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

```r
# install.packages("devtools")
library(devtools)
```

```
## Loading required package: usethis
```

```r
create_package("/Users/bwl/STAT360")
```

```
## v Setting active project to '/Users/bwl/STAT360'
```

```
## Package: STAT360
## 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 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.

```r
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)
#'
#' @param formula an R formula
#' @param data a data frame containing the data for the model
#' @param control an object of class 'mars.control'
#' @param ... other arguments -- currently not used
#'
#' @return an object of class 'mars'
#' @export
#'
#' @examples
#' mm <- mars(wage ~ age,data=ISLR::Wage)
#' @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

```r
# 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.
2. 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
6. 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
8. call `use_package()` to add any package dependencies/imports to DESCRIPTION
9. call `devtools::check()` again to make sure the package will still build
10. 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