Building R Packages

What are R Packages?

R Packages

- 1. Sets of Functions
- 2. Set of Functions + Documentation
- 3. Set of Functions + Documentation + Data
- 4. Set of Functions + Documentation + Data + Vignettes
- 5. Set of Functions + Documentation + Data + Vignettes + Versions
- 6. Set of Functions + Documentation + Data + Vignettes + Versions + Dependencies

Windows and RTools

If you have a Windows machine, you need to install RTools from https://cran.r-project.org/bin/windows/Rtools/ (choose the frozen one).

Using Rtools40 on Windows

Starting with R 4.0.0 (released April 2020), R for Windows uses a brand new toolchain bundle called rtools40.

This version of Rtools upgrades the mingw-w64 gcc toolchains to version 8.3.0, and introduces a new build system based on msys2, which makes easier to build and maintain R itself as well as the system libraries needed by R packages on Windows. For information about the latter, follow the links at the bottom of this document.

This documentation is about rtools40, the current version used for R 4.0.0 and newer. For information about previous versions of Rtools that can be used with R 3.6.3 or older, please visit this page.

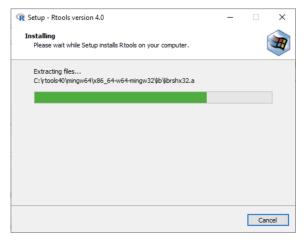
Installing Rtools40

Note that rtools40 is only needed build R packages with C/C++/Fortran code from source. By default, R for Windows installs the precompiled "binary packages" from CRAN, for which you do not need rtools!

To use rtools40, download the installer from CRAN:

- On Windows 64-bit: rtools40-x86 64.exe (recommended: includes both i386 and x64 compilers)
- On Windows 32-bit: rtools40-i686.exe (i386 compilers only)

Note for RStudio users: please check you are using the latest version of RStudio (at least 1.2.5042) to work with rtools40.



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Starting Up

Use RStudio and the devtools and usethis packages. It's easier.

```
install.packages(c("devtools", "usethis"))
```

In RStudio, File -> New Project -> New Directory -> R Package with devtools (scroll down), with a name:

- must start with letter
- no underscores
- periods allowable or use CamelCase
- · can have numbers

Checking Package Names

Try the available::available function:

```
available::available("ggplot")

## — ggplot

## Name valid: 
## Available on CRAN: 
## Available on Bioconductor: 
## Available on GitHub: 
## Abbreviations: http://www.abbreviations.com/ggplot

## Wikipedia: https://en.wikipedia.org/wiki/ggplot

## Wiktionary: https://en.wiktionary.org/wiki/ggplot

## Urban Dictionary:

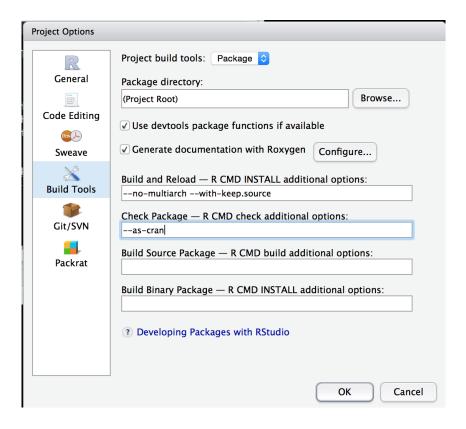
## Not found.

## Sentiment:???
```

Setting Up

Go to Build -> Configure Build Tools

Add --as-cran to "Check Package" (useful later)

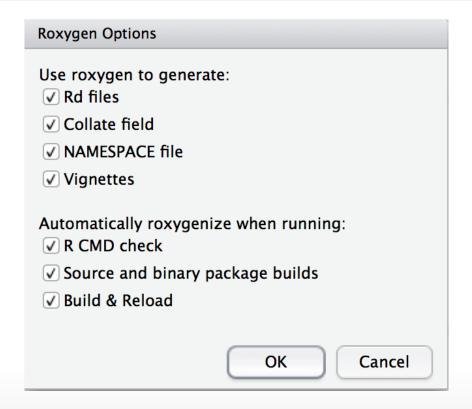


Setting Up

Documentation is a pain, but it's worthwhile. Writing out argument definitions makes it easier to identify if argument names make sense.

