* part5_dictionary_personsummary_*

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com) Jairo Hidalgo Migueles [jairo@jhmigueles.com](mailto:jairo@jhmigueles.com)

```
g.report.part6
```

Generate report from milestone data produced by g.part6

## Description

Creates report from milestone data produced by g.part6. Not intended for direct use by package user

## Usage

```
g.report.part6(metadatadir = c(), f0 = c(), f1 = c(),
    params_cleaning = NULL, params_output,
    verbose = TRUE)
```


## Arguments

metadatadir Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by g.part1. The folderstructure is normally created by g.part 1 and GGIR will recognise what the value of metadatadir is.
f0 File index to start with (default =1). Index refers to the filenames sorted in alphabetical order
f1 File index to finish with (defaults to number of files available, i.e., $\mathrm{f} 1=0$ )
params_cleaning
See details in GGIR.
params_output Parameters object, see GGIR
verbose See details in GGIR.

## Value

Function does not produce data, but only writes reports in csv format
The following files are stored in the root of the results folder: part6_summary.csv
See package vignette "HouseHoldCoanalysis".

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>
g.shell.GGIR Wrapper function around function GGIR

## Description

This function used to be the central function in the package, but has been renamed GGIR. You can still use function call g.shell.GGIR but all arguments will be passed on to function GGIR. We have done this to preserve consistency with older use cases of the GGIR package. All documentation can now be found in GGIR.

## Usage

g.shell.GGIR(...)

## Arguments

... Any of the parameters used by GGIR.

## Value

The function provides no values, it only ensures that other functions are called and that their output is stored. See GGIR.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## GGIR Shell function for analysing an accelerometer dataset.

## Description

This function is designed to help users operate all steps of the analysis. It helps to generate and structure milestone data, and produces user-friendly reports. The function acts as a shell with calls to g.part1, g.part2, g.part3, g.part4 and g.part5.

## Usage

```
GGIR(mode = 1:5,
    datadir = c(),
    outputdir = c(),
    studyname = c(),
    f0 = 1, f1 = 0,
    do.report = c(2, 4, 5, 6),
    configfile = c(),
    myfun = c(),
    verbose = TRUE, ...)
```


## Arguments

mode $\quad$ Numeric (default $=1: 5$ ). Specify which of the five parts need to be run, e.g., mode $=1$ makes that g.part 1 is run; or mode $=1: 5$ makes that the whole GGIR pipeline is run, from g.part1 to g.part5. Optionally mode can also include the number 6 to tell GGIR to run g.part6 which is currently under development.
datadir Character $($ default $=c())$. Directory where the accelerometer files are stored, e.g., "C:/mydata", or list of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin", "C:/mydata/myfile2.bin").
outputdir $\quad$ Character $($ default $=c())$. Directory where the output needs to be stored. Note that this function will attempt to create folders in this directory and uses those folder to keep output.
studyname $\quad$ Character $($ default $=c())$. If the datadir is a folder, then the study will be given the name of the data directory. If datadir is a list of filenames then the studyname as specified by this input argument will be used as name for the study.
Numeric (default $=1$ ). File index to start with (default $=1$ ). Index refers to the filenames sorted in alphabetical order.
f1 Numeric (default $=0$ ). File index to finish with (defaults to number of files available).

| do.report | Numeric (default $=c(2,4,5,6))$. For which parts to generate a summary spreadsheet: $2,4,5$, and/or 6 . Default is $c(2,4,5,6)$. A report will be generated based on the available milestone data. When creating milestone data with multiple machines it is advisable to turn the report generation off when generating the milestone data, value $=c()$, and then to merge the milestone data and turn report generation back on while setting overwrite to FALSE. |
| :---: | :---: |
| configfile | Character $($ default $=c())$. Configuration file previously generated by function GGIR. See details. |
| myfun | List $($ default $=c())$. External function object to be applied to raw data. See package vignette for detailed tutorial with examples on how to use the function embedding: https://cran.r-project.org/package=GGIR/vignettes/ExternalFunction.html |
| verbose | Boolean (default = TRUE). to indicate whether console message should be printed. Note that warnings and error are always printed and can be suppressed with suppressWarning() or suppressMessages(). |
|  | Any of the parameters used GGIR. Given the large number of parameters used in GGIR we have grouped them in objects that start with "params_". These are documented in the details section. You cannot provide these objects as argument to function GGIR, but you can provide the parameters inside them as input to function GGIR. |

## Details

Once you have used function GGIR and the output directory (outputdir) will be filled with milestone data and results. Function GGIR stores all the explicitely entered argument values and default values for the argument that are not explicitely provided in a csv-file named config.csv stored in the root of the output folder. The config.csv file is accepted as input to GGIR with argument configfile to replace the specification of all the arguments, except datadir and outputdir.

