

# Package ‘GGIR’

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**Type** Package

**Title** Raw Accelerometer Data Analysis

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**Description** A tool to process and analyse data collected with wearable raw acceleration sensors as described in Migueles and colleagues (JMPB 2019), and van Hees and colleagues (JApplPhysiol 2014; PLoS ONE 2015). The package has been developed and tested for binary data from 'GENEActiv' <<https://activinsights.com/>>, binary (.gt3x) and .csv-export data from 'Actigraph' <<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.

**URL** <https://github.com/wadpac/GGIR/>,  
<https://groups.google.com/forum/#!forum/RpackageGGIR>,  
<https://wadpac.github.io/GGIR/>

**BugReports** <https://github.com/wadpac/GGIR/issues>

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**Suggests** testthat, covr, knitr, rmarkdown, actlifecounts, readxl

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**VignetteBuilder** knitr

**NeedsCompilation** no

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GGIR-package	<i>A package to process multi-day raw accelerometer data</i>
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## Description

Disclaimer: If you are a new GGIR user then please see [package vignette](#) for an introduction to 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:	3.0-9
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Discussion group:	<a href="https://groups.google.com/forum/#!forum/rpackageggir">https://groups.google.com/forum/#!forum/rpackageggir</a>

## Author(s)

- Vincent T van Hees <[v.vanhees@accelting.com](mailto: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 <[jairoh@ugr.es](mailto:jairoh@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.2018-0063.
- 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 cosinorAnalyses

### 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>

---

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	Sample frequency in Hertz, the default here is low to minimize file size
Nmin	Number of minutes (minimum is 720)
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
start_time	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	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	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	<i>Example output from g.calibrate</i>
----------------	--

---

**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	<i>Example output from g.getmeta</i>
--------------	--------------------------------------

---

**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	<i>Example output from g.inspectfile</i>
------------------	--

---

**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	<i>Metalong object as part of part 1 milestone data</i>
---------------	---

---

**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 calibration 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 <a href="#">g.part1</a>
params_general	See <a href="#">g.part1</a>
params_cleaning	See <a href="#">g.part1</a>
inspectfileobject	Output from the function <a href="#">g.inspectfile</a> .
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 <code>suppressWarning()</code> or <code>suppressMessages()</code> .
...	Any argument used in the previous version of <code>g.calibrate</code> , which will now be used to overrule the arguments specified with the parameter objects.

**Value**

scale	scaling correction values, e.g. <code>c(1,1,1)</code>
offset	offset correction values, e.g. <code>c(0,0,0)</code>
tempoffset	correction values related to temperature, e.g. <code>c(0,0,0)</code>
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 windows with no movement
meantempcal	mean temperature corresponding to the data as used for autocalibration. Only applies to data where temperate data is collected and available to GGIR, such as GENEActiv, Axivity, and in some instances ad-hoc .csv data.

**Author(s)**

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

---

g.getmeta	<i>Function to extract meta-data (features) from data in accelerometer file</i>
-----------	---

---

### 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 GENEActiv 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

```
g.getmeta(datafile, params_metrics = c(), params_rawdata = c(),
          params_general = c(), params_cleaning = c(), daylimit = FALSE,
          offset = c(0, 0, 0), scale = c(1, 1, 1), tempoffset = c(0, 0, 0),
          meantempcal = c(), myfun = c(), inspectfileobject = c(),
          verbose = TRUE, ...)
```

### Arguments

datafile	name of accelerometer file
params_metrics	See details in <a href="#">GGIR</a> .
params_rawdata	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
params_cleaning	See details in <a href="#">GGIR</a> .
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 = 1/scale)
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(averagetemperature,3), scale = 1/tempoffset)
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)
myfun	External function object to be applied to raw data. See details <a href="#">applyExtFunction</a> .
inspectfileobject	Output from the function <a href="#">g.inspectfile</a> .