Click Generate documents with Roxygen. If that is gray, install roxygen2:

install.packages("roxygen2")



Click "Configure" - click all the boxes.

Modifying the Skeleton

Things to do:

- Delete R/hello.R (skip if used devtools)
- Delete man/hello.Rd (skip if used devtools)
- 3. In RStudio, Build → Configure Build Tools → Generate Documentation with Roxygen, make sure that's clicked. I click Vignettes and Build and Reload.
- 4. In RStudio, add --as-cran under the "Check options" in Build → Configure Build Tools.
- 5. Delete the NAMESPACE file (skip devtools). If building fails, add an empty file with # Generated by roxygen2: do not edit by hand at the top and rerun.

DESCRIPTION file

In the RStudio project, go to "Go to file/function" search bar on the menu bar.

- This searches through the files in the package.
 - Also searches for **function names** and can go to that function in the file

Type "DESCRIPTION" and open that file.

DESCRIPTION file

- "Title What the Package Does (Title Case)
- "Author: YOURNAME"
- "Maintainer: YOURNAME your@email.com" -> we will remove this
- "Description: Use paragraph prose here. Don't start with word package" Use four spaces when indenting paragraphs within the Description.
- "License:", one of GPL-2 GPL-3 LGPL-2 LGPL-2.1 LGPL-3 AGPL-3 Artistic-2.0 BSD_2_clause BSD_3_clause MIT

License

```
usethis::use_gpl3_license("John Muschelli")
```

https://www.r-project.org/Licenses/

Authors

I add this to the DESCRIPTION file:

Use Authors@R even if there is only one author.

DESCRIPTION file: additional fields

- You do not use library or require in functions in a package
- Imports: package1, package2
 - packages with specific functions called in package
 - Anything other than base package needs to be imported (stats, methods)
- Depends: package3, package5
 - packages with ALL functions loaded from package
- Suggests: package4, package6
 - used in **examples** or **vignettes**

```
usethis::use_package("tidyr", type = "Imports")
usethis::use_package("dplyr", type = "Suggests")
```

Description

Change the Description so that it's a sentence and it ends with a period. Also I keep putting single quotes around weird words (like science-specific). Make sure to put links in angle brackets (<http...>). Use DOIs if you can. If you go too long on a line, indent it with 4 spaces " ".

Roxygen2

Roxygen allows for functions and documentation in the same file. Let's make a function:

```
top = function(x, n) {
    xx = x[1:n, 1:n]
    hist(xx)
    print(xx)
}
```

Save this to top.R file in R/ (where R functions are).

Roxygen2

Highlight the following code:

```
top = function(x, n) {
```

Go to Code -> Insert Roxygen Skeleton

Output:

```
#' Title
#'

#' @param x
#' @param n
#'

#' @return
#' @export
#'
#' @examples
```

You can add @title and @description tags:

```
#' @title
#' @description
#'
#' @param x
#' @param n
#'
#' @return
#' @export
#'
#' @examples
```

- @param stands for a parameter/argument for that function.
- @return denotes what the function returns. This is required.
- · @export when people install your package, can they use this function
 - non-exported functions are usually helpers, really small, or not fully formed yet
- @examples code to show how the function works. Wrap functions in \dontrun{} if not wanted to run

```
#' @title Print the top of a matrix
#' @description \code{top} is a small function to not just present the first rows
#' of a matrix, but also the first number of columns
# "
#' @param x a \code{matrix}
#' @param n Number of rows and columns to display of the matrix
# "
#' @return A \code{NULL}
#' @export
#' @examples
#' mat = matrix(rnorm(100), nrow = 10)
\#' top(mat, n = 4)
#' \dontrun{
#' top(mat, n = 10)
# " }
```

Functions: a little style

- 1. Create a file for each function (preference) or at least group. Name file function name.
- 2. Optional arguments: Set to NULL and use is.null() to test
- 3. Put functions together: use # ' @rdname.
- 4. See # ' @inheritParams for different functions with the same arguments.
- 5. Add verbose argument for printing
- 6. Use message (not cat) for printing. Someone can use suppressMessages to stop the printing.
- 7. Pass . . . to a main function for additional options for the user.
- 8. Have examples
- 9. Learn do.call(FUNCTION, args = list_of_arguments)
- 10. Notify/warn/message whenever you have to.

