Statistics 360: Advanced R for Data Science Lecture 5

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R packages, CRAN and GitHub

- ► CRAN and the R package system is the key to R's success, allowing contributions from hundreds of scientists outside the "R core team".
- CRAN has strict quality-control checks and requirements.
- ▶ It is possible to distribute R packages via GitHub without the CRAN checks.

R packages

- What is an R package?
 - An easy-to-install collection of R functions, documentation and example data.
- Why make an R package?
 - Share your work with others
 - Get credit for your work
 - Make it easier for you to use your own work.

Making R packages

- ► The definitive source for making R packages is the "Writing R Extensions" document, which you can find at https://mirror.rcg.sfu.ca/mirror/CRAN/doc/manuals/r-release/R-exts.html
- A minimal R package is a folder containing DESCRIPTION and NAMESPACE files and an R subfolder.
- Other possible subfolders are data, demo, exec, inst, man, po, src, tests, tools and vignettes.
- Other possible files in the directory are INDEX, configure, cleanup, LICENSE, LICENCE and NEWS.

Getting Started

▶ A helper function called package.skeleton() in the utils package that comes with R takes a package name and list of R objects as arguments and sets up directories, files and skeleton help files.

```
rm(list=ls()) # clear workspace
#source("../mars.R") # location of my mars source file
#package.skeleton(name="mars",ls()) # Can't be re-run
```

```
Creating directories ...
Creating DESCRIPTION ...
Creating NAMESPACE ...
Creating Read-and-delete-me ...
Saving functions and data ...
Making help files ...
Done.
Further steps are described in './mars/Read-and-delete-me'.
```

\$ more mars/Read-and-delete-me

- * Edit the help file skeletons in 'man', possibly combining help files for multiple functions.
- * Edit the exports in 'NAMESPACE', and add necessary imports.
- * Put any C/C++/Fortran code in 'src'.
- * If you have compiled code, add a useDynLib() directive to 'NAMESPACE'.
- * Run R CMD build to build the package tarball.
- * Run R CMD check to check the package tarball.

Read "Writing R Extensions" for more information.

devtools

- ► A package called devtools provides more help.
 - devtools was started by Hadley Wickham and has since expanded into a collection of tools for developing R packages, described in a work-in-progress book called "R Packages" https://r-pkgs.org/index.html
 - We'll skim Chapter 2, which provides an overview of functionality.
- In addition to automating more of the process, devtools uses roxygen2 to create documentation from structured comments in your R source files.
 - ► C++ programmers will recognize the approach from Doxygen

Getting started with create_package()

- Call create_package() to initialize an R package directory and new R project
- Recommended that this new directory **not** be part of an existing R project or be under version control.
- Launches a new RStudio session in the newly-created directory/project
 - Switch to this new session.
 - If needed, call use_git() to initialize a git repository

```
# install.packages("devtools")
library(devtools)
## Loading required package: usethis
create_package("/Users/bwl/mypackages/MARS")
## v Setting active project to '/Users/bwl/mypackages/MARS'
## v Leaving 'DESCRIPTION' unchanged
## Package: MARS
## Title: What the Package Does (One Line, Title Case)
## Version: 0.0.0.9000
## Authors@R (parsed):
       * First Last <first.last@example.com> [aut, cre] (YOUR-ORCID-ID)
##
## Description: What the package does (one paragraph).
## License: `use_mit_license()`, `use_gpl3_license()` or friends to pick a
       license
##
## Encoding: UTF-8
## Roxygen: list(markdown = TRUE)
## RoxygenNote: 7.2.3
## v Leaving 'NAMESPACE' unchanged
## v Setting active project to '<no active project>'
```

Start your R source files

- ▶ Copy your R source files to the R sub-folder.
 - For now there is just mars.R, but later we will have predict.mars.R, plot.mars.R, etc.
- Call load_all() to load the source into your R session.
 - Rather than just source in your R code, load_all() loads the R functions as a package.
 - Gives a better sense of how the code will behave when loaded by a user.

```
library(devtools) # call in new R session
# load_all()
```

check

- ▶ R CMD check from the command line
- check()
- ▶ Will throw a warning of a non-standard licence

Edit DESCRIPTION file

Add your name, collaborators, etc.

Add a licence

Helper functions add a copy of the relevant license to your package main directory and update the Licence field of the DESCRIPTION file.

```
# use_gpl3_license()
```

Add documentation comments to your source file.

Open your .R file(s), place your cursor in one of the functions and click Code->Insert Roxygen Skeleton to insert a skeleton of the roxygen2-style comments above the function.

```
#' Title
#'
#' @param formula
#' @param data
#' @param control
#' @param ...
#'
#' @return
#' @export
#'
#' @examples
```

```
Multivariate Adaptive Regression Splines (MARS)
#'
#' Oparam formula an R formula
  Oparam data a data frame containing the data for the model
#' @param control an object of class 'mars.control'
  Oparam ... other arguments -- currently not used
#'
  Oreturn an object of class 'mars'
#'
  @export
#'
  @examples
  mm <- mars(wage ~ age,data=ISLR::Wage)</pre>
#'
  @import stats
#' @import ISLR
```

Call document()

- Calling document() will create the .Rd file from your comments, and add mars as an "export" and stats and ISLR as "imports" in the mars NAMESPACE file.
 - See below for Imports
- > document()
 Updating mars documentation
 Loading mars
 Writing NAMESPACE
 Writing mars.Rd
 > load_all()
 Loading mars
 > ?mars
 Rendering development documentation for 'mars'

Add other packages you depend on

- Reference: R packages, Chapter 8
- Your package's reliance on other packages can be as
 - "Depends" you expect that users will always want to call library(package) when loading your package – not common these days
 - "Imports" you need to call functions from the NAMESPACE of another package. Add these to the DESCRIPTION file with use_package().
 - "Suggests" Packages that are not crucial, but helpful (e.g., used in examples). Add these to the DESCRIPTION file yourself
 - "Enhances" Packages that are enhanced by yours not common

```
# use_package("stats")
# use_package("ISLR") # used in mars example
# document()
```

check

- ► Call check() to check your package using the same checks as R CMD check from the command line.
- Should return no errors, warnings or notes.

check()

install

► Call install() to install the package in your R library.

install()

Implementation of an R package recap

- 1. call create_package() to initialize an R package directory and a new project.
- Copy your R scripts to the R directory of your new package and call load_all() to load them into your R session.
- 3. call devtools::check() to check that the package builds.
- 4. edit the **DESCRIPTION** file
- 5. add a licence
- start your documentation by inserting an Roxygen skeleton for your main mars() function then fill in your title, function arguments, return value, examples and any import you require.
- 7. call document() to generate an.Rd file and update NAMESPACE
- call use_package() to add any package dependencies/imports to DESCRIPTION
- call devtools::check() again to make sure the package will still build
- copy your R package files to the directory under version control for your course project, then commit the new files and push them to the GitHub repo.

Topics not covered

- ➤ You have enough to get you started, but you will need to learn more as you go . . .
- Our MARS implementation will use the S3 object oriented programming system.
 - You will need to read about documenting classes, generics and methods in Chapter 10, SS 7
- ► Your final project document will include a "vignette"
 - ► Chapter 11
- You will need to write "unit tests" using the testthat package
 - Chapter 12