# Package 'GGIR'

# August 28, 2016

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# Description

GGIR is an R-package to process multi-day raw accelerometer data. It was developed in the context of research on human daily physical activity with wearable tri-axial acceleration sensors. The term raw accelerometry refers to data being expressed in m/s2 or gravitational acceleration as opposed to the previous generation accelerometers which stored only processed summary measures.

The package has been developed with and for the accelerometer brands Genea and Geneactive. Additionally, it should work for csv data from Geneactiv and Actigraph accelerometer data. Although, I have tested this less thoroughly compared with the binary data formats from Genea and Geneactiv.

Note for Actigraph users: please do not export timestamps to the csv-file as this causes memory issues. To cope with the absense of timestamps the code will re-caculate timestamps from the sample frequency and the start time and date as presented in the file header.

Function g.inspectfile assessess to which monitor brand the file belongs and extracts the file header; function g.calibrate helps to investigate calibration error based on free-living data and proposes correction factors; function g.getmeta extracts the signal features; g.impute takes that information, identifies unreliable signal sections (e.g. monitor not worn or signal clips near its extreme) and replaces these sections by imputed values; and finally g.analyse takes the output from all the functions, runs a basic descriptive analysis and then summarises the output both per measurement and per day of measurement.

To enhance the feasibility of using these individual functions I am providing a couple of shell functions to ease implementing the above functions in study data by less experienced R-users. Here, the main shell function is g.shell.GGIR and allows for automating the full analysis of a dataset including all necessary calls to the functions above. Function g.shell.GGIR relies on functions g.part1 and g.part2 also part of this package. In summary, the user is expected to specify the location of the acceleerometer data and the desired output folder. Next, data is loaded and pre- processed with g.getmeta and g.calibrate. Next, the output is converted to a conveniently portable .RData-format away from the R workspace. Next, these .RData files are used as input for g.part2.

Note that g.part1 generates a folder structure to help the user keep track of various output files and milestone data. The folder structure entails: One master folder with a name output\_xx where xx is equal to the name of the original data folder. Inside the output\_xx folder there will be one folder named meta including all the milestone and a folder results with all the results. Inside the

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meta folder the following subfolders are created: basic, ms2.out, ms3out, and ms4out for respectively g.part1, g.part2, g.part3, and g.part4 milestone data.

The reason why g.part1 and g.part2 are not merged as one generic shell function is because g.part1 takes much longer to run 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. So, function g.shell.GGIR basically is the central point for operating both g.part1 and g.part2 and most users should not really need to interact with g.part1 or g.part2 directly. More recently I expanded the package with g.part3 and g.part4 which provide functionality for estimating sleep and sustained inactivity bouts.

If you want to use this package for a different data format (e.g. from a different accelerometer brand) then please provide me with: the R-code to read the data and example files for testing purposes.

Please note that there is google discussion group for this package (link below).

You can thank me 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 in your own scientific journal/conference publications.

#### **Details**

 Package:
 GGIR

 Type:
 Package

 Version:
 1.2-10

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 2016-8-28

 License:
 LGPL (>= 2.0, < 3)</td>

Discussion group: https://groups.google.com/forum/#!forum/rpackageggir

### Author(s)

- Vincent T van Hees <vincentvanhees@gmail.com>
- Zhou Fang co-developed function g.calibrate
- Jing Hua Zhao <jinghua.zhao@mrc-epid.cam.ac.uk> co-developed function g.binread
- Severine Sabia tested and provided feedback on various functions

### 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

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 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)
#plot data
g.plot(IMP, M = data.getmeta, I = data.inspectfile, durplot=4)
```

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

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# **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)
```

g.analyse	function to analysse meta-data generated by g.getmeta and g.impute

### **Description**

Analyses the output from other functions within the packages to generate a basic descriptive summary for each accelerometer data file. Analyses include: Average acceleration per day, per measurement, L5M5 analyses (assessment of the five hours with lowest acceleration and with highest acceleration). Further, the traditionally popular variable MVPA is automatically extracted in six variants: without bout criteria in combination with epoch = epoch length as defined in g.getmeta (first value of the input argument windowsizes), 1 minute, and 5 minutes, and for bout durations 1 minute, 5 minutes or 10 minutes in combination with the epoch length as defined in g.getmeta.

### Usage

```
g.analyse(I, C, M, IMP, qlevels = c(), qwindow = c(0, 24), quantiletype = 7, L5M5window = c(0, 24), M5L5res = 10, includedaycrit = 16, ilevels = c(), winhr = 5, idloc = 1,snloc=1,mvpathreshold = c(),boutcriter=c(),mvpadur=c(1,5,10), selectdaysfile=c(),window.summary.size=10,dayborder=0,bout.metric = 1, closedbout=FALSE,desiredtz = c())
```

### **Arguments**

I	the output from function g.inspectfile
С	the output from function g.calibrate
М	the output from function g.getmeta
IMP	the output from function g.impute
qlevels	array 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, see g.getmeta.
qwindow	start and end time, in 24 hour clock hours, over which distribution in metric values need to be extracted. Default value = $c(0,24)$ will consider all 24 hours.
quantiletype	type of quantile function to use (default recommended). For details, see quantile function in STATS package
L5M5window	start and end time, in 24 hour clock hours, over which L5M5 needs to be calculated.
M5L5res	resoltion of L5 and M5 analysis in minutes (default: 10 minutes)
includedaycrit	minimum required number of valid hours in day specific analysis (NOTE: there is no minimum required number of hours per day in the summary of an entire measurement, every available hour is used to make the best possible inference on average metric value per average day)
ilevels	Levels for acceleration value frequency distribution in mg, e.g. $c(0,100,200)$ There is no constriction to the number of levels.