The practical value of this is that it eases the replication of analysis, because instead of having to share you R script, sharing your config.csv file will be sufficient. Further, the config.csv file contribute to the reproducibility of your data analysis.
Note: When combining a configuration file with explicitely provided argument values, the explicitely provided argument values will overrule the argument values in the configuration file. If a parameter is neither provided via the configuration file nor as input then GGIR uses its default paramter values which can be inspected with command print(load_params()), and if you are specifically interested in a certain subgroup of parameters, e.g., physical activity, then you can do print(load_params()\$params_phyact). These defaults are part of the GGIR code and cannot be changed by the user.
The parameters that can be used in GGIR are:
params_general: A list of parameters used across all GGIR parts that do not fall in any of the other categories.
overwrite Boolean (default = FALSE). Do you want to overwrite analysis for which milestone data exists? If overwrite $=$ FALSE, then milestone data from a previous analysis will be used if available and visual reports will not be created again.
dayborder Numeric $($ default $=0)$. Hour at which days start and end (dayborder $=4$ would mean 4 am ).
do.parallel Boolean (default $=$ TRUE). Whether to use multi-core processing (only works if at least 4 CPU cores are available).
maxNcores Numeric (default = NULL). Maximum number of cores to use when argument do.parallel is set to true. GGIR by default uses either the maximum number of available cores or the number of files to process (whichever is lower), but this argument allows you to set a lower maximum.
acc.metric Character (default = "ENMO"). Which one of the acceleration metrics do you want to use for all acceleration magnitude analyses in GGIR part 5 and the visual report? For example: "ENMO", "LFENMO", "MAD", "NeishabouriCount_y", or "NeishabouriCount_vm". Only one acceleration metric can be specified and the selected metric needs to have been calculated in part 1 (see g.part1) via arguments such as do. enmo $=$ TRUE or do. mad $=$ TRUE.
part5_agg2_60seconds Boolean (default = FALSE). Whether to use aggregate epochs to 60 seconds as part of the GGIR g.part5 analysis. Aggregation is doen by averaging. Note that when working with count metrics such as Neishabouri counts this means that the threshold can stay the same as in part 2, because again the threshold is expressed relative to the original epoch size, even if averaged per minute. For example if we want to use a cut-point 100 count per minute then we specify mvpathreshold $=100 *(5 / 60)$ as well as 'threshold.mod $=100 *$ (5/60) regardless of whether we set part5_agg2_60seconds to TRUE or FALSE.
print.filename Boolean (default = FALSE). Whether to print the filename before analysing it (in case do.parallel = FALSE). Printing the filename can be useful to investigate problems (e.g., to verify that which file is being read).
desiredtz Character (default $=$ " ", i.e., system timezone). Timezone in which device was configured and experiments took place. If experiments took place in a different timezone, then use this argument for the timezone in which the experiments took place and argument configtz to specify where the device was configured. Use the "TZ identifier" as specified at https://en.wikipedia.org/wiki/Zone.tab to set desiredtz, e.g., "Europe/London".
configtz Character (default $="$ ", i.e., system timezone). At the moment only functional for GENEActiv .bin, AX3 cwa, ActiGraph .gt3x, and ad-hoc csv file format. Timezone in which the accelerometer was configured. Only use this argument if the timezone of configuration and timezone in which recording took place are different. Use the "TZ identifier" as specified at https://en.wikipedia.org/wiki/Zone.tab to set configtz, e.g., "Europe/London".
sensor.location Character (default = "wrist"). To indicate sensor location, default is wrist. If it is hip, the HDCZA algorithm for sleep detection also requires longitudinal axis of sensor to be between -45 and +45 degrees.
windowsizes Numeric vector, three values (default $=c(5,900,3600)$ ). To indicate the lengths of the windows as in c (window1, window2, window3): window1 is the short epoch length in seconds, by default 5 , and this is the time window over which acceleration and angle metrics are calculated; window2 is the long epoch length in seconds for which non-wear and signal clipping are defined, default 900 (expected to be a multitude of 60 seconds); window3 is the window length of data used for non-wear detection and by default 3600 seconds. So, when window3 is larger than window2 we use overlapping windows, while if window2 equals window3 non-wear periods are assessed by non-overlapping windows.
idloc Numeric (default $=1$ ). If idloc $=1$ the code assumes that ID number is stored in the obvious header field. Note that for ActiGraph data the ID is never stored in the file header. For value set to $2,5,6$, and 7 , GGIR looks at the filename and extracts the character string preceding the first occurance of a "_" (idloc = 2), " " (space, idloc = 5), "." (dot, idloc = 6), and "-" (idloc $=7$ ), respectively. You may have noticed that idloc 3 and 4 are skipped, they were used
for one study in 2012, and not actively maintained anymore, but because it is legacy code not omitted.
expand_tail_max_hours Numeric (default = NULL). This parameter has been replaced by recordingEndSleepHour.
recordingEndSleepHour Numeric (default $=$ NULL). Time (in hours) at which the recording should end (or later) to expand the g.partl output with synthetic data to trigger sleep detection for last night. Using argument recordingEndSleepHour implies the assumption that the participant fell asleep at or before the end of the recording if the recording ended at or after recordingEndSleepHour hour of the last day. This assumption may not always hold true and should be used with caution. The synthetic data for metashort entails: timestamps continuing regularly, zeros for acceleration metrics other than EN, one for EN. Angle columns are created in a way that it triggers the sleep detection using the equation: round $(\sin ((1:$ length_expansion) $/(900 /$ epochsize $))) * 15$. To keep track of the tail expansion g.part1 stores the length of the expansion in the RData files, which is then passed via g.part2, g.part3, and g.part4 to g.part5. In g.part5 the tail expansion size is included as an additional variable in the csv-reports. In the g.part4 csv-report the last night is omitted, because we know that sleep estimates from the last night will not be trustworthy. Similarly, in the g.part5 output columns related to the sleep assessment will be omitted for the last window to avoid biasing the averages. Further, the synthetic data are also ignored in the visualizations and time series output to avoid biased output.
dataFormat Character (default = "raw"). To indicate what the format is of the data in datadir. Alternatives: ukbiobank_csv, actiwatch_csv, actiwatch_awd, actigraph_csv, and sensewear_xls, which correspond to epoch level data files from, respecitively, UK Biobank in csv format, Actiwatch in csv format, Actiwatch in awd format, ActiGraph csv format, and Sensewear in xls format (also works with xlsx). Here, the assumed epoch size for UK Biobank csvdata is 5 seconds. The epoch size for the other non-raw data formats is flexible, but make sure that you set first value of argument windowsizes accordingly. Also when working with non-raw data formats specify argument extEpochData_timeformat as documented below. For ukbiobank_csv nonwear is a column in the data itself, for actiwatch_csv, actiwatch_awd, actigraph_csv, and sensewear_xls non-wear is detected as 60 minute rolling zeros. The length of this window can be modified with the third value of argument windowsizes expressed in seconds.
maxRecordingInterval Numeric $($ default $=$ NULL). To indicate the maximum gap in hours between repeated measurements with the same ID for the recordings to be appended. So, the assumption is that the ID can be matched, make sure argument idloc is set correctly. If argument maxRecordingInterval is set to NULL (default) recordings are not appended. If recordings overlap then GGIR will use the data from the latest recording. If recordings are separated then the timegap between the recordings is filled with data points that resemble monitor not worn. The maximum value of maxFile gap is 504 ( 21 days). Only recordings from the same accelerometer brand are appended. The part 2 csv report will show number of appended recordings, sampling rate for each, time overlap or gap and the names of the filenames of the respective recording.
extEpochData_timeformat Character (default $=$ "\%d-\%m-\%Y $\% \mathrm{H}: \% \mathrm{M}: \% \mathrm{~S} ")$. To specify the time format used in the external epoch level data when argument dataFormat is set to "actiwatch_csv", "actiwatch_awd", "actigraph_csv" or "sensewear_xls". For example "\%Y-\%m\%d \%I:\%M:\%S \%p" for "2023-07-11 01:24:01 PM" or "\%m/\%d/\%Y \%H:\%M:\%S" "2023-07-11 13:24:01"
params_rawdata: A list of parameters used to related to reading and pre-processing raw data,
excluding parameters related to metrics as those are in the params_metrics object.
backup.cal.coef Character (default $=$ "retrieve"). Option to use backed-up calibration coefficient instead of deriving the calibration coefficients when analysing the same file twice. Argument backup.cal.coef has two usecase. Use case 1: If the auto-calibration fails then the user has the option to provide back-up calibration coefficients via this argument. The value of the argument needs to be the name and directory of a csv-spreadsheet with the following column names and subsequent values: "filename" with the names of accelerometer files on which the calibration coefficients need to be applied in case auto-calibration fails; "scale.x", "scale.y", and "scale.z" with the scaling coefficients; "offset.x", "offset.y", and "offset.z" with the offset coefficients, and; "temperature.offset.x", "temperature.offset.y", and "temperature.offset.z" with the temperature offset coefficients. This can be useful for analysing short lasting laboratory experiments with insufficient sphere data to perform the auto-calibration, but for which calibration coefficients can be derived in an alternative way. It is the users responsibility to compile the csv-spreadsheet. Instead of building this file the user can also Use case 2: The user wants to avoid performing the auto-calibration repeatedly on the same file. If backup.cal.coef value is set to "retrieve" (default) then GGIR will look out for the "data_quality_report.csv" file in the outputfolder QC, which holds the previously generated calibration coefficients. If you do not want this happen, then deleted the data_quality_report.csv from the QC folder or set it to value "redo".
minimumFileSizeMB Numeric (default $=2$ ). Minimum File size in MB required to enter processing. This argument can help to avoid having short uninformative files to enter the analyses. Given that a typical accelerometer collects several MBs per hour, the default setting should only skip the very tiny files.
do.cal Boolean (default = TRUE). Whether to apply auto-calibration or not by g.calibrate. Recommended setting is TRUE.
imputeTimegaps Boolean (default = TRUE). To indicate whether timegaps larger than 1 sample should be imputed. Currently only used for .gt3x data and ActiGraph .csv format, where timegaps can be expected as a result of Actigraph's idle sleep.mode configuration.
spherecrit Numeric (default $=0.3$ ). The minimum required acceleration value (in g ) on both sides of 0 g for each axis. Used to judge whether the sphere is sufficiently populated
minloadcrit Numeric (default $=168$ ). The minimum number of hours the code needs to read for the autocalibration procedure to be effective (only sensitive to multitudes of 12 hrs , other values will be ceiled). After loading these hours only extra data is loaded if calibration error has not been reduced to under 0.01 g .
printsummary Boolean (default $=$ FALSE). If TRUE will print a summary of the calibration procedure in the console when done.
chunksize Numeric (default $=1$ ). Value to specify the size of chunks to be loaded as a fraction of an approximately 12 hour period for auto-calibration procedure and as fraction of 24 hour period for the metric calculation, e.g., 0.5 equals 6 and 12 hour chunks, respectively. For machines with less than 4 Gb of RAM memory or with $<2 \mathrm{~GB}$ memory per process when using do.parallel = TRUE a value below 1 is recommended. The value is constrained by GGIR to not be lower than 0.05 . Please note that setting 0.05 will not produce output when 3rd value of parameter windowsizes is 3600 .
dynrange Numeric (default = NULL). Provide dynamic range of 8 gravity.
interpolationType Integer (default $=1$ ). To indicate type of interpolation to be used when resampling time series (mainly relevant for Axivity sensors), $1=$ linear, $2=$ nearest neighbour.
rmc.file Character (default $=$ NULL). Filename of file to be read if it is in the working directory, or full path to the file otherwise.
rmc.nrow Numeric (default $=$ NULL). Number of rows to read, same as nrow argument in read.csv and nrows in fread. The whole file is read by default (i.e., rmc.nrow = Inf).
rmc.skip Numeric (default $=0$ ). Number of rows to skip, same as skip argument in read.csv and in fread.
rmc.dec Character (default $=$ "."). Decimal used for numbers, same as dec argument in read.csv and in fread.
rmc.firstrow.acc Numeric (default $=$ NULL). First row (number) of the acceleration data.
rmc.firstrow.header Numeric (default = NULL). First row (number) of the header. Leave blank if the file does not have a header.
rmc.header.length Numeric $($ default $=$ NULL $)$. If file has header, specify header length (number of rows).
rmc.col.acc Numeric, three values (default $=c(1,2,3)$ ). Vector with three column (numbers) in which the acceleration signals are stored.
rmc.col.temp Numeric (default = NULL). Scalar with column (number) in which the temperature is stored. Leave in default setting if no temperature is available. The temperature will be used by g.calibrate.
rmc.col.time Numeric (default = NULL). Scalar with column (number) in which the timestamps are stored. Leave in default setting if timestamps are not stored.
rmc.unit.acc Character (default = "g"). Character with unit of acceleration values: "g", "mg", or "bit".
rmc.unit.temp Character (default = "C"). Character with unit of temperature values: (K)elvin, (C)elsius, or (F)ahrenheit.
rmc.unit.time Character (default = "POSIX"). Character with unit of timestamps: "POSIX", "UNIXsec" (seconds since origin, see argument rmc.origin), "character", or "ActivPAL" (exotic timestamp format only used in the ActivPAL activity monitor).
rmc.format.time Character (default $=$ " Character giving a date-time format as used by strptime. Only used for rmc.unit.time: character and POSIX.
rmc.bitrate Numeric (default $=$ NULL). If unit of acceleration is a bit then provide bit rate, e.g., 12 bit.
rmc.dynamic_range Numeric or character (default $=$ NULL). If unit of acceleration is a bit then provide dynamic range deviation in g from zero, e.g., $+/-6 \mathrm{~g}$ would mean this argument needs to be 6 . If you give this argument a character value the code will search the file header for elements with a name equal to the character value and use the corresponding numeric value next to it as dynamic range.
rmc.unsignedbit Boolean (default $=$ TRUE). If unsignedbit $=$ TRUE means that bits are only positive numbers. if unsignedbit $=$ FALSE then bits are both positive and negative.
rmc.origin Character (default $=$ "1970-01-01"). Origin of time when unit of time is UNIXsec, e.g., 1970-1-1.
rmc.desiredtz Character (default $=$ NULL). Timezone in which experiments took place. This argument is scheduled to be deprecated and is now used to overwrite desiredtz if not provided.
rmc.configtz Character (default = NULL). Timezone in which device was configured. This argument is scheduled to be deprecated and is now used to overwrite configtz if not provided.
rmc.sf Numeric (default $=$ NULL). Sample rate in Hertz, if this is stored in the file header then that will be used instead (see argument rmc.headername.sf).
rmc.headername.sf Character (default $=$ NULL). If file has a header: Row name under which the sample frequency can be found.
rmc.headername.sn Character (default = NULL). If file has a header: Row name under which the serial number can be found.
rmc.headername.recordingid Character (default $=$ NULL). If file has a header: Row name under which the recording ID can be found.
rmc.header.structure Character (default $=$ NULL). Used to split the header name from the header value, e.g., ":" or " ".
rmc.check4timegaps Boolean (default = FALSE). To indicate whether gaps in time should be imputed with zeros. Some sensing equipment provides accelerometer with gaps in time. The rest of GGIR is not designed for this, by setting this argument to TRUE the gaps in time will be filled with zeros.
rmc.col.wear Numeric (default $=$ NULL). If external wear detection outcome is stored as part of the data then this can be used by GGIR. This argument specifies the column in which the wear detection (Boolean) is stored.
rmc.doresample Boolean (default = FALSE). To indicate whether to resample the data based on the available timestamps and extracted sample rate from the file header.
rmc.noise Numeric (default $=13$ ). Noise level of acceleration signal in mg-units, used when working ad-hoc .csv data formats using read.myacc.csv. The read.myacc.csv does not take rmc.noise as argument, but when interacting with GGIR or g.part1 rmc.noise is used.
rmc.scalefactor.acc Numeric value (default 1) to scale the acceleration signals via multiplication. For example, if data is provided in $\mathrm{m} / \mathrm{s} 2$ then by setting this to $1 / 9.81$ we would derive gravitational units.
frequency_tol Number $($ default $=0.1)$ as passed on to readAxivity from the GGIRread package. Represents the frequency tolerance as fraction between 0 and 1 . When the relative bias per data block is larger than this fraction then the data block will be imputed by lack of movement with gravitational oriationed guessed from most recent valid data block. Only applicable to Axivity .cwa data.
params_metrics: A list of parameters used to specify the signal metrics that need to be extract in GGIR g.part1.
do.anglex Boolean (default $=$ FALSE). If TRUE, calculates the angle of the X axis relative to the horizontal:

$$
\operatorname{angle} X=\left(\tan ^{-1} \frac{a c c_{\text {rollmedian }(x)}}{\left(\operatorname{acc}_{\text {rollmedian }(y)}\right)^{2}+\left(\operatorname{acc}_{\text {rollmedian }(z)}\right)^{2}}\right) * 180 / \pi
$$

do.angley Boolean (default $=$ FALSE). If TRUE, calculates the angle of the Y axis relative to the horizontal:

$$
\operatorname{angle} Y=\left(\tan ^{-1} \frac{a c c_{\text {rollmedian }(y)}}{\left(\operatorname{acc} c_{\text {rollmedian }(x)}\right)^{2}+\left(\operatorname{acc}_{\text {rollmedian }(z)}\right)^{2}}\right) * 180 / \pi
$$

do.anglez Boolean (default $=$ TRUE). If TRUE, calculates the angle of the Z axis relative to the horizontal:

$$
\text { angle } Z=\left(\tan ^{-1} \frac{a c c_{\text {rollmedian }(z)}}{\left(\operatorname{acc} c_{\text {rollmedian }(x)}\right)^{2}+\left(\operatorname{acc}_{\text {rollmedian }(y)}\right)^{2}}\right) * 180 / \pi
$$

do.zex Boolean (default $=$ FALSE). If TRUE, calculates metric zero-crossing count for x -axis. For computation specifics see source code of function g.applymetrics
do.zcy Boolean (default = FALSE). If TRUE, calculates metric zero-crossing count for y-axis. For computation specifics see source code of function g.applymetrics
do.zcz Boolean (default = FALSE). If TRUE, calculates metric zero-crossing count for z-axis. For computation specifics see source code of function g.applymetrics
do.enmo Boolean (default $=$ TRUE). If TRUE, calculates the metric:

$$
E N M O=\sqrt{a c c_{x}^{2}+a c c_{y}^{2}+a c c_{z}^{2}}-1
$$

(if ENMO $<0$, then ENMO $=0$ ).
do.lfenmo Boolean (default = FALSE). If TRUE, calculates the metric ENMO over the low-pass filtered accelerations (for computation specifics see source code of function g.applymetrics). The filter bound is defined by the parameter hb .
do.en Boolean (default $=$ FALSE). If TRUE, calculates the Euclidean Norm of the raw accelerations:

$$
E N=\sqrt{a c c_{x}^{2}+a c c_{y}^{2}+a c c_{z}^{2}}
$$

do.mad Boolean (default $=$ FALSE $)$. If TRUE, calculates the Mean Amplitude Deviation:

$$
M A D=\frac{1}{n} \Sigma\left|r_{i}-\bar{r}\right|
$$

do.enmoa Boolean (default $=$ FALSE). If TRUE, calculates the metric:

$$
E N M O a=\sqrt{a c c_{x}^{2}+a c c_{y}^{2}+a c c_{z}^{2}}-1
$$

(if ENMOa $<0$, then ENMOa $=\mid$ ENMOal).
do.roll_med_acc_x Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.roll_med_acc_y Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.roll_med_acc_z Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.dev_roll_med_acc_x Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.dev_roll_med_acc_y Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.dev_roll_med_acc_z Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.bfen Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.hfen Boolean (default $=$ FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.hfenplus Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.lfen Boolean (default $=$ FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.lfx Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.lfy Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.lfz Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.hfx Boolean (default $=$ FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.hfy Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.hfz Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.bfx Boolean (default = FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.bfy Boolean (default $=$ FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.bfz Boolean (default $=$ FALSE). If TRUE, calculates the metric. For computation specifics see source code of function g.applymetrics.
do.brondcounts Boolean (default = FALSE). this option has been deprecated (October 2022) due to issues with the activityCounts package that we used as a dependency. If TRUE, calculated the metric via R package activityCounts. We called them BrondCounts because there are large number of activity counts in the physical activity and sleep research field. By calling them _brondcounts_ we clarify that these are the counts proposed by Jan Br\&\#248;nd and implemented in R by Ruben Brondeel. The _brondcounts_ are intended to be an imitation of the counts produced by one of the closed source ActiLife software by ActiGraph.
do.neishabouricounts Boolean (default = FALSE). If TRUE, calculates the metric via R package actilifecounts, which is an implementation of the algorithm used in the closed-source software ActiLife by ActiGraph (methods published in doi: 10.1038/s41598-022-16003-x). We use the name of the first author (instead of ActiLifeCounts) of the paper and call them NeishabouriCount under the uncertainty that ActiLife will implement this same algorithm over time. To use the Neishabouri counts for the physical activity intensity classification in part 5 (i.e., metric over the threshold.lig, threshold.mod, and threshold.vig would be applied), the acc.metric argument needs to be set as one of the following: "NeishabouriCount_x", "NeishabouriCount_y", "NeishabouriCount_z", "NeishabouriCount_vm" to use the counts in the $\mathrm{x}-, \mathrm{y}$-, z -axis or vector magnitude, respectively.
lb Numeric (default $=0.2$ ). Lower boundary of the frequency filter (in Hertz) as used in the filter-based metrics.
hb Numeric (default $=15$ ). Higher boundary of the frequency filter (in Hertz) as used in the filter-based metrics.
n Numeric $($ default $=\mathrm{n})$. Order of the frequency filter as used in the filter-based metrics.
zc.lb Numeric (default $=0.25$ ). Used for zero-crossing counts only. Lower boundary of cut-off frequency filter.
zc.hb Numeric (default = 3). Used for zero-crossing counts only. Higher boundary of cut-off frequencies in filter.
zc.sb Numeric (default $=0.01$ ). Stop band used for calculation of zero crossing counts. Value is the acceleration threshold in $g$ units below which acceleration will be rounded to zero.
zc.order Numeric (default = 2). Used for zero-crossing counts only. Order of frequency filter.
zc.scale Numeric (default =1) Used for zero-crossing counts only. Scaling factor to be applied after counts are calculated (GGIR part 3).
actilife_LFE Boolean (default = FALSE). If TRUE, calculates the NeishabouriCount metric with the low-frequency extension filter as proposed in the closed source ActiLife software by ActiGraph. Only applicable to the metric NeishabouriCount.
params_cleaning: A list of parameters used across all GGIR parts releated to masking or imputing data, abbreviated as "cleaning".
do.imp Boolean (default = TRUE). Whether to impute missing values (e.g., suspected of monitor non-wear or clippling) or not by g.impute in GGIR g.part2. Recommended setting is TRUE.
TimeSegments2ZeroFile Character (default = NULL). Takes path to a csv file that has columns "windowstart" and "windowend" to refer to the start and end time of a time windows in format "2024-10-12 20:00:00", and "filename" of the GGIR milestone data file without the "meta_" segment of the name. GGIR part 2 uses this to set all acceleration values to zero and the nonwear classification to zero (meaning sensor worn). Motivation: When the accelerometer is not worn during the night GGIR automatically labels them as invalid, while the user may like to treat them as zero movement. Disclaimer: This functionality was developed in 2019. With hindsight it is not generic enough and in need for revision. Please contact GGIR maintainers if you would like us to invest time in improving this functionality.
data_cleaning_file Character (default = NULL). Optional path to a csv file you create that holds four columns: ID, day_part5, relyonguider_part4, and night_part4. ID should hold the participant ID. Columns day_part5 and night_part4 allow you to specify which day(s) and night(s) need to be excluded from g.part5 and g.part4, respectively. When including multiple day(s)/night(s) create a new line for each day/night. So, this will be done regardless of whether the rest of GGIR thinks those day(s)/night(s) are valid. Column relyonguider_part4 allows you to specify for which nights g.part4 should fully rely on the guider. See also package vignette.
excludefirstlast.part5 Boolean (default = FALSE). If TRUE then the first and last window (wakingwaking, midnight-midnight, or sleep onset-onset) are ignored in g.part5.
excludefirstlast Boolean (default $=$ FALSE). If TRUE then the first and last night of the measurement are ignored for the sleep assessment in g.part4.
excludefirst.part4 Boolean (default = FALSE). If TRUE then the first night of the measurement are ignored for the sleep assessment in g.part4.
excludelast.part4 Boolean (default $=$ FALSE). If TRUE then the last night of the measurement are ignored for the sleep assessment in g.part4.
includenightcrit Numeric (default $=16$ ). Minimum number of valid hours per night ( 24 hour window between noon and noon), used for sleep assessment in g.part4.
minimum_MM_length.part5 Numeric (default = 23). Minimum length in hours of a MM day to be included in the cleaned g.part5 results.
study_dates_file Character $($ default $=c())$. Full path to csv file containing the first and last date of the expected wear period for every study participant (dates are provided per individual). Expected format of the activity diary is: First column headers followed by one row per recording. There should be three columns: first column is recording ID, which needs to match with the ID GGIR extracts from the accelerometer file; second column should contain the first date of the study; and third column the last date of the study. Date columns should be by default in format "23-04-2017", or in the date format specified by argument
study_dates_dateformat (below). If not specified (default), then GGIR would use the first and last day of the recording as beginning and end of the study. Note that these dates are used on top of the data_masking_strategy selected.
study_dates_dateformat Character (default = " To specify the date format used in the study_dates_file as used by strptime.
strategy Deprecated and replaced by data_masking_strategy. If strategy is specified then its value is passed on and used for data_masking_strategy.
data_masking_strategy Numeric (default = 1). How to deal with knowledge about study protocol. data_masking_strategy $=1$ means select data based on hrs.del.start and hrs.del.end. data_masking_strategy $=2$ makes that only the data between the first midnight and the last midnight is used. data_masking_strategy $=3$ selects the most active X days in the file where X is specified by argument ndayswindow, where the days are a series of 24 -h blocks starting any time in the day ( X hours at the beginning and end of this period can be deleted with arguments hrs.del.start and hrs.del.end) data_masking_strategy $=4$ to only use the data after the first midnight. data_masking_strategy $=5$ is similar to data_masking_strategy $=$ 3, but it selects $X$ complete calendar days where $X$ is specified by argument ndayswindow ( X hours at the beginning and end of this period can be deleted with arguments hrs.del.start and hrs.del.end).
hrs.del.start Numeric (default = 0). How many HOURS after start of experiment did wearing of monitor start? Used in GGIR g.part2 when data_masking_strategy $=1$.
hrs.del.end Numeric (default $=0$ ). How many HOURS before the end of the experiment did wearing of monitor definitely end? Used in GGIR g.part2 when data_masking_strategy = 1.
maxdur Numeric (default $=0$ ). How many DAYS after start of experiment did experiment definitely stop? (set to zero if unknown).
ndayswindow Numeric (default = 7). If data_masking_strategy is set to 3 or 5, then this is the size of the window as a number of days. For data_masking_strategy 3 value can be fractional, e.g. 7.5, while for data_masking_strategy 5 it needs to be an integer.
includedaycrit.part5 Numeric (default $=2 / 3$ ). Inclusion criteria used in part 5 for number of valid hours during the waking hours of a day, when value is smaller than or equal to 1 used as fraction of waking hours, when value above 1 used as absolute number of valid hours required. Do not confuse this argument with argument includedaycrit which is only used in GGIR part 2 and applies to the entire day.
segmentWEARcrit.part5 Numeric (default $=0.5$ ). Fraction of qwindow segment expected to be valid in part 5 , where 0.3 indicates that at least 30 percent of the time should be valid.
segmentDAYSPTcrit.part5 Numeric vector or length 2 (default $=c(0.9,0)$ ). Inclusion criteria for the proportion of the segment that should be classified as day (awake) and spt (sleep period time) to be considered valid. If you are interested in comparing time spent in behaviour then it is better to set one of the two numbers to 0 , and the other defines the proportion of the segment that should be classified as day or spt, respectively. The default setting would focus on waking hour segments and includes all segments that overlap for at least 90 percent with waking hours. In order to shift focus to the SPT you could use $c(0,0.9)$ which ensures that all segments that overlap for at least 90 percent with the SPT are included. Setting both to zero would be problematic when comparing time spent in behaviours between days or individuals: A complete segment would be averaged with an incomplete segments (someone going to bed or waking up in the middle of a segment) by which it is no longer clear whether the person is less active or sleeps more during that segment. Similarly it is not clear whether the person has more wakefulness during SPT for a segment or woke up or went to bed during the segment.
includedaycrit Numeric (default $=16$ ). Minimum required number of valid hours in calendar day specific to analysis in part 2. If you specify two values as in $c(16,16)$ then the first value will be used in part 2 and the second value will be used in part 5 and applied as a criterion on the full part 5 window. Note that this is then applied in addition to parameter includedaycrit.part5 which only looks at valid data during waking hours.
max_calendar_days Numeric (default $=0$ ). The maximum number of calendar days to include (set to zero if unknown).
nonWearEdgeCorrection Boolean (default $=$ TRUE). If TRUE then the non-wear detection around the edges of the recording (first and last 3 hours) are corrected following description in vanHees2013 as has been the default since then. This functionality is advisable when working with sleep clinic or exercise lab data typically lasting less than a day.
nonwear_approach Character (default = "2023"). Whether to use the traditional version of the non-wear detection algorithm (nonwear_approach $=" 2013 "$ ) or the new version (nonwear_approach $=$ " 2023 "). The 2013 version would use the longsize window (windowsizes[3], one hour as default) to check the conditions for nonwear identification and would flag as nonwear the mediumsize window (windowsizes[2], 15 min as default) in the middle. The 2023 version differs in which it would flag as nonwear the full longsize window. For the 2013 method the longsize window is centered in the centre of the mediumsize window, while in the 2023 method the longsizewindow is aligned with its left edge to the left edge of the mediumsize window.
params_phyact: A list of parameters releated to physical activity as used in GGIR g.part2 and GGIR g.part5.
mvpathreshold Numeric (default $=100$ ). Acceleration threshold for MVPA estimation in GGIR g.part2. This can be a single number or an vector of numbers, e.g., mvpathreshold $=c(100$, 120). In the latter case the code will estimate MVPA separately for each threshold. If this variable is left blank, e.g., mvpathreshold $=c()$, then MVPA is not estimated.
mppadur Numeric $($ default $=10)$. The bout duration $(s)$ for which MVPA will be calculated. Only used in GGIR g.part2.
boutcriter Numeric (default $=0.8$ ). A number between 0 and 1 , it defines what fraction of a bout needs to be above the mvpathreshold, only used in GGIR g.part2.
threshold.lig Numeric (default $=40$ ). In g.part5: Threshold for light physical activity to separate inactivity from light. Value can be one number or an vector of multiple numbers, e.g., threshold. $\operatorname{lig}=c(30,40)$. If multiple numbers are entered then analysis will be repeated for each combination of threshold values. Threshold is applied to the first metric in the milestone data, so if you have only specified do. enmo = TRUE then it will be applied to ENMO.
threshold.mod Numeric $($ default $=100)$. In g.part5: Threshold for moderate physical activity to separate light from moderate. Value can be one number or an vector of multiple numbers, e.g., threshold. $\bmod =c(100,120)$. If multiple numbers are entered then analysis will be repeated for each combination of threshold values. Threshold is applied to the first metric in the milestone data, so if you have only specified do. enmo $=$ TRUE then it will be applied to ENMO.