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

### Value

metalong	dataframe with long epoch meta-data: EN, non-wear score, clipping score, temperature
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$metashort[1:5,])

## End(Not run)
```

---

`g.imputeTimegaps`      *Impute gaps in three axis raw accelerometer data*

---

### 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 `_g_`

### Usage

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

### Arguments

<code>x</code>	Data.frame with raw accelerometer data, and a timestamp column with millisecond resolution.
<code>sf</code>	Sample frequency in Hertz
<code>k</code>	Minimum time gap length to be imputed
<code>impute</code>	Boolean to indicate whether the time gaps identified should be imputed
<code>PreviousLastValue</code>	Automatically identified last value in previous chunk of data read.
<code>PreviousLastTime</code>	Automatically identified last timestamp in previous chunk of data read.
<code>epochsize</code>	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	<i>function to inspect accelerometer file for brand, sample frequency and header</i>
---------------	--

---

**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 <a href="#">g.getmeta</a>
params_rawdata	See <a href="#">g.part1</a>
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 = .csv, 3 = .wav, 4 = .cwa, 5 = ad-hoc .csv, 6 = .gt3x)
sf	samplefrequency in Hertz
filename	filename

**Author(s)**

Vincent T van Hees <[v.vanhees@accltelling.com](mailto:v.vanhees@accltelling.com)>

---

`g.loadlog`*Load and clean sleeplog information*

---

### 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

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

### 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](mailto:v.vanhees@accelting.com)>

### Examples

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

## End(Not run)
```



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 [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 <a href="#">applyExtFunction</a> .
params_metrics	See details in <a href="#">GGIR</a> .
params_rawdata	See details in <a href="#">GGIR</a> .
params_cleaning	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	If you are working with a non-standard csv formatted files, <a href="#">g.part1</a> also takes any input arguments needed for function <a href="#">read.myacc.csv</a> and argument <code>rmc.noise</code> from <a href="#">get_nw_clip_block_params</a> . First test these argument with function <a href="#">read.myacc.csv</a> directly. To ensure compatibility with R scripts written for older GGIR versions, the user can also provide parameters listed in the <code>params_</code> 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.part1` is used via `GGIR` you have the option to specify a configuration file, which will overrule the default parameter values. Further, as user you can set parameter values as input argument to both `g.part1` 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)
```

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 [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(),
        myfun = c(), params_cleaning = c(), params_247 = c(),
        params_phyact = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> 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 <a href="#">applyExtFunction</a> .
params_cleaning	See details in <a href="#">GGIR</a> .
params_247	See details in <a href="#">GGIR</a> .
params_phyact	See details in <a href="#">GGIR</a> .
params_output	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	To ensure compatibility with R scripts written for older <a href="#">GGIR</a> versions, the user can also provide parameters listed in the <code>params_</code> 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 specify 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.part3

*Detection of sustained inactivity periods as needed for sleep detection 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> 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 <a href="#">applyExtFunction</a> .
params_sleep	See details in <a href="#">GGIR</a> .
params_metrics	See details in <a href="#">GGIR</a> .
params_output	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	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 specify 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 explanation).
- `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>

**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.part3 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 description of the sleep detection.

**Usage**

```
g.part4(datadir = c(), metadatadir = c(), f0 = f0, f1 = f1, 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> 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	List of parameters used for sleep analysis (GGIR part 3, 4, and 5): see documentation <a href="#">g.part3</a> .
params_metrics	List of parameters used for metrics extraction (GGIR part 1): see documentation <a href="#">g.part1</a> .
params_cleaning	See details in <a href="#">GGIR</a> .
params_output	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	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 includes 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](mailto: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)
```

---

g.part5 *Merge output from physical activity and sleep analysis into one report*

---

### 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

