gnumaker

Version: 0.0.0.9009

Overview

gnumaker makes it easy to create and use GNU Makefiles to aid a reproducible work flow for data analysis projects.

GNU Make is the defacto standard for efficiently rerunning appropriate steps in the data analysis or reporting process if a particular file is changed. Only the necessary steps are rerun.

Rather than creating a new system for setting up and building output from statistical software syntax files, **gnumaker** leverages off existing GNU Make rules. These rules, for R, Sweave, R Markdown, Stata, SAS and other syntax files are available at r-makefile-definitions on Github. These are described in P Baker (2020) Using GNU Make to Manage the Workflow of Data Analysis Projects, *Journal of Statistical Software* (Accepted).

For those not familiar with GNU Make, **gnumaker** allows simple dependencies between files to be specified to produce a working Makefile and the associated directed acyclic graph (DAG). I'd welcome Github issues containing error reports or feature requests. Alternatively, you can email the package maintainer at drpetebaker at gmail dot com.

Installation

You can install the development version of **gnumaker** from GitHub with:

```
## if you don't have devtools installed, automatically install it from CRAN
if (!requireNamespace("devtools", quietly = TRUE))
   install.packages("devtools")
devtools::install_github("petebaker/gnumaker")
```

Usage

There are currently four key functions in **gnumaker**. These are:

- create_makefile() creates a gnu_makefile object given the specified dependencies between syntax, data and output files,
- write_makefile() writes a gnu_makefile object to a Makfile on disk,
- info_rules() provides information about data analysis GNU Make rules for various target and dependency filename extensions, and
- plot() plots the DAG of a gnu_makefile object.

Example

Suppose we have a data file simple.csv and use read.R to read and clean the data. After storing the cleaned data in a .RData file, we then employ linmod.R to plot and analyse the data. Next, using the stored results, two reports report1.pdf and report2.docx are produced from report1.Rmd and report2.Rmd. The workflow may be encapsulated in a Makefile which is then employed to manage the process and generate or regenerate any intermediate files when the data or syntax changes.

Rather than writing GNU Make commands, we can employ the **gnumaker** package to create an appropriate Makefile called Makefile.demo with:

To plot the corresponding DAG, which shows the relationships between target files (wheat coloured circles) which we can (re)generate from a minimal set of syntax and data files (green rectangles), use:

plot(gm1)

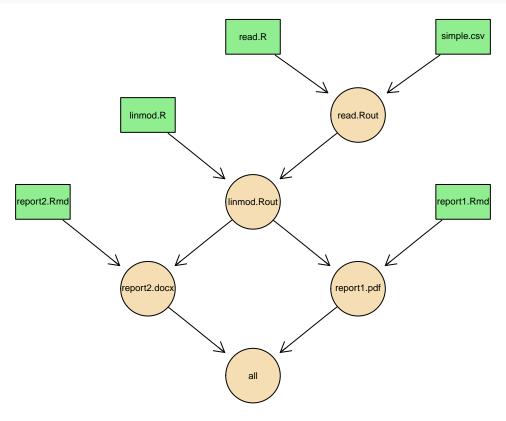


Figure 1: DAG of Makefile for simple example. The DAG of the gnu_makefile object can be produced with plot(gm1). Using the minimal set of files (shown in green rectangles), then GNU Make allows us to (re)generate all other files shown as wheat coloured circles)

Details

Using the **gnumaker** package we simply need to provide a list of targets to the the **create_makefile** function where the components specify a target as a name and dependency file(s) as a character vector. The package uses the GNU Make pattern rules in **r-rules.mk** to choose file names for targets but we can override the defaults.

For instance, in this example the first two dependency files are simple.csv and read.R so we provide the first target as the first component of the list as read = c("read.R", "simple.csv"), where the name read can be anything we like. Note that the target filename for read will be derived from the first failename in the list read.R. The second target depends on the read target and linmod.R and so we specify this with linmod = c("linmod.R", "read") and so on.

Using the function create_makefile to create a gnu_makefile object, target file names are substituted using

defaults and the appropriate Make commands are rearranged using the DAG of the relationships. For instance, the default target file for the first dependency in the read component, which is read.R, becomes read.Rout but we can change the default target file extension for all .R files using the default.exts argument and specify say a HTML target file with default.exts = list(R = "html").