winhr window size in hours of L5 and M5 analysis (dedault = 5 hours)

idloc If value = 1 (default) the code assumes that ID number is stored in the obvi-

ous header field. If value = 2 the code uses the character string preceding the

character' in the filename as the ID number

snloc If value = 1 (default) the code assumes that device serial number is stored in the

obvious header field. If value = 2 the code uses the character string between the

first and second character '\_' in the filename as the serial number

mvpathreshold Threshold for MVPA estimation. This can be a single number or an array of

numbers, e.g. c(100,120). In the later case the code will estimate MVPA seperately for each threshold. If this variable is left blank c() then MVPA is not

estimated

boutcriter The variable boutcriter is a number between 0 and 1 and defines what fraction

of a bout needs to be above the mvpathreshold

mypadur default = c(1,5,10). Three bout duration for which MVPA will be calculated

selectdaysfile Functionality designed for the London Centre of Longidutinal studies. Csv file

holding the relation between device serial numbers and measurement days of

interest.

dayborder Hour at which days start and end (default = 0), value = 4 would mean 4am

window.summary.size

Functionality designed for the London Centre of Longidutinal studies. Size in

minutes of the summary window

bout.metric This argument used to be called mypa.2014 and had TRUE or FALSE as its

value. However, it has now become clear that this aspect of the analyses is still very much open for debate. Therefore, I have changed it into an argument where you can specify a metric for bout detection based on a number. A description of

these bout metrics can be found in the new function g.getbout

closedbout If TRUE then count breaks in a bout towards the bout duration. If FALSE then

only count time spent above the threshold towards the bout duration.

desiredtz see g.getmeta

### **Details**

The value summary is a dataframe and comes with the following variables:

- ID Participant id extracted from file header
- device\_sn Device serial number extracted from file header
- dodylocation Body location extracted from file header
- filename Name of the accelerometer file
- start\_time Timestamp when experiment started
- startday Name of day when experiment started
- samplefreqSample frequency (Hz)
- device Name of the device brand, e.g. Geneactiv

• clipping\_score Fraction of 15 minute windows per file for which the acceleration in one of the three axis was close to the maximum for at least 80 percent of the time. This should be 0

- meas\_dur\_dys Measurement duration (days)
- complete\_24hrcycle Fraction of 15 minute windows per 24 hours for which valid data is available at any day of the measurement
- meas\_dur\_def\_proto\_day Measurement duration (days) minus the hours that are ignored at the beginning and end of the measurement motived by protocol design
- wear\_dur\_def\_proto\_day Measurement duration according to protocol (days) minus invalid time periods
- calib\_err Estimated based on all non-movement periods in the measurement after applying the autocalibration
- calib\_status Summary statement about the status of the calibration error minimisation
- ENMO ENMO is the main summary measure of acceleration. The value presented is the average ENMO over all the available data normalised per 24 hour cycles, with invalid data imputed by the average at similar timepoints on different days of the week. In addition to ENMO it is possible to extract other acceleration metrics (i.e. BFEN, HFEN, HFENplus)
- pX\_ENMO\_mg\_0-24h This variable represents the Xth percentile in the distribution of short epoch acceleration values of the average day within the time interval as specified.
- L5hr\_ENMO\_mg\_0-24 Starting time in hours of the least active five\* hours within the time interval as specified (\* window size is modifiable in g.getmeta)
- L5\_ENMO\_mg Average acceleration over L5
- M5hr\_ENMO\_mg\_0-24 Starting time in hours of the most active five\* hours in the day within the time interval as specified (\* window size is modifiable in g.getmeta)
- M5\_ENMO\_mg\_0-24 Average acceleration over M5
- Accelerationa 1am-6am value of ENMO (mg) Average acceleration between 1am and 6am
- N valid WEdays Number of valid weekend days
- N valid WDdays Number of valid week days
- AD\_... The variable ... was calculated per day and then averaged over all the available days
- WE\_... The variable ... was calculated per day and then averaged over weekend days only
- WD\_... The variable ... was calculated per day and then averaged over week days only
- WWE\_... The variable ... was calculated per day and then averaged over weekend days. Double weekend days are averaged This is only relevant for experiments that last for more than seven days
- WWD\_... The variable ... was calculated per day and then averaged over week days. Double weekend days were averaged. This is only relevant for experiments that last for more than seven days)
- ...\_MVPA\_E5S\_B1M80\_T100 MVPA calculated based on 5 second epoch setting bout duration
   1 Minute and inclusion criterion of more than 80 percent. This is only done for metric ENMO at the moment, and only if mvpathreshold is not left blank
- ...\_mean\_ENMO... ENMO or other metric was first calcualte per day and then average according to AD, WD, WWE, WWD

• data exclusion strategy A log of the decision made when calling g.impute: value=1 mean ignore specific hours; value=2 mean ignore all data before the first midnight and after the last midnight

- n hours ignored at the start of the measurement (if strategy = 1) A log of the decision made when calling g.impute
- n hours ignored at the end of the measurement (if strategy = 1) A log of the decision made when calling g.impute
- n days of measurement after which data is ignored (if strategy = 1) A log of the decision made when calling g.impute

The value daysummary is a dataframe and comes with the following variables:

- ID Participant id extracted from file header
- filename File name
- calender\_date Calender data
- bodylocation Body location (if known)
- N valid hours Number of hours with valid data
- N hours Number of hours of measurement
- Day of the weekDay of the week
- Day of measurement Day number relative to start of the measurement
- L5\_ENMO\_mg\_0-24h Magnitude of average acceleration during the least active five hours calculated with metric ENMO. Within the time window as specified
- L5hr\_ENMO\_mg\_0-24h Starting hour of L5 on a scale from 0 to 24, where 14.5 means 14:30. Within the time window as specified
- M5\_ENMO\_mg\_0-24h Magnitude of average acceleration during the most active five hours calculated with metric ENMO. Within the time window as specified
- M5hr\_ENMO\_mg\_0-24h Starting hour of M5 on a scale from 0 to 24, where 14.5 means 14:30. Within the time window as specified
- mean\_ENMO\_mg\_1-6am Mean acceleration between 1am and 6am
- mean\_ENMO\_mg\_24hr Mean acceleration over 24 hour period
- pX\_ENMO\_mg\_0-24h Percentile in the short epoch distribution with invalid data imputed. Within the time window as specified
- [A,B)\_ENMO\_mg\_0-24h Time spent in minutes between (and including) acceleration value A in mg and (excluding) acceleration value B in mg. This is only done for metric ENMO at the moment, and only done if ilevels is not left blank
- MVPA\_E5S\_B1M80\_T100 MVPA calculated based on 5 second epoch setting bout duration 1
   Minute and inclusion criterion of more than 80 percent. This is only done for metric ENMO
   at the moment, and only if mvpathreshold is not left blank

# Value

summary summary for the file that was analysed (see details)

daysummary summary per day for the file that was analysed (see details)

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### Author(s)

Vincent T van Hees <vincentvanhees@gmail.com>

### **Examples**

```
data(data.getmeta)
 data(data.inspectfile)
 data(data.calibrate)
 ## Not run:
   #inspect file:
   I = g.inspectfile(datafile)
   #autocalibration:
   C = g.calibrate(datafile)
   #get meta-data:
   M = g.getmeta(datafile, desiredtz = "Europe/London", windowsizes = c(5, 900, 3600),
   daylimit = FALSE, offset = c(0, 0, 0), scale = c(1, 1, 1), tempoffset = c(0, 0, 0))
 ## End(Not run)
 #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)
g.binread
                         function to read binary files as produced by the accelerometer named
                          'Genea', not to be confused with the 'GENEActiv' (see package GEN-
                         EAread for this)
```

# **Description**

For reading the binary data as collected with a Genea accelerometer (Unilever Discover, UK). For reading GENEActive binary data, see package GENEAread.

# Usage

```
g.binread(binfile, start = 0, end = 0)
```

### **Arguments**

binfile	filename (required)
start	start point for reading data, this can either be a timestamp "year-month-day hr:min:sec" or a page number (optional)
end	end point for reading data, this can either be a timestamp "year-month-day hr:min:sec" or a page number (optional)

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### **Details**

If only start is defined then g.binread will read all data beyond start until the end of the file is reached

### Value

rawxyz matrix with raw x, y, and, z acceleration values

header file header

timestamps 1 timestamps for rawxyz in seconds since 1970-01-01 00:00

timestamps for rawxyz in day time format

batt.voltage matrix with battery voltage and corresponding timestamps

### Author(s)

Vincent T van Hees <vincentvanhees@gmail.com> Jing Hua Zhao <jinghua.zhao@mrc-epid.cam.ac.uk>

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, use.temp = TRUE, spherecrit = 0.3, minloadcrit = 72,
printsummary = TRUE, chunksize=c(), windowsizes=c(5,900,3600), selectdaysfile=c(),
dayborder=0)
```

# **Arguments**

datafile	name of accelerometer file
use.temp	use temperature sensor data if available (Geneactive only)
spherecrit	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	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	if TRUE will print a summary when done

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chunksize number between 0.2 and 1 to specificy the size of chunks to be loaded as a

fraction of a 12 hour period, e.g. 0.5 equals 6 hour chunks. The default is 1 (12 hrs). For machines with less than 4Gb of RAM memory a value below 1 is

recommended.

windowsizes see g.getmeta selectdaysfile see g.part1 dayborder see g.part1

### Value

scale scaling correction values, e.g. c(1,1,1) offset offset correction values, e.g. c(0,0,0)

tempoffset correction values related to temperature, e.g. c(0,0,0)

cal.error.start

absolute difference between Euclidean norm during all non-movement windows

and 1 g before autocalibration

cal.error.end absolute difference between Euclidean norm during all non-movement windows

and 1 g after autocalibration

spheredata average, standard deviation, Euclidean norm and temperature (if available) for

all ten second non-movement windows as used for the autocalibration procedure

npoints number of 10 second no-movement windows used to populate the sphere

nhoursused number of hours of measurement data scanned to find the ten second time win-

dows with no movement

mean temperature corresponding to the data as used for autocalibration. Only

applies to data collected with GENEActiv monitor.

### Author(s)

Vincent T van Hees <vincentvanhees@gmail.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)
```

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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,closedbout=FALSE,bout.metric=1,ws3=5)

### **Arguments**

x vector of zeros and/or ones to be screened for bouts of ones

boutduration duration of bout in epochs

boutcriter Minimum percentage of boutduration for which the epoch values are expected

to meet the threshold criterium

closedbout TRUE if you want breaks in bouts to be counted towards time spent in bouts

(argument only active for bout.metric 1 and 2)

bout.metric If value=1 the code uses the MVPA bout definition as has been available since