threshold.vig Numeric (default $=400$ ). In g.part5: Threshold for vigorous physical activity to separate moderate from vigorous. Value can be one number or an vector of multiple numbers, e.g., threshold.vig $=c(400,500)$. If multiple numbers are entered then analysis will be repeated for each combination of threshold values. Threshold is applied to the first metric in the milestone data, so if you have only specified do. enmo $=$ TRUE then it will be applied to ENMO.
boutdur.mypa Numeric (default $=\mathrm{c}(1,5,10)$ ). Duration(s) of MVPA bouts in minutes to be extracted. It will start with the identification of the longest to the shortest duration. In the default setting, it will start with the 10 minute bouts, followed by 5 minute bouts in the rest of the data, and followed by 1 minute bouts in the rest of the data.
boutdur.in Numeric (default $=\mathrm{c}(10,20,30)$ ). Duration(s) of inactivity bouts in minutes to be extracted. Inactivity bouts are detected in the segments of the data which were not labelled as sleep or MVPA bouts. It will start with the identification of the longest to the shortest duration. In the default setting, it will start with the identification of 30 minute bouts, followed by 20 minute bouts in the rest of the data, and followed by 10 minute bouts in the rest of the data. Note that we use the term inactivity instead of sedentary behaviour for the lowest intensity level of behaviour. The reason for this is that GGIR does not attempt to classifying the activity type sitting at the moment, by which we feel that using the term sedentary behaviour would fail to communicate that.
boutdur.lig Numeric (default $=\mathrm{c}(1,5,10)$ ). Duration(s) of light activity bouts in minutes to be extracted. Light activity bouts are detected in the segments of the data which were not labelled as sleep, MVPA, or inactivity bouts. It will start with the identification of the longest to the shortest duration. In the default setting, this will start with the identification of 10 minute bouts, followed by 5 minute bouts in the rest of the data, and followed by 1 minute bouts in the rest of the data.
boutcriter.mvpa Numeric (default $=0.8$ ). A number between 0 and 1, it defines what fraction of a bout needs to be above the threshold.mod.
boutcriter.in Numeric (default $=0.9$ ). A number between 0 and 1, it defines what fraction of a bout needs to be below the threshold. lig.
boutcriter.lig Numeric (default $=0.8$ ). A number between 0 and 1 , it defines what fraction of a bout needs to be between the threshold.lig and the threshold.mod.
frag.metrics Character (default = NULL). Fragmentation metric to extract. Can be "mean", "TP", "Gini", "power", or "CoV", "NFragPM", or all the above metrics with "all". See package vignette for description of fragmentation metrics.
part6_threshold_combi Character (default = "40_100_120") to indicate the threshold combination derived in part 5 to be used for part 6
params_sleep: A list of parameters used to configure the sleep analysis as performend in GGIR g.part3 and g.part4.
relyonguider Boolean (default $=$ FALSE). Sustained inactivity bouts (sib) that overlap with the guider are labelled as sleep. If relyonguider $=$ FALSE and the sib overlaps only partially with the guider then it is the sib that defines the edge of the SPT window and not the guider. If relyonguider = TRUE and the sib overlaps only partially with the guider then it is the guider that defines the edge of the SPT window and not the sib. If participants were instructed NOT to wear the accelerometer during waking hours and ignorenonware=FALSE then set to relyonguider=TRUE, in all other scenarios set to FALSE.
relyonsleeplog Boolean (default $=$ FALSE). Do not use, now replaced by argument relyonguider. Values provided to argument relyonsleeplog will be passed on to argument relyonguider to not preserve functionality of old R scripts.
def.noc.sleep Numeric (default = 1). The time window during which sustained inactivity will be assumed to represent sleep, e.g., def.noc.sleep $=c(21,9)$. This is only used if no sleep log entry is available. If left blank def.noc. sleep $=c()$ then the 12 hour window centred at the least active 5 hours of the 24 hour period will be used instead. Here, L5 is hardcoded
and will not change by changing argument winhr in function g .part2. If def.noc.sleep is filled with a single integer, e.g., def. noc. sleep=c(1) then the window will be detected with based on built in algorithms. See argument HASPT. algo from HASPT for specifying which of the algorithms to use.
sleepwindowType Character (default = "SPT"). To indicate type of information in the sleeplog, "SPT" for sleep period time. Set to "TimeInBed" if sleep log recorded time in bed to enable calculation of sleep latency and sleep efficiency.
nnights Numeric (default = NULL). This argument has been deprecated.
loglocation Character (default = NULL). Path to csv file with sleep log information. See package vignette for how to format this file.
colid Numeric (default $=1$ ). Column number in the sleep $\log$ spreadsheet in which the participant ID code is stored.
coln1 Numeric $($ default $=2)$. Column number in the sleep $\log$ spreadsheet where the onset of the first night starts.
ignorenonwear Boolean (default = TRUE). If TRUE then ignore detected monitor non-wear periods to avoid confusion between monitor non-wear time and sustained inactivity.
constrain2range Deprecated, used to be a Boolean (default = TRUE) Whether or not to constrain the range of threshold used in the diary free sleep period time window detection.
HASPT.algo Character (default = "HDCZA"). To indicate what algorithm should be used for the sleep period time detection. Default "HDCZA" is Heuristic algorithm looking at Distribution of Change in Z-Angle as described in van Hees et al. 2018. Other options included: "HorAngle", which is based on HDCZA but replaces non-movement detection of the HDCZA algorithm by looking for time segments where the angle of the longitudinal sensor axis has an angle relative to the horizontal plane between -45 and +45 degrees. And "NotWorn" which is also the same as HDCZA but looks for time segments when a rolling average of acceleration magnitude is below 5 per cent of its standard deviation, see Cookbook vignette in the Annexes of https://wadpac.github.io/GGIR/ for more detailed guidance on how to use "NotWorn".
HDCZA_threshold Numeric (default = $\mathbf{c}()$ ) If HASPT. algo is set to "HDCZA" and HDCZA_threshold is NOT NULL, (e.g., HDCZA_threshold $=0.2$ ), then that value will be used as threshold in the 6th step in the diagram of Figure 1 in van Hees et al. 2018 Scientific Report (doi: $10.1038 / \mathrm{s} 41598-018-31266-\mathrm{z}$ ). However, doing so has not been supported by research yet and is only intended to facilitate methodological research, so we advise sticking with the default in line with the publication. Further, if HDCZA_threshold is set to a numeric vector of length 2, e.g. c( 10,15 ), that will be used as percentile and multiplier for the above mentioned 6th step.
HASPT.ignore.invalid Boolean (default = FALSE). To indicate whether invalid time segments should be ignored in the Sleep Period Time detection. If FALSE (default), the imputed angle or activity metric during the invalid time segments is used in the Sleep Period Time detection. If TRUE, invalid time segments are ignored for the Sleep Period Time detection (i.e., considered to be out of the Sleep Period Time). If NA, then invalid time segments are considered to be no movement segments.
HASIB.algo Character (default = "vanHees2015"). To indicate which algorithm should be used to define the sustained inactivity bouts (i.e., likely sleep). Options: "vanHees2015", "Sadeh1994", "Galland2012".
Sadeh_axis Character (default = "Y"). To indicate which axis to use for the Sadeh1994 algorithm, and other algortihms that relied on count-based Actigraphy such as Galland2012.
sleeplogsep Character (default $=$ NULL). This argument is deprecated.
nap_model Character (default $=$ NULL). To specify classification model. Currently the only option is "hip3yr", which corresponds to a model trained with hip data in 3-3.5 olds trained with parent diary data.
longitudinal_axis Integer (default = NULL). To indicate which axis is the longitudinal axis. If not provided, the function will estimate longitudinal axis as the axis with the highest 24 hour lagged autocorrelation. Only used when sensor.location = "hip" or HASPT. algo = "HorAngle".
anglethreshold Numeric (default =5). Angle threshold (degrees) for sustained inactivity periods detection. The algorithm will look for periods of time (timethreshold) in which the angle variability is lower than anglethreshold. This can be specified as multiple thresholds, each of which will be implemented, e.g., anglethreshold $=c(5,10)$.
timethreshold Numeric (default $=5$ ). Time threshold (minutes) for sustained inactivity periods detection. The algorithm will look for periods of time (timethreshold) in which the angle variability is lower than anglethreshold. This can be specified as multiple thresholds, each of which will be implemented, e.g., timethreshold $=c(5,10)$.
possible_nap_window Numeric (default $=c(9,18)$ ). Numeric vector of length two with range in clock hours during which naps are assumed to take place, e.g., possible_nap_window = $c(9,18)$. Currently used in the context of an explorative nap classification algortihm that was trained in 3.5 year olds.
possible_nap_dur Numeric $($ default $=c(15,240)$ ). Numeric vector of length two with range in duration (minutes) of a nap, e.g., possible_nap_dur $=c(15,240)$. Currently used in the context of an explorative nap classification algortihm that was trained in 3.5 year olds.
sleepefficiency.metric Numeric (default $=1$ ). If 1 (default), sleep efficiency is calculated as detected sleep time during the SPT window divided by log-derived time in bed. If 2 , sleep efficiency is calculated as detected sleep time during the SPT window divided by detected duration in sleep period time plus sleep latency (where sleep latency refers to the difference between time in bed and sleep onset). sleepefficiency.metric is only considered when argument sleepwindowType = "TimeInBed"
possible_nap_edge_acc Numeric (default = Inf). Maximum acceleration before or after the SIB for the nap to be considered. By default this will allow all possible naps.
params_247: A list of parameters releated to description of 24/7 behaviours that do not fall under conventional physical activity or sleep outcomes, these parameters are used in GGIR g.part2 and GGIR g.part5:
qwindow Numeric or character (default $=c(0,24)$ ). To specify windows over which all variables are calculated, e.g., acceleration distribution, number of valid hours, LXMX analysis, MVPA. If numeric, qwindow should have length two, e.g., qwindow $=c(0,24)$, all variables will only be calculated over the full 24 hours in a day. If qwindow $=c(8,24)$ variables will be calculated over the window $0-8,8-24$ and $0-24$. All days in the recording will be segmented based on these values. If you want to use a day specific segmentation in each day then you can set qwindow to be the full path to activity diary file (character). Expected format of the activity diary is: First column headers followed by one row per recording, first column is recording ID, which needs to match with the ID GGIR extracts from the accelerometer file. Followed by date column in format "23-04-2017", where date format is specified by argument qwindow_dateformat (below). Use the character combination date, Date or DATE in the column name. This is followed by one or multiple columns with start times for the
activity types in that day format in hours:minutes:seconds. The header of the column will be used as label for each activity type. Insert a new date column before continuing with activity types for next day. Leave missing values empty. If an activity $\log$ is used then individuals who do not appear in the activity log will still be processed with value qwindow $=c(0,24)$. Dates with no activity log data can be skipped, no need to have a column with the date followed by a column with the next date. If times in the activity diary are not multiple of the short window size (epoch length), the next epoch is considered (e.g., with epoch of 5 seconds, 8:00:02 will be redefined as 8:00:05 in the activity $\log$ ). When using the qwindow functionality in combination with GGIR part 5 then make sure to check that arguments segmentWEARcrit. part5 and segmentDAYSPTcrit.part5 are specified to your research needs.
qwindow_dateformat Character (default = " To specify the date format used in the activity log as used by strptime.
M5L5res Numeric (default $=10$ ). Resolution of L5 and M5 analysis in minutes.
winhr Numeric (default $=5$ ). Vector of window size(s) (unit: hours) of LX and MX analysis, where look for least and most active consecutive number of X hours.
qlevels Numeric (default = NULL). Vector of percentiles for which value needs to be extracted. These need to be expressed as a fraction of 1 , e.g., $c(0.1,0.5,0.75)$. There is no limit to the number of percentiles. If left empty then percentiles will not be extracted. Distribution will be derived from short epoch metric data. Argument qlevels can for example be used for the MX-metrics (e.g. Rowlands et al) as discussed in the main package vignette
ilevels Numeric (default $=$ NULL). Levels for acceleration value frequency distribution in $\mathrm{m} g$, e.g., ilevels $=c(0,100,200)$. There is no limit to the number of levels. If left empty then the intensity levels will not be extracted. Distribution will be derived from short epoch metric data.
iglevels Numeric $($ default $=$ NULL $)$. Levels for acceleration value frequency distribution in $\mathrm{m} g$ used for intensity gradient calculation (according to the method by Rowlands 2018). By default this is argument is empty and the intensity gradient calculation is not done. The user can either provide a single value (any) to make the intensity gradient use the bins iglevels $=c(\operatorname{seq}(0,4000, b y=25), 8000)$ or the user could specify their own distribution. There is no constriction to the number of levels.
IVIS_windowsize_minutes Numeric (default =60). Window size of the Intradaily Variability (IV) and Interdaily Stability (IS) metrics in minutes, needs to be able to add up to 24 hours.