Finally we specify the first target (usually all) as two reports report1.pdf and report2.docx using target.all = c("rep1", "rep2") which by default would be report1.html and report2.html but which we specify as report1.pdf and report2.docx by specifying the option all.exts = list(rep1 = "pdf", rep2 = "docx").

Once we have constructed a suitable <code>gnu_makefile</code> object then we write it to disk with <code>write_makefile</code>. To run all R script files and analyses in order we simply type <code>make</code> in a terminal or set up RStudio or our IDE to use GNU Make as the build mechanism which allows us to (re)run analyses by pressing the appropriate Build button.

The Makefile, with optional comment for linmod target, is specified using:

A Makefile Makefile.demo is produced with write_makefile(gm1)

```
write_makefile(gm1, file = "Makefile.demo")
#> File: Makefile.demo written at Sun Jul 19 15:59:42 2020
# File: Makefile.demo
# Created at: Sun Jul 19 15:59:42 2020
# Produced by gnumaker: 0.0.0.9009 on R version 4.0.2 (2020-06-22)
# Before running make, please check file and edit if necessary
# .PHONY all target: run when 'make' is invoked without a target
.PHONY: all
all: report1.pdf report2.docx
# report1.pdf depends on report1.Rmd, linmod.Rout
report1.pdf: report1.Rmd linmod.Rout
# report2.docx depends on report2.Rmd, linmod.Rout
report2.docx: report2.Rmd linmod.Rout
# plots and analysis using 'linmod.R'
linmod.Rout: linmod.R read.Rout
# read.Rout depends on read.R, simple.csv
read.Rout: read.R simple.csv
# include GNU Makfile rules. Most recent version available at
# https://github.com/petebaker/r-makefile-definitions
include ~/lib/r-rules.mk
```

```
# remove all target, output and extraneous files
.PHONY: cleanall
cleanall:
    rm -f *~ *.Rout *.RData *.docx *.pdf *.html *-syntax.R *.RData
```

We can use the function info_rules to determine the possible target files for dependency files. For instance, what target files have Makefile rules for an .R R syntax file?

```
info_rules("R")
#> Possible filename extensions for 'R':
#> [1] "docx" "html" "odt" "pdf" "Rout" "rtf"
#>
#> Default: 'Rout'
#>
#> Example rules:
#> example1.Rout: example1.R dep_file2 dep_file3
#> or
#> example1.Rout: {@:.Rout=.R} dep_file2 dep_file3
#>
#> NB: For further help on Makefile rules, type 'make help' in a terminal once
#> an appropriate 'Makefile' is present in the current directory
```

For .Rmd R Markdown files, use

```
info_rules("Rmd")
#> Possible filename extensions for 'Rmd':
#> [1] "_beamer-handout.Rmd" "_beamer.pdf"
                                                   "_ioslides.html"
#> [4] "_slidy.html" "_tufte.pdf"
                                                   "-syntax.R"
#> [7] "docx"
                             "html"
                                                   "odt"
#> [10] "pdf"
                              "pptx"
                                                    "rtf"
#>
#> Default: 'html'
#>
#> Example rules:
#> example1.html: example1.Rmd dep file2 dep file3
#> example1.html: {@:.html=.Rmd} dep_file2 dep_file3
#>
#> Other options are available for R Markdown files, such as:
#>
#> example1_ioslides.html: example1.Rmd dep_file2 dep_file3
#> example1_beamer.pdf: example1.Rmd dep_file2 dep_file3
#> to produce ioslide and beamer presentation formats.
#>
#> An R syntax file can be produced with
#> make example1-syntax.R
#> and a similar rule can be specified if required with
#> example1-syntax.R: example1.Rmd dep_file2 dep_file3
#> NB: For further help on Makefile rules, type 'make help' in a terminal once
       an appropriate 'Makefile' is present in the current directory
```

For more examples, see the gnumaker vignette (under construction).

Note

 ${f gnumaker}$ is under construction and could change (and improve) rapidly at various times but this depends on work/life balance.