2014 (see papers by Sabia AJE 2014 and da Silva IJE 2014). Here, the algorithm looks for 10 minute windows in which more than XX percent of the epochs are above mvpathreshold, and then counts the entire window as mvpa. If value=2 the code looks for a group or groups of epochs with a value above mvpathreshold that span a time window of at least mvpadur minutes in which more than boutcriter percent of the epochs are above the threshold. The motivation for the defition 1 was: A person who spends 10 minutes in MVPA with a 2 minute break in the middle is equally active as a person who spends 8 minutes in MVPA without taking a break. Therefore, both should be counted equal and counted as 10 minute MVPA bout. The motivation for the definition 2 is: not counting breaks towards MVPA simplifies interpretation and still counts the two persons in the example as each others equal. If value=3, using sliding window across the data to test bout criteria per window and do not allow for breaks of 1 minute or

longer.

epoch length in seconds, only needed for bout.metric =3, because it needs to

measure how many epochs equal 1 minute breaks

#### Value

ws3

x Vector with binary numbers indicator where bouts where detected

boutcount Vector with binary numbers indicator where bouts where detected and counted

towards time spent in bouts, see argument closedbout for clarification

### Author(s)

Vincent T van Hees <vincentvanhees@gmail.com>

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### **Examples**

```
y = g.getbout(x=round(runif(1000,0.4,1)),boutduration = 120,boutcriter=0.9,
    closedbout=FALSE,bout.metric=3,ws3=5)
```

g.getmeta

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. Further, the function should be able to cope with csv-format data procuded by GENEActiv and Actigraph

### Usage

```
g.getmeta(datafile, desiredtz = c(),
windowsizes = c(5, 900, 3600),daylimit = FALSE,
offset = c(0,0,0), scale = c(1,1,1),
tempoffset = c(0,0,0),do.bfen = FALSE, do.enmo = TRUE,
do.lfenmo = FALSE, do.en = FALSE,
do.hfen = FALSE, do.hfenplus = FALSE,
do.teLindert2013=FALSE,do.anglex=FALSE,do.angley=FALSE,do.anglez=FALSE,
do.roll_med_acc_x=FALSE,do.roll_med_acc_y=FALSE,do.roll_med_acc_z=FALSE,
do.dev_roll_med_acc_x=FALSE,do.dev_roll_med_acc_y=FALSE,do.dev_roll_med_acc_z=FALSE,
do.enmoa = FALSE,lb = 0.2,hb = 15, n = 4,meantempcal=c(),chunksize=c(),
selectdaysfile=c(),dayborder=0,...)
```

### **Arguments**

= 1/scale

datafile name of accelerometer file desiredtz desired timezone: see also http://en.wikipedia.org/wiki/Zone.tab windowsizes Three values to indicate the lengths of the windows as in c(window1, window2, window3): window1 is the short epoch length in seconds and by default 5 this is the time window over which acceleration and angle metrics are calculated, window 2 is the long epoch length in seconds for which non-wear and signal clipping are defined, default 900. However, 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. daylimit number of days to limit (roughly), if set to FALSE no daylimit will be applied offset offset correction value per axis, usage: value = scale(value,center = -offset, scale

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scale	scaling correction value per axis, usage: value = scale(value,center = -offset, scale = 1/scale)
tempoffset	temperature offset correction value per axis, usage: value = scale(value,center = -offset, scale = 1/scale) + scale(temperature, center = rep(averagetemperate,3), scale = 1/tempoffset)
do.bfen	if TRUE, calculate metric BFEN with band-pass filter configuration set by 1b and hb
do.enmo	if TRUE (default), calculate metric ENMO with negative values rounded to zero
do.lfenmo	if TRUE, calculate metric LFENMO with low-pass filter configuration set by hb
do.en	if TRUE, calculate metric EN
do.hfen	if TRUE, calculate metric HFEN with low-pass filter configuration set by hb
do.hfenplus	if TRUE, calculate metric HFENplus with band-pass filter configuration set by 1b and hb
do.teLindert20	
	if TRUE, calculate the 5 sec epoch Actiwatch count replication as described by te Lindert et al 2013 in the journal SLEEP (volume 36, issue 5, page 781)
do.anglex	if TRUE, calculate the angle of the x-axis relative to the horizontal plane (degrees) utilizing all three axes
do.angley	if TRUE, calculate the angle of the y-axis relative to the horizontal plane (degrees) utilizing all three axes
do.anglez	if TRUE, calculate the angle of the z-axis relative to the horizontal plane (degrees) utilizing all three axes
do.enmoa	if TRUE (default), calculate metric ENMOa which is equal to metric ENMO but with the absolute taken from the Euclidean norm minus one.
do.roll_med_ac	
d	if TRUE, calculate rolling median for the x axis
do.roll_med_ac	if TRUE, calculate rolling median for the y axis
do.roll_med_ac	·
	if TRUE, calculate rolling median for the z axis
do.dev_roll_me	d_acc_x if TRUE, calculate deviations from rolling median for the x axis
do.dev_roll_me	d_acc_y if TRUE, calculate deviations from rolling median for the y axis
do.dev_roll_me	d_acc_z if TRUE, calculate deviations from rolling median for the z axis
lb	lower boundary of the frequency filter (in Hertz)
hb	upper boundary of the frequency filter (in Hertz)
n	order of the frequency filter
meantempcal	mean temperature corresponding to the data as used for autocalibration. If autocalibration is not done or if temperature was not available then leave blank (default)

16 g.getmeta

chunksize number between 0.2 and 1 to specificy the size of chunks to be loaded as a

fraction of a 24 hour period, e.g. 0.5 equals 12 hour chunks. The default is 1 (24 hrs). For machines with less than 4Gb of RAM memory a value below 1 is

recommended.

selectdaysfile see g.part1 dayborder see g.part1

Please ignore. Only used by the code internally when called from within g.part1

with selectdaysfile specific.