```
g.part5(datadir = c(), metadatadir = c(), f0 = c(), f1 = c(),
        params_sleep = c(), params_metrics = c(),
        params_247 = c(), params_phyact = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <b>GGIR</b> 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 <a href="#">GGIR</a> .
params_metrics	See details in <a href="#">GGIR</a> .
params_247	See details in <a href="#">GGIR</a> .
params_phyact	See details in <a href="#">GGIR</a> .
params_cleaning	See details in <a href="#">GGIR</a> .
params_output	See details in <a href="#">GGIR</a> .
params_general	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	To ensure compatibility with R scripts written for older <b>GGIR</b> 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 includes 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**

```
g.part5.analyseRest(sibreport = NULL, dsummary = NULL,
  ds_names = NULL, fi = NULL,
  di = NULL, time = NULL, tz = NULL,
  possible_nap_dur = 0,
  possible_nap_edge_acc = Inf)
```

**Arguments**

sibreport	sibreport data.frame produced by <a href="#">g.sibreport</a>
dsummary	matrix created internally by <a href="#">g.part5</a>
ds_names	character vector with variable names corresponding to dsummary created internally by <a href="#">g.part5</a>
fi	Numeric scalar to indicate variable index, created internally by <a href="#">g.part5</a>
di	Numeric scalar to indicate recording index, created internally by <a href="#">g.part5</a>
time	Daytime section of time column from the ts object, created internally by <a href="#">g.part5</a> ,
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**

```
g.part6(datadir = c(), metadatadir = c(), f0 = c(), f1 = c(),
        params_general = c(), params_phyact = c(), params_247 = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> 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_general	See details in <a href="#">GGIR</a> .
params_phyact	See details in <a href="#">GGIR</a> .
params_247	See details in <a href="#">GGIR</a> .
verbose	See details in <a href="#">GGIR</a> .
...	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 includes ... (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(metatadir=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(metatadir = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> will recognise what the value of metadatadir is.
dofirstpage	Boolean to indicate whether a first page with histograms summarizing the whole measurement should be added
viewingwindow	See <a href="#">GGIR</a>
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 <a href="#">GGIR</a>
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 <a href="#">g.part1</a> )

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](mailto:v.vanhees@accelting.com)> Matthew R Patterson <[mpatterson@shimmersensing.com](mailto:mpatterson@shimmersensing.com)>

### Examples

```

## Not run:
# generate plots for the first 10 files:
g.plot5(metadataadir="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.part2](#)*

---

### Description

Creates report from milestone data produced by [g.part2](#). Not intended for direct use by package user

### Usage

```

g.report.part2(metadataadir = c(), f0 = c(), f1 = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> 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)
maxdur	see <a href="#">g.part2</a>
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 <a href="#">GGIR</a>
verbose	See details in <a href="#">GGIR</a> .

**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](mailto: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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> will recognise what the value of metadatadir is.
loglocation	see <a href="#">g.part4</a>

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)
data_cleaning_file	see <a href="#">GGIR</a>
sleepwindowType	see <a href="#">GGIR</a>
params_output	Parameters object, see <a href="#">GGIR</a>
verbose	See details in <a href="#">GGIR</a> .

### 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)>

---

g.report.part5      *Generate report from milestone data produced by [g.part5](#)*

---

### Description

Creates report from milestone data produced by [g.part5](#). Not intended for direct use by package user

### Usage

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

### Arguments

metadataadir	Directory that holds a folder 'meta' and inside this a folder 'basic' which contains the milestone data produced by <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> will recognise what the value of metadataadir 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)
loglocation	see <a href="#">g.part4</a>
params_cleaning	See details in <a href="#">GGIR</a> .
LUX_day_segments	see <a href="#">g.part5</a>
params_output	Parameters object, see <a href="#">GGIR</a>
verbose	See details in <a href="#">GGIR</a> .

### 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](mailto: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(metadataadir, params_output)
```

### Arguments

`metadataadir` 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 metadataadir 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)

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g.report.part6

*Generate report from milestone data produced by [g.part6](#)*

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### Description

Creates report from milestone data produced by [g.part6](#). Not intended for direct use by package user