NAMESPACE

The NAMESPACE files tells the R package what to import and export. In Roxygen:

- @export adds this to the NAMESPACE file
 - when package is installed, users can call this function
- @import in roxygen, if you want to import a package, you say @import PACKAGENAME
 - imports **ALL** functions from that package
 - if package is listed under Depends in DESCRIPTION, then the whole package is loaded when you load your package
 - otherwise it simply exposes them for your package to use them, but not the user, users still have to do library (PACKAGENAME)

NAMESPACE

- @importFrom in roxygen, if you want to import a function, you say @import PACKAGENAME func1 func2
 - only imports these functions. Better way of doing things.
 - if pkgA has function A and pkgB has functions A and B, if @import pkgA A, @import pkgB B, then if you call A(), R knows it's from pkgA
 - you must import anything explicitly other than from the base package, includying anything from stats (e.g. quantile) or graphics (e.g. hist)

Add @importFrom graphics hist to your top.R file

Build and Reload

- Go to Build -> Build and Reload the package
 - First time you may see some warnings (no NAMESPACE file!)
 - Rerunning should get rid of these
 - look in the folders
- Then try Build -> Check Package

Using Data

The data/ directory is where data goes, it must be named .RData or rda. The use data function can do this for you:

```
usethis::use_data(DATAOBJECT, compress = "xz")
```

The output will be DATAOBJECT.rda in the data folder. You can use this in your package

Making Data

That's not reproducible!

The data-raw directory can be data you want to create (such as simulated data).

This will have scripts with use data at the end to make the data.

usethis::use data raw()

Documenting Data

Note how DATAOBJECT is the name of the object/rda. Now we can document the data as follows:

```
#' @title Some object to document
#'

#' @description A list containing things
#'

#' @format A list with 7 elements, which are:

#' \describe{

#' \item{x}{first thing}

#' \item{y}{second thing}

#' }

"DATAOBJECT"
```

Different kinds of data

The inst/ directory will copy any of the contents to the installed directory path. So if blah.csv was in inst/ then it will be in the directory.

Most times, however, people put data in inst/extdata to separate folders out.

You can use find.package to find the installed directory:

```
find.package("readr")
```

[1] "/Users/johnmuschelli/Library/R/4.0/library/readr"

To get files, though, you should use system.file:

```
system.file("extdata", package = "readr")
```

[1] "/Users/johnmuschelli/Library/R/4.0/library/readr/extdata"

Different kinds of data

If you pass in multiple characters, it assumes you put it together with file.path:

```
system.file("extdata", "challenge.csv", package = "readr")

## [1] "/Users/johnmuschelli/Library/R/4.0/library/readr/extdata/challenge.csv"
```

The mustwork argument is useful for making sure the file exists:

```
system.file("extdata", "asdfsdf.csv", package = "readr", mustWork = TRUE)
```

Error in system.file("extdata", "asdfsdf.csv", package = "readr", mustWork = TRUE): no file

Using the file system

```
1. file.path > paste for paths
2. Use file.exists. Use any and all
3. file.remove if you need to delete things
4. Make temporary (empty, non-existent) files, with extension:
  tempfile(fileext = ".csv").
5. dir.create to create and unlink to destroy directories.
6. Using tempdir for stuff that's intermediate.
7. Tempdir sub-directory: tdir = tempfile(); dir.create(tdir);
  on.exit({ unlink(tdir) })
8. file.copy and file.rename
9. download.file Or httr::GET.
```

Vignettes

http://r-pkgs.had.co.nz/vignettes.html

A package has data + code + dependencies. A vignette can tie this together to tell you **how** to use the package. Typically it is an analysis.

THIS IS EXACTLY WHAT A REPRODUCIBLE PAPER IS!

usethis::use_vignette("my-vignette")

· Can make private packages that are the workflow for your paper

Unit tests

The testthat package is great for unit testing. Put test scripts in tests/testthat, always named test-DESCRIPTOR.R. To set up testthat:

```
usethis::use_testthat()
```

And a specific test:

```
usethis::use test("name of test")
```

General Rule: Any package issue turns into a test.