#### Value

metalong dataframe with long epoch meta-data: EN, non-wear score, clipping score, tem-

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 <vincentvanhees@gmail.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

# **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)
```

g.impute 17

g.impute	function to identify invalid periods in the meta-data as generated by g.getmeta and to impute these invalid periods with the average of sim-
	ilar timepoints on other days of the measurement

# Description

Functions takes the output from g.getmeta and information about the study protocol to label impute invalid time segments in the data.

# Usage

```
g.impute(M, I, strategy = 1, hrs.del.start = 0, hrs.del.end = 0, maxdur = 0,
ndayswindow = 7,desiredtz="Europe/London")
```

# Arguments

М	output from g.getmeta
I	output from g.inspectfile
strategy	how to deal with knowledge about study protocol. value = 1 means select data based on hrs.del.start, hrs.del.end, and maxdur. Value = 2 makes that only the data between the first midnight and the last midnight is used for imputation. Value = 3 only selects the most active $X$ days in the files. $X$ is specified by argument ndayswindow
hrs.del.start	how many HOURS after start of experiment did wearing of monitor start?
hrs.del.end	how many HOURS before the end of the experiment did wearing of monitor definitely end?
maxdur	How many DAYS after start of experiment did experiment definitely stop? (set to zero if unknown = default)
ndayswindow	If strategy is set to 3 then this is the size of the window as a number of days
desiredtz	see g.getmeta

# Value

metashort	imputed short epoch variables
rout	matrix to clarify when data was imputed for each long epoch time window and the reason for imputation. Value = 1 indicates imputation. Columns $1 = \text{monitor}$ non wear, column $2 = \text{clipping}$ , column $3 = \text{additional nonwear}$ , column $4 = \text{protocol based exclusion}$ and column $5 = \text{sum of column } 1,2,3$ and $4$ .
averageday	matrix with n columns for n metrics values and m rows for m short epoch time windows in an average 24 hours period

# Author(s)

Vincent T van Hees <vincentvanhees@gmail.com>

18 g.inspectfile

# **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)

#impute meta-data:
IMP = g.impute(M=data.getmeta, I=data.inspectfile)
```

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)
```

# Arguments

datafile name of data file

# Value

header fileheader

monn monitor name (genea, geneactive)

monc monitor brand code (1 = genea; 2 = geneactive, 3 = actigraph)

dformn data format (bin, csv)

dformc data format code (1 = bin, 2 = csv)

sf samplefrequency in Hertz

filename filename

### Author(s)

Vincent T van Hees <vincentvanhees@gmail.com>

g.part1 function to load and pre-process acceleration files

### **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.part1 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 g.shell.GGIR provides the main shell that allows for operating g.part1 and g.part2.

### Usage

# **Arguments**

datadir	Directory where the accelerometer files are stored or list of accelerometer file- names and directories
outputdir	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 organise output
f0	File index to start with (default = $1$ ). Index refers to the filenames sorted in increasing order
f1	File index to finish with (defaults to number of files available)
windowsizes	see g.getmeta
desiredtz	see g.getmeta

chunksize	see g.getmeta
studyname	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 will be used as name for the analysis
do.bfen	if TRUE, calculate metric BFEN with band-pass filter configuration set by 1b and hb, see $g.getmeta$
do.enmo	if TRUE (default), calculate metric ENMO, see g.getmeta
do.lfenmo	if TRUE, calculate metric LFENMO with low-pass filter configuration set by hb,see g.getmeta
do.en	if TRUE, calculate metric EN, see g.getmeta
do.hfen	if TRUE, calculate metric HFEN with low-pass filter configuration set by hb, see g.getmeta
do.hfenplus	if TRUE, calculate metric HFENplus with band-pass filter configuration set by 1b and hb, see g.getmeta
do.teLindert20	13
	if TRUE, calculate the 5 sec epoch Actiwatch count replication as described by te Lindert et al 2013 in the journal SLEEP (volume 36, issue 5, page 781)
do.anglex	if TRUE, calculate the angle of the x-axis relative to the horizontal plane (degrees) utilizing all three axes
do.angley	if TRUE, calculate the angle of the y-axis relative to the horizontal plane (degrees) utilizing all three axes
do.anglez	if TRUE, calculate the angle of the z-axis relative to the horizontal plane (degrees) utilizing all three axes
do.enmoa	if TRUE (default), calculate metric ENMOa which is equal to metric ENMO but with the absolute taken from the Euclidean norm minus one.
do.roll_med_aco	
d	see g.getmeta
do.roll_med_aco	see g.getmeta
do.roll_med_acc	
uo,, ollou_uo,	see g.getmeta
do.dev_roll_med	
	see g.getmeta
do.dev_roll_med	
d. d	see g.getmeta
do.dev_roll_med	see g.getmeta
do.cal	Whether to apply auto-calibration or not, see g.calibrate. Default and recom-
	mended setting is TRUE
1b	lower boundary of the frequency filter (in Hertz)
hb	upper boundary of the frequency filter (in Hertz), see g.getmeta
n	order of the frequency filter, see g.getmeta

use.temp see g.calibrate use temperature sensor data if available (Geneactive only)

spherecrit see g.calibrate 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 see g.calibrate 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 be reduced to under 0.01 g.

see g.calibrate if TRUE will print a summary when done printsummary

Whether to print the filename before before analysing it (default is FALSE). print.filename

Printing the filename can be useful to investigate problems (e.g. to verify that

which file is being read).

overwrite Overwrite previously generated milestone data by this function for this particu-

lar dataset. If FALSE then it will skip the previously processed files (default =

FALSE).

backup.cal.coef

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 csy-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. The argument is intended for analysing short lasting laboratory experiments with insufficient sphere data, but for which calibration coefficients can be derived in an alternative way. It is the users responsibility to compile the csv-spreadsheet.

selectdaysfile Optional functionality. Character pointing at a csv file holding the relationship between device serial numbers (first column) and measurement dates of interest (second and third column). The date format should be dd/mm/yyyy. And the first row if the csv file is assumed to have a character variable names, e.g. "serialnumber" "Day1" and "Day2" respectively. Raw data will be extracted and stored in the output directory in a new subfolder named 'raw'.

dayborder

Hour at which days start and end (default = 0), value = 4 would mean 4am

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

g.part2

function to analyse and summarize pre-processed output from g.part1

# **Description**

Loads the output from g.part1 and then applies g.impute and g.analyse, after which the output is converted to .RData-format which will be used by g.shell.GGIR to generate reports. The variables in these reports are the same variables as described in g.analyse.

## Usage