### Usage

```
g.report.part6(metadataadir = 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 <a href="#">g.part1</a> . The folderstructure is normally created by <a href="#">g.part1</a> and <a href="#">GGIR</a> 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_cleaning	See details in <a href="#">GGIR</a> .
params_output	Parameters object, see <a href="#">GGIR</a>
verbose	See details in <a href="#">GGIR</a> .

### 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".

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g.shell.GGIR

*Wrapper function around function GGIR*

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### 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](#).

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 GGIR

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*Shell function for analysing an accelerometer dataset.*


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**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	Numeric (default = 1:5). Specify which of the five parts need to be run, e.g., mode = 1 makes that <a href="#">g.part1</a> is run; or mode = 1:5 makes that the whole GGIR pipeline is run, from <a href="#">g.part1</a> to <a href="#">g.part5</a> . Optionally mode can also include the number 6 to tell GGIR to run <a href="#">g.part6</a> 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	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	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.
f0	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).

<code>do.report</code>	Numeric (default = <code>c(2, 4, 5, 6)</code> ). For which parts to generate a summary spreadsheet: 2, 4, 5, and/or 6. Default is <code>c(2, 4, 5, 6)</code> . 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, <code>value = c()</code> , and then to merge the milestone data and turn report generation back on while setting <code>overwrite</code> to <code>FALSE</code> .
<code>configfile</code>	Character (default = <code>c()</code> ). Configuration file previously generated by function <code>GGIR</code> . See details.
<code>myfun</code>	List (default = <code>c()</code> ). External function object to be applied to raw data. See package vignette for detailed tutorial with examples on how to use the function embedding: <a href="https://cran.r-project.org/package=GGIR/vignettes/ExternalFunction.pdf">https://cran.r-project.org/package=GGIR/vignettes/ExternalFunction.pdf</a>
<code>verbose</code>	Boolean (default = <code>TRUE</code> ). to indicate whether console message should be printed. Note that warnings and error are always printed and can be suppressed with <code>suppressWarning()</code> or <code>suppressMessages()</code> .
<code>...</code>	Any of the parameters used <code>GGIR</code> . Given the large number of parameters used in <code>GGIR</code> 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 <code>GGIR</code> , but you can provide the parameters inside them as input to function <code>GGIR</code> .

## 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 explicitly entered argument values and default values for the argument that are not explicitly 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 explicitly provided argument values, the explicitly 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 parameter 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 done 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 `mvpthreshold = 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 GENActiv .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.
- window sizes** 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 occurrence 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.part1](#) 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:  $\text{round}(\sin((1:\text{length\_expansion}) / (900/\text{epochsize}))) * 15$ . 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, respectively, 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 csv-data 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 window sizes 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 window sizes 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 %H:%M:%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 = 72). 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 between 0.2 and 1 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 4Gb of RAM memory or with < 2GB memory per process when using `do.parallel = TRUE` a value below 1 is recommended.

**dynrange** Numeric (default = NULL). Provide dynamic range of 8 gravity.

**interpolationType** Integer (default = 1). To indicate type of interpolation to be used when re-sampling time series (mainly relevant for Axivity sensors), 1=linear, 2=nearest neighbour.

**rnc.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., +/-6g 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 m/s<sup>2</sup> 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 orientation 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:

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

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

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

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

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

**do.zcx** 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:

$$ENMO = \sqrt{acc_x^2 + acc_y^2 + acc_z^2} - 1$$

(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:

$$EN = \sqrt{acc_x^2 + acc_y^2 + acc_z^2}$$

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