Unit tests

The testthat package is great for unit testing. Put test scripts in tests/testthat, always named test-DESCRIPTOR.R. To set up testthat:

```
testthat::context("OVERALL DESCRIPTION OF TESTS IN THIS FILE")
testthat::test_that("Description of this test", { MYCODE })
testthat::expect_equal(OUTPUT, 1234.34535)
testthat::expect_identcal(OUTPUT1, OUTPUT2)
testthat::expect_true(SOME_OUTPUT)
testthat::expect_silent({ no_warning_error_code })
testthat::expect_message({ some_warn }, "a[test]regexp")
```

Continuous integration (Thoughtworks.com)

Continuous Integration (CI) is a development practice that requires developers to integrate code into a shared repository several times a day. Each check-in is then verified by an automated build, allowing teams to detect problems early.

Building CI and README

```
usethis::use_git()
usethis::use_github() # must have GITHUB_PAT set up
usethis::use_github(protocol = "https") # must have GITHUB_PAT set up
usethis::use_readme_rmd()
usethis::use_github_action_check_standard()
usethis::use_appveyor()
usethis::use_travis()
```

GitHub Actions - Build all 3 OSes

https://github.com/r-lib/actions



README.md

GitHub Actions for the R language



This repository stores GitHub Actions for R projects, which can be used to do a variety of CI tasks. It also has a number of example workflows which use these actions.

- 1. r-lib/actions/setup-r Sets up R
- 2. r-lib/actions/setup-pandoc Sets up pandoc
- 3. r-lib/actions/setup-tinytex Sets up LaTeX with tinytex
- 4. r-lib/actions/pr-fetch Fetches changes of a PR associated with an event
- 5. r-lib/actions/pr-push Pushes changes to a PR associated with an event
- 6. r-lib/actions/run-rchk Runs rchk tests to detect memory protection errors in C source code.

Examples

See the r-lib/actions/examples directory for a variety of example workflows using these actions.

14/1 . (* 11 1

Configuring Travis

In .travis.yml, add the following lines:

```
os:
   - linux
   - osx

warnings_are_errors: true
after_success:
   - Rscript -e 'covr::codecov(type = "all")'
```

Configuring Appveyor

In appveyor.yml, add the following lines:

```
environment:
   global:
    WARNINGS_ARE_ERRORS: 1
```

Adding to the README. Rmd

Add the following lines, changing GITHUB_USERNAME/REPO to the correct version

```
[![Travis-CI Build Status](https://travis-ci.com/GITHUB_USERNAME/REPO.svg?branch=master)](https://ci.appveyor.com/api/projects/status/github/GITHUB_USERNAME/REPO.svg?branch=master)]
```

or can use:

```
usethis::use_badge("Travis-CI Build Status",
src = "https://travis-ci.com/GITHUB_USERNAME/REPO.svg?branch=master",
href = "https://travis-ci.com/GITHUB_USERNAME/REPO")
```

to the README. Rmd.

S3, S4, Reference Classes

- S3 simple just say class (x) = "myS3Class"
 - Usually List of objects
 - http://adv-r.had.co.nz/S3.html
- · S4 more complex see new
 - name, representation (slots), and inheritance (does it act like an array/list)
 - http://adv-r.had.co.nz/S4.html
- Reference Classes
 - very different, class\$method()
 - http://adv-r.had.co.nz/R5.html

S3, S4 Methods

```
· S3:bar <- function(y) UseMethod("bar", y)

- bar.myS3Class will allow you to use bar(x)

· S4:setGeneric("myGeneric", function(x)
    standardGeneric("myGeneric"))

setMethod("myGeneric", signature(x = "myS3Class"), function(x, y) {
    x@slot + y
})
</pre>
```

Compiled Code: C and C++

See http://r-pkgs.had.co.nz/src.html

- The src/ folder has compiled code.
- cleanup generally deletes intermediate or downloaded files run in configure.
- · configure runs code before make is run
- Makevars or a Makefile gives direction for compiling code
- There are configure.win and Makevars.win for Windows-specific setup.
- See.call for calling these compiled functions.

Using Rcpp is a different framework.

Questions?