```
g.part2(datadir=c(),metadatadir=c(),f0=c(),f1=c(),strategy = 1, hrs.del.start = 0.5, hrs.del.end = 0.5, maxdur = 7, includedaycrit = 16, L5M5window = c(0,24), M5L5res = 10, winhr = 5, qwindow=c(0,24), qlevels = c(0.1), ilevels = c(0,10), mvpathreshold = c(100),boutcriter = 0.8,ndayswindow=7,idloc=1, do.imp=TRUE,storefolderstructure=FALSE,overwrite=FALSE,epochvalues2csv=FALSE, mvpadur=c(1,5,10),selectdaysfile=c(),window.summary.size=10,dayborder=0, bout.metric=2,closedbout=FALSE,desiredtz="Europe/London")
```

#### **Arguments**

datadir	Directory where the accelerometer files are stored or list, e.g. "C:/mydata" of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin","C:/mydata/myfile2.bin").
metadatadir	Directory where the output from g.part1 was stored
f0	File index to start with (default = 1). Index refers to the filenames sorted in increasing order
f1	File index to finish with (defaults to number of files available)

how to deal with knowledge about study protocol. value = 1 to select data based strategy on hrs.del.start, hrs.del.end, and maxdur. Value = 2 to only use the data between the first midnight and the last midnight, value = 3 only selects the most active X days in the files. X is specified by argument ndayswindow See also g.impute hrs.del.start how many HOURS after start of experiment did wearing of monitor start?, see g.impute hrs.del.end how many HOURS before the end of the experiment did wearing of monitor definitely end?, see g.impute maxdur how many DAYS after start of experiment did experiment definitely stop? (set to zero if unknown = default), see g.impute includedaycrit minimum required number of valid hours in day specific analysis (NOTE: there is no minimum required number of hours per day in the summary of an entire measurement, every available hour is used to make the best possible inference on average metric value per week) L5M5window start and end time, in 24 hour clock hours, over which L5M5 needs to be calculated. The calculation is done based on the average day M5L5res resoltion of L5 and M5 analysis in minutes (default: 10 minutes) winhr window size in hours of L5 and M5 analysis (dedault = 5 hours) qwindow start and end time, in 24 hour clock hours, over which distribution in metric values need to be extracted. Value = c(0.24) will consider all 24 hours. glevels array 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, see g.getmeta. ilevels Levels for acceleration value frequency distribution in mg, e.g. c(0,100,200) There is no constriction to the number of levels. Threshold for MVPA estimation. Threshold needs to be based on metric ENMO. mvpathreshold This can be a single number or an array of numbers, e.g. c(100,120). In the later case the code will estimate MVPA seperately for each threshold. If this variable is left blank c() then MVPA is not estimated boutcriter The variable boutcriter is a number between 0 and 1 and defines what fraction of a bout needs to be above the mypathreshold If strategy is set to 3 then this is the size of the window as a number of days ndayswindow idloc If value = 1 (default) the code assumes that ID number is stored in the obvious header field. If value = 2 the code uses the character string preceding the character' in the filename as the ID number do.imp Whether to impute missing values (e.g. suspected of monitor non-wear) or not by g.impute. Default and recommended setting is TRUE

storefolderstructure

overwrite

Store folder structure of the accelerometer data

Overwrite previously generated milestone data by this function for this particular dataset. If FALSE then it will skip the previously processed files (default =

FALSE).

epochvalues2csv

If TRUE then epoch values are exported to a CSV spreadsheet. Here, non-wear

time is imputed where possible (default = FALSE).

mvpadur default = c(1,5,10). Three bout duration for which MVPA will be calculated

selectdaysfile Functionality designed for the London Centre of Longidutinal studies. Csv file

holding the relation between device serial numbers and measurement days of

interest.

dayborder Hour at which days start and end (default = 0), value = 4 would mean 4am

window.summary.size

Functionality designed for the London Centre of Longidutinal studies. Size in

minutes of the summary window

bout.metric This argument used to be called mvpa.2014 and had TRUE or FALSE as its

value. However, it has now become clear that this aspect of the analyses is still very much open for debate. Therefore, I have changed it into an argument where you can specify a metric for bout detection based on a number. A description of

these bout metrics can be found in the new function g.getbout

closedbout See g.getbout desiredtz see g.getmeta

### 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 <vincentvanhees@gmail.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.part3	Detection of sustained inactivity periods as needed for sleep detection in g.part4.

# **Description**

Function called by g.shell.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 g.shell.GGIR only.

### Usage

