

# Package: PEcAn.RothC (via r-universe)

June 5, 2026

**Type** Package

**Title** PEcAn Package for Integration of the RothC Model

**Version** 0.0.0.9000

**Description** This module provides functions to link the Rothamstead soil carbon model ``RothC'' to PEcAn. Uses RothC  $\geq$  2.0, available on GitHub  
<[https://github.com/Rothamsted-Models/RothC\\_Code](https://github.com/Rothamsted-Models/RothC_Code)> as well as directly from the RothC team  
<<https://www.rothamsted.ac.uk/rothamsted-carbon-model-rothc>>.

**URL** <https://pecanproject.github.io>

**BugReports** <https://github.com/PecanProject/pecan/issues>

**Depends** R ( $\geq$  4.1)

**Imports** dplyr, lubridate ( $\geq$  1.6.0), ncdf4, PEcAn.logger, PEcAn.utils ( $\geq$  1.8.0), rlang, stats, utils

**Suggests** testthat ( $\geq$  3.0.0), withr

**SystemRequirements** RothC

**OS\_type** unix

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**Copyright** Authors

**Encoding** UTF-8

**RoxygenNote** 7.3.3

**Config/testthat/edition** 3

**Config/pak/sysreqs** libicu-dev libnetcdf-dev libssl-dev libudunits2-dev

**Repository** <https://pecanproject.r-universe.dev>

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**RemoteUrl** <https://github.com/PecanProject/pecan>

**RemoteRef** HEAD

**RemoteSha** 3654c6a1cc83cd02402ca26bd012ed1e111bef1b

**RemoteSubdir** models/rothc

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met2model.RothC	<i>Extract monthly weather from CF file for input to RothC</i>
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### Description

Input files need to be named '`<in.path>/<in.prefix>.YYYY.nc`'

### Usage

```
met2model.RothC(
  in.path,
  in.prefix,
  outfolder,
  start_date,
  end_date,
  overwrite = FALSE
)
```

### Arguments

<code>in.path</code>	path on disk where CF files live
<code>in.prefix</code>	prefix for each file
<code>outfolder</code>	location where model specific output is written.
<code>start_date, end_date</code>	When to start and end output. Specify as exact dates, but output will be padded to whole months.
<code>overwrite</code>	logical: replace output files if they already exist?

### Details

Output files are named '`<outfolder>/<in.prefix>.YY-mm.YY-mm.dat`' with one line per month and columns for temperature, rainfall, and evaporation.

Note that the created file contains only weather data and not any of the soil or management data needed for RothC's single combined input file. See '`write.config.RothC()`' for assembly into a model-ready `RothC_input.dat`'.

**Value**

data frame summarizing file metadata

**Author(s)**

Chris Black

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model2netcdf.RothC     *Convert RothC output to PEcAn-formatted netCDF*

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**Description**

Convert RothC output to PEcAn-formatted netCDF

**Usage**

model2netcdf.RothC(outdir, sitelat, sitelon, start\_date, end\_date)

**Arguments**

outdir	Location of model output
sitelat	Latitude of the site
sitelon	Longitude of the site
start_date	Start time of the simulation
end_date	End time of the simulation

**Author(s)**

Chris Black

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read\_restart.RothC     *Read restart template for SDA*

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**Description**

Reads restart files from RothC. Not implemented yet.

**Usage**

read\_restart.RothC(outdir, runid, stop.time, settings, var.names, params)

**Arguments**

outdir	Output directory
runid	Run ID
stop.time	Year that is being read
settings	PEcAn settings object
var.names	Variable names to be extracted
params	Any parameters required for state calculations

**Value**

Forecast numeric matrix

**Author(s)**

Chris Black

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write.config.RothC      *Writes a RothC config file.*

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**Description**

Requires a pft xml object, a list of trait values for a single model run, and the name of the file to create

**Usage**

```
write.config.RothC(defaults, trait.values, settings, run.id)
```

**Arguments**

defaults	list of defaults to process
trait.values	vector of samples for a given trait
settings	list of settings from pecan settings file
run.id	id of run

**Value**

configuration file for MODEL for given run

**Author(s)**

Chris Black

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write\_restart.RothC    *Write restart template for SDA*

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**Description**

Not implemented yet.

**Usage**

```
write_restart.RothC(  
  outdir,  
  runid,  
  start.time,  
  stop.time,  
  settings,  
  new.state,  
  RENAME,  
  new.params,  
  inputs  
)
```

**Arguments**

outdir	outout directory
runid	run id
start.time	Time of current assimilation step
stop.time	Time of next assimilation step
settings	pecan settings list
new.state	Analysis state matrix returned by sda.enkf
RENAME	flag to either rename output file or not
new.params	optional, additional params to pass write.configs that are deterministically related to the parameters updated by the analysis
inputs	new input paths updated by the SDA workflow, will be passed to write.configs

**Author(s)**

Chris Black

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