

# Package ‘RCflux’

December 6, 2018

**Title** Calculate Closed-Chamber Gas Fluxes  
**Version** 0.5  
**Description** Calculate closed-chamber gas fluxes.  
**Depends** R (>= 3.2.3),  
plyr,  
dplyr,  
stringr,  
yaml,  
tools,  
nlme,  
ggplot2,  
HMR,  
readxl  
**License** MIT  
**LazyData** true  
**RoxygenNote** 6.1.1

## R topics documented:

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RCflux-package	<i>RCflux calculates fluxes from closed chamber data</i>
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## Description

Gas concentration data may come from a GC or other gas analysers Data formats for several sources can be read in: [list here] Output is written to CSV and PNG files. Calculates gas fluxes from closed chamber data.

**Details**

The only function you're likely to need from **RCflux** is [calcFlux](#). Refer to the vignettes for details of how to use it - use `vignette()`.

**Author(s)**

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calcFlux	<i>This function calculates gas fluxes from GC sample data</i>
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**Description**

This function converts Agilent GC output.

**Usage**

```
calcFlux(filelist)
```

**Arguments**

filelist            A file listing the Agilent GC output files to be processed.

**See Also**

[convert\\_GC\\_output](#) for the higher-level function which calls this.

**Examples**

```
calcFlux("./filelist.txt")
```

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convert_GC_output	<i>Converts Agilent GC output to a standard format</i>
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**Description**

This function converts Agilent GC output.

**Usage**

```
convert_GC_output(filelistIn)
```

**Arguments**

filelistIn            A file listing the Agilent GC output files to be processed.

**See Also**

[calcFlux](#) for the higher-level function which calls this.

**Examples**

```
convert_GC_output("f:/0Peter/misc/stats/GCflux/filelist_aGCxls.txt")
convert_GC_output("S:/DISE_Instrumentation/GC/Agilent/processing/filelist_aGCxls.txt")
```

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read_GC_input	<i>Reads a GC input file</i>
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**Description**

This function reads a GC input file.

**Usage**

```
read_GC_input(fname)
```

**Arguments**

fname	A GC input file.
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**See Also**

[convert\\_GC\\_output](#) for the higher-level function which calls this.

**Examples**

```
read_GC_input(fname)
```

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read_GC_output	<i>This function reads Agilent GC output.</i>
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**Description**

Reads Agilent GC output

**Usage**

```
read_GC_output(fname)
```

**Arguments**

fname	A GC output file from the Agilent GC.
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**See Also**

[convert\\_GC\\_output](#) for the higher-level function which calls this.

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standardiseNames	<i>Standardises names in a data frame</i>
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**Description**

This function standardises names in a data frame.

**Usage**

```
standardiseNames(df)
```

**Arguments**

df	A data frame.
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**See Also**

[convert\\_GC\\_output](#) for the higher-level function which calls this.

**Examples**

```
tdf <- data.frame(Plot = c(1, 2), Compound.Name = c("CO2", "CH4"))
tdf
names(tdf)
tdf <- standardiseNames(tdf)
tdf
names(tdf)
```

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