IVIS_epochsize_seconds Numeric (default = NULL). This argument is deprecated.
IVIS.activity.metric Numeric (default =2). Metric used for activity calculation. Value = 1, uses continuous scaled acceleration. Value $=2$, tries to collapse acceleration into a binary score of rest versus active to try to simulate the original approach.
IVIS_acc_threshold Numeric (default $=20$ ). Acceleration threshold to distinguish inactive from active.
qM5L5 Numeric (default = NULL). Percentiles (quantiles) to be calculated over L5 and M5 window.
MX.ig.min.dur Numeric (default $=10$ ). Minimum MX duration needed in order for intensity gradient to be calculated.
LUXthresholds Numeric (default $=c(0,100,500,1000,3000,5000,10000)$ ). Vector with numeric sequence corresponding to the thresholds used to calculate time spent in LUX ranges.
LUX_cal_constant Numeric (default = NULL). If both LUX_cal_constant and LUX_cal_exponent are provided LUX values are converted based on formula $y=$ constant $* \exp (x * \operatorname{exponent})$

LUX_cal_exponent Numeric (default = NULL). If both LUX_cal_constant and LUX_cal_exponent are provided LUX LUX values are converted based on formula $y=$ constant $* \exp (x *$ exponent)
LUX_day_segments Numeric (default = NULL). Vector with hours at which the day should be segmented for the LUX analysis.
L5M5window Argument deprecated after version 1.5-24. This argument used to define the start and end time, in 24 hour clock hours, over which L5M5 needs to be calculated. Now this is done with argument qwindow.
cosinor Boolean (default = FALSE). Whether to apply the cosinor analysis from the ActCR package.
part6CR Boolean (default $=$ FALSE) to indicate whether circadian rhythm analysis should be run by part 6.
part6HCA Boolean (default $=$ FALSE) to indicate whether Household Co Analysis should be run by part 6 .
part6Window Character vector with length two (default $=c($ "start", "end")) to indicate the start and the end of the time series to be used for circadian rhythm analysis in part 6 . In other words, this parameters is not used for Household co-analysis. Alternative values are: "Wx", "Ox", "Hx", where "x" is a number to indicat the xth wakeup, onset or hour of the recording. Negative values for " $x$ " are also possible and will count relative to the end of the recording. For example, c("W1", "W-1") goes from the first till the last wakeup, c("H5", "H-5") ignores the first and last 5 hours, and $\mathrm{c}($ "O2", "W10") goes from the second onset till the 10th wakeup time.
params_output: A list of parameters used to specify whether and how GGIR stores its output at various stages of the process.
storefolderstructure Boolean (default = FALSE). Store folder structure of the accelerometer data.
do.part2.pdf Boolean (default $=$ TRUE). In g.part2: Whether to generate a pdf for g.part2.
do.part3.pdf Boolean (default $=$ TRUE). In g.part3: Whether to generate a pdf for g.part3.
timewindow Character (default $=\mathrm{c}($ "MM", "WW")). In g.part5: Timewindow over which summary statistics are derived. Value can be "MM" (midnight to midnight), "WW" (waking time to waking time), "OO" (sleep onset to sleep onset), or any combination of them.
save_ms5rawlevels Boolean (default = FALSE). In g.part5: Whether to save the time series classification (levels) as csv or RData files (as defined by save_ms5raw_format). Note that time stamps will be stored in the column timenum in UTC format (i.e., seconds from 1970-01-01). To convert timenum to time stamp format, you need to specify your desired time zone, e.g., as.POSIXct (mdat\$timenum, tz = "Europe/London").
save_ms5raw_format Character (default = "csv"). In g.part5: To specify how data should be stored: either "csv" or "RData". Only used if save_ms5rawlevels = TRUE.
save_ms5raw_without_invalid Boolean (default = TRUE). In g.part5: To indicate whether to remove invalid days from the time series output files. Only used if save_ms5rawlevels = TRUE.
epochvalues2csv Boolean (default = FALSE). In g.part2: If TRUE then epoch values are exported to a csv file. Here, non-wear time is imputed where possible.
do.sibreport Boolean (default $=$ FALSE). In g.part4: To indicate whether to generate report for the sustained inactivity bouts (SIB). If set to TRUE and when an advanced sleep diary is
available in part 4 then part 5 will use this to generate summary statistics on the overlap between self-reported nonwear and napping with SIB. Here, SIB can be filter based on argument possible_nap_edge_acc and the first value of possible_nap_dur
do.visual Boolean (default $=$ TRUE). In g.part4: If TRUE, the function will generate a pdf with a visual representation of the overlap between the sleeplog entries and the accelerometer detections. This can be used to visually verify that the sleeplog entries do not come with obvious mistakes.
outliers.only Boolean (default = FALSE). In g.part4: Only used if do. visual = TRUE. If FALSE, all available nights are included in the visual representation of the data and sleeplog. If TRUE, then only nights with a difference in onset or waking time larger than the variable of argument criterror will be included.
criterror Numeric (default $=3$ ). In g.part4: Only used if do.visual $=$ TRUE and outliers.only $=$ TRUE. criterror specifies the number of minimum number of hours difference between sleep log and accelerometer estimate for the night to be included in the visualisation.
visualreport Boolean (default = TRUE). If TRUE, then generate visual report based on combined output from g.part2 and g.part4. Please note that the visual report was initially developed to provide something to show to study participants and not for data quality checking purposes. Over time we have improved the visual report to also be useful for QC-ing the data. However, some of the scorings as shown in the visual report are created for the visual report only and may not reflect the scorings in the main GGIR analyses as reported in the quantitative csvreports. Most of our effort in the past 10 years has gone into making sure that the csv-report are correct, while the visualreport has mostly been a side project. This is unfortunate and we hope to find funding in the future to design a new report specifically for the purpose of QC-ing the analyses done by GGIR.
viewingwindow Numeric (default = 1). Centre the day as displayed around noon (viewingwindow $=1$ ) or around midnight (viewingwindow $=2$ ) in the visual report generated with visualreport = TRUE.
week_weekend_aggregate.part5 Boolean (default = FALSE). In g.part5: To indicate whether week and weekend-days aggregates should be stored. This is turned off by default as it generates a large number of extra columns in the output report.
dofirstpage Boolean (default $=$ TRUE). To indicate whether a first page with histograms summarizing the whole measurement should be added in the file summary reports generated with visualreport = TRUE.
sep_reports Character (default = ","). Value used as sep argument in fwrite for writing csv reports.
dec_reports Character (default $=$ "."). Value used as dec argument in fwrite for writing csv reports.
sep_config Character (default = ","). Value used as sep argument in fwrite for writing csv config file.
dec_config Character (default = "."). Value used as dec argument in fwrite for writing csv config file.
visualreport_without_invalid Boolean (default = TRUE). If TRUE, then reports generated with visualreport = TRUE only show the windows with sufficiently valid data according to includedaycrit when viewingwindow $=1$ or includenightcrit when viewingwindow $=2$

## Value

The function provides no values, it only ensures that other functions are called and that their output is stored. Further, a configuration file is stored containing all the argument values used to facilitate reproducibility.