$$MAD = \frac{1}{n} \sum |r_i - \bar{r}|$$

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

$$ENMOa = \sqrt{acc_x^2 + acc_y^2 + acc_z^2} - 1$$

(if ENMOa < 0, then ENMOa = |ENMOa|).

**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.bronddcounts** 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 BronddCounts because there are large number of activity counts in the physical activity and sleep research field. By calling them `_bronddcounts_` we clarify that these are the counts proposed by Jan Brondgeest and implemented in R by Ruben Brondeel. The `_bronddcounts_` 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 x-, 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 = 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 related 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 clipping) or not by `g.impute` in GGIR `g.part2`. Recommended setting is TRUE.

**TimeSegments2ZeroFile** Data frame (default = NULL). Optional data.frame to specify which time segments need to be ignored for the imputation, and acceleration metrics to be imputed by zeros. The data.frame is expected to contain two columns named windowstart and windowend, with the start- and end time of the time segment in POSIXlt class.

**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 (waking-waking, 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** Numeric (default = 1). See `data_masking_strategy` below.

**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 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 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).

**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 descrip-

tion 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 (`window-sizes[3]`, one hour as default) to check the conditions for nonwear identification and would flag as nonwear the mediumsize window (`window-sizes[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 related to physical activity as used in GGIR [g.part2](#) and GGIR [g.part5](#).

**mvpthreshold** Numeric (default = 100). Acceleration threshold for MVPA estimation in GGIR [g.part2](#). This can be a single number or an array of numbers, e.g., `mvpthreshold = c(100, 120)`. In the latter case the code will estimate MVPA separately for each threshold. If this variable is left blank, e.g., `mvpthreshold = c()`, then MVPA is not estimated.

**mvpadur** Numeric (default = 10). The bout duration(s) for which MVPA will be calculated. Only used in GGIR [g.part2](#).

**boutcritier** Numeric (default = 0.8). A number between 0 and 1, it defines what fraction of a bout needs to be above the `mvpthreshold`, 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 array of multiple numbers, e.g., `threshold.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 array of multiple numbers, e.g., `threshold.mod = 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 array 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.mvpa** Numeric (default = `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 = `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 = 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", "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 performed in GGIR [g.part3](#) and [g.part4](#).

**relyonguider** Boolean (default = FALSE). If TRUE, then sleep onset and waking time are defined based on timestamps derived from the guider. If participants were instructed NOT to wear the accelerometer during waking hours then set to 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.

**nights** 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 the 5 minute rolling average of counts is below 20 per cent of its standard deviation.
- HDCZA\_threshold** Numeric (default = 0.2) If HASPT.algo is set to "HDCZA" then HDCZA\_threshold will be used as threshold instead of 6th step in the diagram of Figure 1 in van Hees et al. 2018 Scientific Report (doi: 10.1038/s41598-018-31266-z). We have now simplified this step to a constant number, which can be modified via HDCZA\_threshold.
- 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 algorithms 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 algorithm 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 algorithm 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 related 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, LXX 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). 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. 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 mg, 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 mg 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(seq(0, 4000, by=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  $y = \text{constant} * \exp(x * \text{exponent})$

**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  $y = \text{constant} * \exp(x * \text{exponent})$

**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 indicate 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 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 = 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 csv-reports. 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](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
- 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:
     #-----
     window sizes = 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),
      window sizes = 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.ISO8601          Check whether character timestamp is in iso8601 format.
```