```
g.part3(metadatadir=c(),f0,f1,anglethreshold = 5,timethreshold = 5,
ignorenonwear=FALSE,overwrite=FALSE,desiredtz="Europe/London")
```

# 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 g.shell.GGIR will recognise what the value of metadatadir is.
f0	File index to start with (default $= 1$ ). Index refers to the filenames sorted in increasing order
f1	File index to finish with (defaults to number of files available)
anglethreshold	Angle threshold (degrees) for sustained inactivity periods detection, default = 5
timethreshold	Time threshold (minutes) for sustained inactivity periods detection, default = 5. This can be specified as multiple thresholds, each of which will be implemented. For example, timethreshold = $c(5,10)$
ignorenonwear	If TRUEE then ignore detected monitor non-wear periods to avoid confusion between monitor non-wear time and sustained inactivity (default = TRUE)
overwrite	Overwrite previously generated milestone data by this function for this particular dataset? If FALSE then it will skip the previously processed files (default = FALSE).
desiredtz	See g.getmeta

# 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 <vincentvanhees@gmail.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

### **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.part3 as either nocturnal sleep or daytime sustained inactivity

### **Description**

Loads output from g.part3 as stored in milestone data and sleep log information (if available) and then uses these information sources to define nocturnal sleep and daytime sustained inactivity.

### Usage

### **Arguments**

datadir Directory where the accelerometer files are stored or list of accelerometer file-

names and directories

metadatadir Directory that holds a folders 'meta' and inside this a folder 'basic' which con-

tains the milestone data produced by g.part1. The folderstructure is normally created by g.part3. When using g.part4 via g.shell.GGIR then g.shell.GGIR will automatically recognise what the value of metadatadir is, so the user does not

need to specify this.

f0 File index to start with (default = 1). Index refers to the filenames sorted in

increasing order

File index to finish with (defaults to number of files available)

idloc If value = 1 (default) the code assumes that ID number is stored in the obvi-

ous header field. If value = 2 the code uses the character string preceding the

character '\_' in the filename as the ID number

loglocation Location of the spreadsheet (csv) with sleep log information. The spreadsheet

needs to have the following structure: one column for participant id, and then followed by alternatingly one column for onset time and one column for waking time. There can be multiple sleeplogs in the same spreadsheet. The first raw of the spreadsheet needs to be filled with column names, it does not matter what these column names are. Timestamps are to be stored without date as in hh:mm:ss. If onset corresponds to lights out or intention to fall asleep, then it is the end-users responsibility to account for this in the interpretation of the results.

colid Column number in the sleep log spreadsheet in which the participant ID code is

stored (default = 1)

coln1 Column number in the sleep log spreadsheet where the onset of the first night

starts

nnights Number of nights for which sleep log information should be available. It as-

sumes that this is constant within a study. If sleep log information is missing for

certain nights then leave these blank

sleeplogidnum Should the participant identifier as stored in the sleeplog be interpretted as a

number (TRUE=default) or a character (FALSE)?

do.visual If g.part4 is run with do.visual == TRUE then 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 Relevant for do.visual == TRUE. Outliers.only == FALSE will visualise all

available nights in the data. Outliers.only == TRUE will visualise only for nights with a difference in onset or waking time larger than the variable of argument

criterror.

excludefirstlast

If TRUE then the first and last night of the measurement are ignored for the sleep

assessment.

criterror Relevant for do.visual == TRUE and outliers.only == TRUE. criterror speci-

fies the number of minimum number of hours difference between sleep log and accelerometer estimate for the night to be included in the visualisation

includenightcrit

Minimum number of valid hours per night (24 hour window between noon and noon)

relyonsleeplog If TRUE then sleep onset and waking time are defined based on timestamps derived from sleep log if FALSE (default) the sleep log is only used to guide the accelerometer-based detection. If participants were instructed NOT to wear the accelerometer during waking hours then set to TRUE, in all other scenarios set to FALSE (FALSE).

def.noc.sleep

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 def.noc.sleep is left blank then the 12 hour window centred at the least active 5 hours of the 24 hour period will be used instead.

storefolderstructure

Store folder structure of the accelerometer data

overwrite

Overwrite previously generated milestone data by this function for this particular dataset. If FALSE then it will skip the previously processed files (default = FALSE).

### Value

The function does not produce values but generates an RData file in the milestone subfolder ms4.out which incudes a dataframe named nightsummary and comes with the following variables:

- id Participant id extracted from file
- night Night number
- acc\_onset Detected onset of sleep expressed as hours since the previous midnight
- acc\_wake Detected waking time (after sleep period) expressed as hours since the previous midnight
- acc\_timeinbed Difference between onset and waking time.
- acc\_def Definition of sustained inactivity by accelerometer
- sleeplog\_onset Sleep onset derived from sleep log or specified by researcher if not sleep log is available
- sleeplog\_wake Waking time derived from sleep log or specified by researcher if not sleep log is available
- sleeplog\_timeinbed Time in bed derived from sleep log or specified by researcher if not sleep log is available
- error\_onset Difference between sleep onset as estimated by accelerometer and estimated by sleeplog/defined by researcher
- error\_wake Difference between waking time as estimated by accelerometer and estimated by sleeplog/defined by researcher
- fraction\_night\_invalid Fraction of the night for which the data was invalid, e.g. monitor not worn or no accelerometer measurement started/ended within the night

acc\_dur\_noc Total sleep duration, which equals the accumulated nocturnal sustained inactivity bouts. This is not the same as the time in bed, which only looks at time difference between falling asleep and waking up.

- acc\_dur\_sibd Accumulated sustained inactivty bouts during the day. These are the periods
  we would label during the night as sleep, but during the day they form a subclass of inactivity,
  which may represent day time sleep or wakefulness while being motionless for a sustained
  period of time
- acc\_n\_noc Number of noturnal sleep periods
- acc\_n\_sibd Number of sustained inactivity periods during the day
- acc\_onset\_ts acc\_onset formatted as a timestamp
- acc\_wake\_ts acc\_wake formatted as a timestamp
- sleeplog\_onset\_ts sleeplog\_onset formatted as a timestamp
- sleeplog\_wake\_ts sleeplog\_wake formatted as a timestamp
- page pdf page on which the visualisation can be found
- daysleeper If 0 then the person is a nightsleeper (sleep period did not overlap with noon) if value=1 then the person is a daysleeper (sleep period did overlap with noon)
- weekday Day of the week on which the night started
- calendardate Calendar data which the night started
- filename Name of the accelerometer file
- cleaningcode 0: no problem; 1: sleeplog not available, 2: not enough valid accelerometer data, 3: no accelerometer data available, 4: there were no nights to be analysed for this person
- sleeplog\_used Whether a sleep log was used (TRUE/FALSE)
- acc\_available Whether accelerometer data was available (TRUE/FALSE). This dataframe is used in g.report.part4 to create two reports one per night and one per person.

Note that function g.shell.GGIR comes with the option for report generation. In relation to function g.part4 it is important to mention that these reports are effectively the variable names mentioned above or derivatives. Please fine below extra clarification on a few of the variable names for which the meaning may not be obvious:

- sleeplog\_used Whether a sleeplog was available (TRUE) or not (FALSE)
- n\_nights\_acc Number of nights of accelerometer data
- n\_WE\_nights\_complete Number of weekend nights complete which means both accelerometer and sleeplog data
- n\_WD\_nights\_complete Number of weekday nights complete which means both accelerometer and sleeplog data
- n\_WEnights\_daysleeper Number of weekend nights on which the person slept until after noon
- n\_WDnights\_daysleeper Number of weekday nights on which the person slept until after noon
- sleeplog\_dur\_AD\_mn Mean sleep duration according to sleeplog accros all days

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 sleeplog\_dur\_AD\_sd Standard deviation of sleep duration according to sleeplog accros all days

- sleeplog\_dur\_WD\_sd Standard deviation of sleep duration according to sleeplog accros weekdays
- sleeplog\_dur\_WE\_sd Standard deviation of sleep duration according to sleeplog accros weekend days

### Author(s)

Vincent T van Hees <vincentvanhees@gmail.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, accepted for publication October 2015

# **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)
```