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## References

- van Hees VT, Gorzelniak L, Dean Leon EC, Eder M, Pias M, et al. (2013) Separating Movement and Gravity Components in an Acceleration Signal and Implications for the Assessment of Human Daily Physical Activity. PLoS ONE 8(4): e61691. doi:10.1371/journal.pone. 0061691
- van Hees VT, Fang Z, Langford J, Assah F, Mohammad A, da Silva IC, Trenell MI, White T, Wareham NJ, Brage S. Auto-calibration of accelerometer data for free-living physical activity assessment using local gravity and temperature: an evaluation on four continents. J Appl Physiol (1985). 2014 Aug 7
- van Hees VT, Sabia S, et al. (2015) A novel, open access method to assess sleep duration using a wrist-worn accelerometer, PLoS ONE, November 2015


## Examples

```
## Not run:
    mode = c(1,2,3,4,5)
    datadir = "C:/myfolder/mydata"
    outputdir = "C:/myresults"
    studyname ="test"
    f0 = 1
    f1 = 2
    GGIR(#-------------------------------
        # General parameters
        #------------------------------
        mode = mode,
        datadir = datadir,
        outputdir = outputdir,
        studyname = studyname,
        f0 = f0,
        f1 = f1,
        overwrite = FALSE,
        do.imp = TRUE,
        idloc = 1,
        print.filename = FALSE,
        storefolderstructure = FALSE,
        #------------------------------
        # Part 1 parameters:
        #------------------------------
        windowsizes = c(5,900,3600),
        do.cal = TRUE,
        do.enmo = TRUE,
```

```
do.anglez = TRUE,
chunksize = 1,
printsummary = TRUE,
#-------------------------------
# Part 2 parameters:
#-----------------------------
data_masking_strategy = 1,
ndayswindow = 7,
hrs.del.start = 1,
hrs.del.end = 1,
maxdur = 9,
includedaycrit = 16,
L5M5window = c(0,24),
M5L5res = 10,
winhr = c(5,10),
qlevels = c(c(1380/1440),c(1410/1440)),
qwindow = c(0,24),
ilevels = c(seq(0,400,by=50), 8000),
mvpathreshold = c(100,120),
#-------------------------------
# Part 3 parameters:
#-------------------------------
timethreshold = c(5,10),
anglethreshold = 5,
ignorenonwear = TRUE,
#-------------------------------
# Part 4 parameters:
#-------------------------------
excludefirstlast = FALSE,
includenightcrit = 16,
def.noc.sleep = 1,
loglocation = "D:/sleeplog.csv",
outliers.only = FALSE,
criterror = 4,
relyonguider = FALSE,
colid = 1,
coln1 = 2,
do.visual = TRUE,
#-------------------------------
# Part 5 parameters:
#-------------------------------
# Key functions: Merging physical activity with sleep analyses
threshold.lig = c(30,40,50),
threshold.mod = c(100,120),
threshold.vig = c(400,500),
excludefirstlast = FALSE,
boutcriter = 0.8,
boutcriter.in = 0.9,
boutcriter.lig = 0.8,
boutcriter.mvpa = 0.8,
boutdur.in = c(10,20,30),
boutdur.lig = c(1,5,10),
boutdur.mvpa = c(1,5,10),
```

```
timewindow = c("WW"),
#-----------------------------------
# Report generation
#---------------------------------
do.report = c(2,4,5))
# For externally derived Actiwatch data in .AWD format:
GGIR(datadir = "/media/actiwatch_awd", # folder with epoch level .AWD file
    outputdir = "/media/myoutput",
    dataFormat = "actiwatch_awd",
    extEpochData_timeformat = "%m/%d/%Y %H:%M:%S",
    mode = 1:5,
    do.report = c(2, 4, 5),
    windowsizes = c(60, 900, 3600), # 60 is the expected epoch length
    visualreport = FALSE,
    outliers.only = FALSE,
    overwrite = TRUE,
    HASIB.algo = "Sadeh1994",
    def.noc.sleep = c()) # <= because we cannot use HDCZA for ZCY
# For externally derived Actiwatch data in .CSV format:
GGIR(datadir = "/media/actiwatch_csv", # folder with epoch level .AWD file
    outputdir = "/media/myoutput",
    dataFormat = "actiwatch_csv",
    extEpochData_timeformat = "%m/%d/%Y %H:%M:%S",
    mode = 1:5,
    do.report = c(2, 4, 5),
    windowsizes = c(15, 900, 3600), # 15 is the expected epoch length
    visualreport = FALSE,
    outliers.only = FALSE,
    HASIB.algo = "Sadeh1994",
    def.noc.sleep = c()) # <= because we cannot use HDCZA for ZCY
# For externally derived UK Biobank data in .CSV format:
GGIR(datadir = "/media/ukbiobank",
    outputdir = "/media/myoutput",
    dataFormat = "ukbiobank_csv",
    extEpochData_timeformat = "%m/%d/%Y %H:%M:%S",
    mode = c(1:2),
    do.report = c(2),
    windowsizes = c(5, 900, 3600), # We know that data was stored in 5 second epoch
        desiredtz = "Europe/London", # We know that data was collected in the UK
        visualreport = FALSE,
        overwrite = TRUE)
# For externally derived ActiGraph count data in .CSV format assuming
# a study protocol where sensor was not worn during the night:
GGIR(datadir = "/examplefiles",
    outputdir = "",
    dataFormat = "actigraph_csv",
    mode = 1:5,
    do.report = c(2, 4, 5),
    windowsizes = c(5, 900, 3600),
```

```
    threshold.in = round(100 * (5/60), digits = 2),
    threshold.mod = round(2500 * (5/60), digits = 2),
    threshold.vig = round(10000 * (5/60), digits = 2),
    extEpochData_timeformat = "%m/%d/%Y %H:%M:%S",
    do.neishabouricounts = TRUE,
    acc.metric = "NeishabouriCount_x",
    HASPT.algo = "NotWorn",
    HASIB.algo = "NotWorn",
    do.visual = TRUE,
    includedaycrit = 10,
    includenightcrit = 10,
    visualreport = FALSE,
    outliers.only = FALSE,
    save_ms5rawlevels = TRUE,
    ignorenonwear = FALSE,
    HASPT.ignore.invalid = FALSE,
    save_ms5raw_without_invalid = FALSE)
        # For externally derived Sensear data in .xls format:
        GGIR(datadir = "C:/yoursenseweardatafolder",
            outputdir = "D:/youroutputfolder",
            mode = 1:5,
            windowsizes = c(60, 900, 3600),
            threshold.in = 1.5,
            threshold.mod = 3,
            threshold.vig = 6,
            dataFormat = "sensewear_xls",
            extEpochData_timeformat = "%d-%b-%Y %H:%M:%S",
            HASPT.algo = "NotWorn",
            desiredtz = "America/New_York",
            overwrite = TRUE,
            do.report = c(2, 4, 5),
            visualreport = FALSE)
```

\#\# End(Not run)
is.IS08601 Check whether character timestamp is in iso8601 format.

## Description

Checks whether timestamp stored in character format is in ISO8601 format or not

## Usage

is.IS08601(x)

## Arguments

x
Timestamps in character format either in ISO8601 or as "yyyy-mm-dd hh:mm:ss".

## Examples

```
x ="1980-1-1 18:00:00"
is.IS08601(x)
```

iso8601chartime2POSIX Convert iso8601 timestamps to POSIX timestamp

## Description

To avoid ambiguities when sharing and comparing timestamps. All timestamps are expressed in iso8601 format: https://en.wikipedia.org/wiki/ISO_8601 However, to generate plots in R we need to convert them back to POSIX

## Usage

iso8601chartime2POSIX( $\mathrm{x}, \mathrm{tz}$ )

## Arguments

x
tz

Vector of timestamps in iso8601 in character format
Timezone of data collection, e.g. "Europe/London". See List_of_tz_database_time_zones on Wikipedia for full list.

## Examples

```
x ="2017-05-07T13:00:00+0200"
tz = "Europe/Amsterdam"
x_converted = iso8601chartime2POSIX(x,tz)
```

load_params Load default parameters

## Description

Loads default paramter values Not intended for direct use by GGIR users.