---

### Description

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

### Usage

```
is.ISO8601(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.ISO8601(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](https://en.wikipedia.org/wiki/ISO_8601) However, to generate plots in R we need to convert them back to POSIX

### Usage

```
iso8601chartime2POSIX(x, tz)
```

**Arguments**

x	Vector of timestamps in iso8601 in character format
tz	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	<i>Load default parameters</i>
-------------	--------------------------------

---

**Description**

Loads default parameter 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 <a href="#">GGIR</a> .

**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 <a href="#">GGIR</a> .

**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](https://en.wikipedia.org/wiki/ISO_8601)

**Usage**

POSIXtime2iso8601(x, tz)

**Arguments**

x	Vector of timestamps in POSIX format
tz	Timezone of data collection, e.g. "Europe/London". See <a href="https://en.wikipedia.org/wiki/List_of_tz_databases">https://en.wikipedia.org/wiki/List_of_tz_databases</a> for full list

**Author(s)**

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

<code>rmc.file</code>	Filename of file to be read if it is in the working directory, or full path to the file otherwise.
<code>rmc.nrow</code>	Number of rows to read, same as <code>nrow</code> argument in <a href="#">read.csv</a> and <code>nrows</code> in <a href="#">fread</a> . The whole file is read by default (i.e., <code>rmc.nrow = Inf</code> ).
<code>rmc.skip</code>	Number of rows to skip, same as <code>skip</code> argument in <a href="#">read.csv</a> and in <a href="#">fread</a> .
<code>rmc.dec</code>	Decimal used for numbers, same as <code>skip</code> argument in <a href="#">read.csv</a> and in <a href="#">fread</a> .
<code>rmc.firstrow.acc</code>	First row (number) of the acceleration data.
<code>rmc.firstrow.header</code>	First row (number) of the header. Leave blank if the file does not have a header.
<code>rmc.header.length</code>	If file has header, specify header length (numeric).
<code>rmc.col.acc</code>	Vector with three column (numbers) in which the acceleration signals are stored

<code>rmc.col.temp</code>	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 <a href="#">g.calibrate</a> .
<code>rmc.col.time</code>	Scalar with column (number) in which the timestamps are stored. Leave in default setting if timestamps are not stored.
<code>rmc.unit.acc</code>	Character with unit of acceleration values: "g", "mg", or "bit"
<code>rmc.unit.temp</code>	Character with unit of temperature values: (K)elvin, (C)elsius, or (F)ahrenheit
<code>rmc.unit.time</code>	Character with unit of timestamps: "POSIX", "UNIXsec" (seconds since origin, see argument <code>rmc.origin</code> ), "character", or "ActivPAL" (exotic timestamp format only used in the ActivPAL activity monitor).
<code>rmc.format.time</code>	Character string giving a date-time format as used by <a href="#">strptime</a> . Only used for <code>rmc.unit.time</code> : character and POSIX.
<code>rmc.bitrate</code>	Numeric: If unit of acceleration is a bit then provide bit rate, e.g. 12 bit.
<code>rmc.dynamic_range</code>	Numeric, if unit of acceleration is a bit then provide dynamic range deviation in g from zero, e.g. +/-6g 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.
<code>rmc.unsignedbit</code>	Boolean, if <code>unsignedbit = TRUE</code> means that bits are only positive numbers. if <code>unsignedbit = FALSE</code> then bits are both positive and negative.
<code>rmc.origin</code>	Origin of time when unit of time is UNIXsec, e.g. 1970-1-1
<code>rmc.desiredtz</code>	Deprecated, please see <code>desiredtz</code> .
<code>rmc.configtz</code>	Deprecated, please see <code>configtz</code> .
<code>rmc.sf</code>	Sample rate in Hertz, if this is stored in the file header then that will be used instead.
<code>rmc.headername.sf</code>	If file has a header: Row name (character) under which the sample frequency can be found.
<code>rmc.headername.sn</code>	If file has a header: Row name (character) under which the serial number can be found.
<code>rmc.headername.recordingid</code>	If file has a header: Row name (character) under which the recording ID can be found.
<code>rmc.header.structure</code>	Character used to split the header name from the header value, e.g. ":" or " "
<code>rmc.check4timegaps</code>	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.

<code>rmc.col.wear</code>	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.
<code>rmc.doresample</code>	Boolean to indicate whether to resample the data based on the available timestamps and extracted sample rate from the file header
<code>rmc.scalefactor.acc</code>	Numeric value (default 1) to scale the acceleration signals via multiplication. For example, if data is provided in m/s <sup>2</sup> then by setting this to 1/9.81 we would derive gravitational units.
<code>interpolationType</code>	Integer to indicate type of interpolation to be used when resampling time series (mainly relevant for Axivity sensors), 1=linear, 2=nearest neighbour.
<code>PreviousLastValue</code>	Automatically identified last value in previous chunk of data read.
<code>PreviousLastTime</code>	Automatically identified last timestamp in previous chunk of data read.
<code>desiredtz</code>	Timezone in which device was worn.
<code>configtz</code>	Timezone in which device was configured. If equal to <code>desiredtz</code> you can leave this in its default value.
<code>header</code>	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.part1` 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 original 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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