g.plot

function to generate a plot for quality check purposes

# **Description**

Function takes meta-data as generated by g.getmeta and g.impute to create a visual representation of imputed time periods

### **Usage**

```
g.plot(IMP, M, I, durplot)
```

## **Arguments**

IMP output from g.impute
 M output from g.getmeta
 I output from g.inspectfile
 durplot number of days to plot

# Value

function only produces a plot, no values

### Author(s)

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### **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)

#impute meta-data:
IMP = g.impute(M = data.getmeta, I = data.inspectfile, strategy = 1, hrs.del.start = 0, hrs.del.end = 0, maxdur = 0)

#plot data
g.plot(IMP, M = data.getmeta, I = data.inspectfile, durplot=4)
```

g.shell.GGIR

Shell function for analysing a accelerometer dataset.

### **Description**

This function is designed to help users operate all steps of the analysis. It helps to generate and structure milestone data, produces user-friendly reports. The function acts as a shell with calls to g.part1, g.part2, g.part3 and g.part4. Please see these specific functions for clarification on optional input arguments.

### Usage

```
g.shell.GGIR(mode=c(1,2),datadir=c(),outputdir=c(),studyname=c(),f0=1,f1=0,do.report=c(2),overwrite=FALSE,visualreport=FALSE,viewingwindow=1,...)
```

# **Arguments**

mode

Specify which of the four parts need to be run, e.g. mode = 1 makes that g.part1 is run. Default setting, mode = c(1,2), makes that both part1 and part2 are ran. Note that if mode = c(1,3) then the code will also set do.anglez = TRUE in order to enable sleep detection. If you run part 1 and 3 seperatedly then you need to remember to set argument do.anglez to TRUE when running part1.

datadir	Directory where the accelerometer files are stored or list, e.g. "C:/mydata" of accelerometer filenames and directories, e.g. c("C:/mydata/myfile1.bin","C:/mydata/myfile2.bin").
outputdir	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	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
f0	File index to start with (default = 1). Index refers to the filenames sorted in increasing order
f1	File index to finish with (defaults to number of files available)
overwrite	Do you want to overwrite analysis for which milestone data exists? If over-write=FALSE then milestone data from a previous analysis will be used if available and visual reports will not be created again.
do.report	For which parts to generate a summary spreadsheet: 2 and/or 4. Default is $c(2)$ . 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.
visualreport	If TRUE then generate visual report based on combined output from part 2 and 4. This is in beta-version at the moment.
viewingwindow	Centre the day as displayed around noon (value = 1) or around midnight (value = 2)
•••	Any input argument needed for functions g.part1, g.part2, g.part3 or g.part4. See respective function documentation for further clarification.

### Value

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

### Author(s)

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### 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)
datadir= "C:/myfolder/mydata"
outputdir= "C:/myresults"
studyname="test"
f0 = 1
f1 = 2
g.shell.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:
           #-----
           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 = TRUE,
includenightcrit = 16,
def.noc.sleep = c(),
loglocation= "D:/sleeplog.csv",
outliers.only = FALSE,
criterror = 4,
relyonsleeplog = FALSE,
sleeplogidnum = TRUE,
colid=1,
coln1=2,
do.visual = TRUE,
nnights = 9,
#-----
# Report generation
#-----
do.report=c(2,4))
```

## End(Not run)

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