## Usage

```
load_params(topic = c("sleep", "metrics", "rawdata", "247",
    "phyact", "cleaning", "output", "general"))
```


## Arguments

topic Character vector with parameter groups to be loaded.

## Value

Lists of parameter objects

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

```
part6AlignIndividuals part6AlignIndividuals
```


## Description

Align individual time series per household where households are identified by the character or number string between the first and second '-' in the filename.

## Usage

part6AlignIndividuals(GGIR_ts_dir = NULL, outputdir = NULL,
path_ggirms $=$ NULL, desiredtz $=" "$, verbose $=$ TRUE)

## Arguments

GGIR_ts_dir Character, path to time series directory in the GGIR output
outputdir Directory where you would like to store the output
path_ggirms path to GGIR created folder named meta, with the milestone data files
desiredtz Character, specifying the timezone database name of the timezone the data was collected in.
verbose See details in GGIR.

## Value

no object is returned, only files are created in the output directory

```
part6PairwiseAggregation
                                    part6PairwiseAggregation
```


## Description

Pairwise aggregation of the time series of a group.

## Usage

part6PairwiseAggregation(outputdir = NULL, desiredtz = "", verbose = TRUE)

## Arguments

outputdir Directory where you would like to store your results
desiredtz Character, specifying the timezone database name of the timezone the data was collected in
verbose See details in GGIR.

## Value

No object is returned, only files are created in the output directory

```
POSIXtime2iso8601 Convert POSIX to iso8601 timestamp
```


## Description

To avoid ambiguities when sharing and comparing timestamps. All timestamps are expressed in iso8601 format: https://en.wikipedia.org/wiki/ISO_8601

## Usage

POSIXtime2iso8601(x, tz)

## Arguments

x
tz

Vector of timestamps in POSIX format
Timezone of data collection, e.g. "Europe/London". See https://en.wikipedia.org/wiki/List_of_tz_databas for full list

## Author(s)

Vincent T van Hees < v.vanhees@accelting.com>

## Examples

```
## Not run:
x ="2017-05-07 13:15:17 CEST"
tz = "Europe/Amsterdam"
x_converted = POSIXtime2iso8601(x,tz)
## End(Not run)
```

    read.myacc.csv Read custom csv files with accelerometer data
    
## Description

Loads csv files with accelerometer data and standardises the output format (incl. unit of measurement, timestamp format, header format, and column locations) to make the data compatible with other GGIR functions.

## Usage

```
read.myacc.csv(rmc.file=c(), rmc.nrow=Inf, rmc.skip = c(), rmc.dec=".",
    rmc.firstrow.acc = c(), rmc.firstrow.header=c(),
    rmc.header.length = c(),
    rmc.col.acc = 1:3, rmc.col.temp = c(),
    rmc.col.time=c(),
    rmc.unit.acc = "g", rmc.unit.temp = "C",
    rmc.unit.time = "POSIX",
    rmc.format.time = "%Y-%m-%d %H:%M:%OS",
    rmc.bitrate = c(), rmc.dynamic_range = c(),
    rmc.unsignedbit = TRUE,
    rmc.origin = "1970-01-01",
    rmc.desiredtz = NULL,
    rmc.configtz = NULL,
    rmc.sf = c(),
    rmc.headername.sf = c(),
    rmc.headername.sn = c(),
    rmc.headername.recordingid = c(),
    rmc.header.structure = c(),
    rmc.check4timegaps = FALSE,
    rmc.col.wear = c(),
    rmc.doresample = FALSE,
    rmc.scalefactor.acc = 1,
    interpolationType=1,
    PreviousLastValue = c(0, 0, 1),
    PreviousLastTime = NULL,
    desiredtz = NULL,
    configtz = NULL,
    header = NULL)
```


## Arguments

| rmc.file | Filename of file to be read if it is in the working directory, or full path to the file <br> otherwise. |
| :--- | :--- |
| rmc.nrow | Number of rows to read, same as nrow argument in read.csv and nrows in fread. <br> The whole file is read by default (i.e., rmc.nrow = Inf). |
| rmc.skip | Number of rows to skip, same as skip argument in read.csv and in fread. |
| rmc.dec | Decimal used for numbers, same as skip argument in read.csv and in fread. |
| rmc.firstrow. acc |  |$\quad$| First row (number) of the acceleration data. |
| :--- |

rmc. header. length
If file has header, specify header length (numeric).
rmc.col.acc Vector with three column (numbers) in which the acceleration signals are stored
rmc.col.temp Scalar with column (number) in which the temperature is stored. Leave in
default setting if no temperature is avaible. The temperature will be used by
g.calibrate.
rmc.col.time Scalar with column (number) in which the timestamps are stored. Leave in
default setting if timestamps are not stored.
rmc.unit.acc Character with unit of acceleration values: "g", "mg", or "bit"
rmc.unit.temp Character with unit of temperature values: (K)elvin, (C)elsius, or (F)ahrenheit
rmc.unit.time Character with unit of timestamps: "POSIX", "UNIXsec" (seconds since origin,
see argument rmc.origin), "character", or "ActivPAL" (exotic timestamp format
only used in the ActivPAL activity monitor).
rmc.format.time
Character string giving a date-time format as used by strptime. Only used for
rmc.unit.time: character and POSIX.
rmc.bitrate Numeric: If unit of acceleration is a bit then provide bit rate, e.g. 12 bit.
rmc.dynamic_range
Numeric, if unit of acceleration is a bit then provide dynamic range deviation in
g from zero, e.g. $+/-6 \mathrm{~g}$ would mean this argument needs to be 6 . If you give this
argument a character value the code will search the file header for elements with
a name equal to the character value and use the corresponding numeric value
next to it as dynamic range.
rmc.unsignedbit
Boolean, if unsignedbit $=$ TRUE means that bits are only positive numbers. if
unsignedbit $=$ FALSE then bits are both positive and negative.
rmc.origin Origin of time when unit of time is UNIXsec, e.g. 1970-1-1
rmc.desiredtz Deprecated, please see desiredtz.
rmc.configtz Deprecated, please see configtz.
rmc.sf Sample rate in Hertz, if this is stored in the file header then that will be used
instead.
rmc.headername.sf
If file has a header: Row name (character) under which the sample frequency can be found.
rmc.headername.sn
If file has a header: Row name (character) under which the serial number can be found.
rmc.headername.recordingid
If file has a header: Row name (character) under which the recording ID can be found.
rmc.header.structure
Character used to split the header name from the header value, e.g. ":" or " "
rmc.check4timegaps
Boolean to indicate whether gaps in time should be imputed with zeros. Some sensing equipment provides accelerometer with gaps in time. The rest of GGIR is not designed for this, by setting this argument to TRUE the the gaps in time will be filled with zeros.
rmc.col.wear If external wear detection outcome is stored as part of the data then this can be used by GGIR. This argument specifies the column in which the wear detection (Boolean) is stored.
rmc.doresample Boolean to indicate whether to resample the data based on the available timestamps and extracted sample rate from the file header
rmc.scalefactor.acc
Numeric value (default 1) to scale the acceleration signals via multiplication. For example, if data is provided in $\mathrm{m} / \mathrm{s} 2$ then by setting this to $1 / 9.81$ we would derive gravitational units.
interpolationType
Integer to indicate type of interpolation to be used when resampling time series (mainly relevant for Axivity sensors), $1=$ linear, $2=$ nearest neighbour.
PreviousLastValue
Automatically identified last value in previous chunk of data read.
PreviousLastTime
Automatically identified last timestamp in previous chunk of data read.
desiredtz Timezone in which device was worn.
configtz Timezone in which device was configured. If equal to desiredtz you can leave this in its default value.
header Header information that was extracted the previous time this file was read, to be re-used instead of being extracted again.

## Details

To use this function in the context of GGIR use all arguments from this function, except rmc.file, rmc.nrow, and rmc.skip as input for function GGIR or g.partl and also specify argument rmc.noise, which is not part of this function but needed to tell GGIR what noise level to expect in the data. The rmc.noise is taken from the params_rawdata object if not explicitly specified by user.

## Value

List with objects data holding the time series of acceleration, and header if it was available in the orignal file.

## Author(s)

Vincent T van Hees [v.vanhees@accelting.com](mailto:v.vanhees@accelting.com)

## Examples

```
# create test files: No header, with temperature, with time
N = 30
sf = 30
x = Sys.time()+((0:(N-1))/sf)
timestamps = as.POSIXlt(x, origin="1970-1-1", tz = "Europe/London")
mydata = data.frame(x = rnorm(N), time = timestamps, y = rnorm(N), z = rnorm(N),
            temp = rnorm(N) + 20)
testfile = "testcsv1.csv"
write.csv(mydata, file= testfile, row.names = FALSE)
loadedData = read.myacc.csv(rmc.file=testfile, rmc.nrow=20, rmc.dec=".",
    rmc.firstrow.acc = 1, rmc.firstrow.header=c(),
    desiredtz = "",
    rmc.col.acc = c(1,3,4), rmc.col.temp = 5, rmc.col.time=2,
    rmc.unit.acc = "g", rmc.unit.temp = "C", rmc.origin = "1970-01-01")
if (file.exists(testfile)) file.remove(testfile)
```


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