





Fundamentals of Package Development

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Welcome!

- Instructors:
 - Andy Teucher
 - andyteucher.ca
 - GitHub: [ateucher](https://github.com/ateucher)
 - Mastodon: [@andyteucher@fosstodon.org](https://fosstodon.org/@andyteucher)
 - Sam Albers
 - samalbers.science
 - GitHub: [boshek](https://github.com/boshek)

Welcome!

- This is a three part course for people looking to learn how to build R packages in an efficient way, make them easy to maintain, and easy for users to use.
- Code of Conduct:
 - https://github.com/ateucher/pkg-dev-psc-2024-04-29/blob/main/CODE_OF_CONDUCT.md
 -  Treat everyone with respect
 -  Everyone should feel welcome

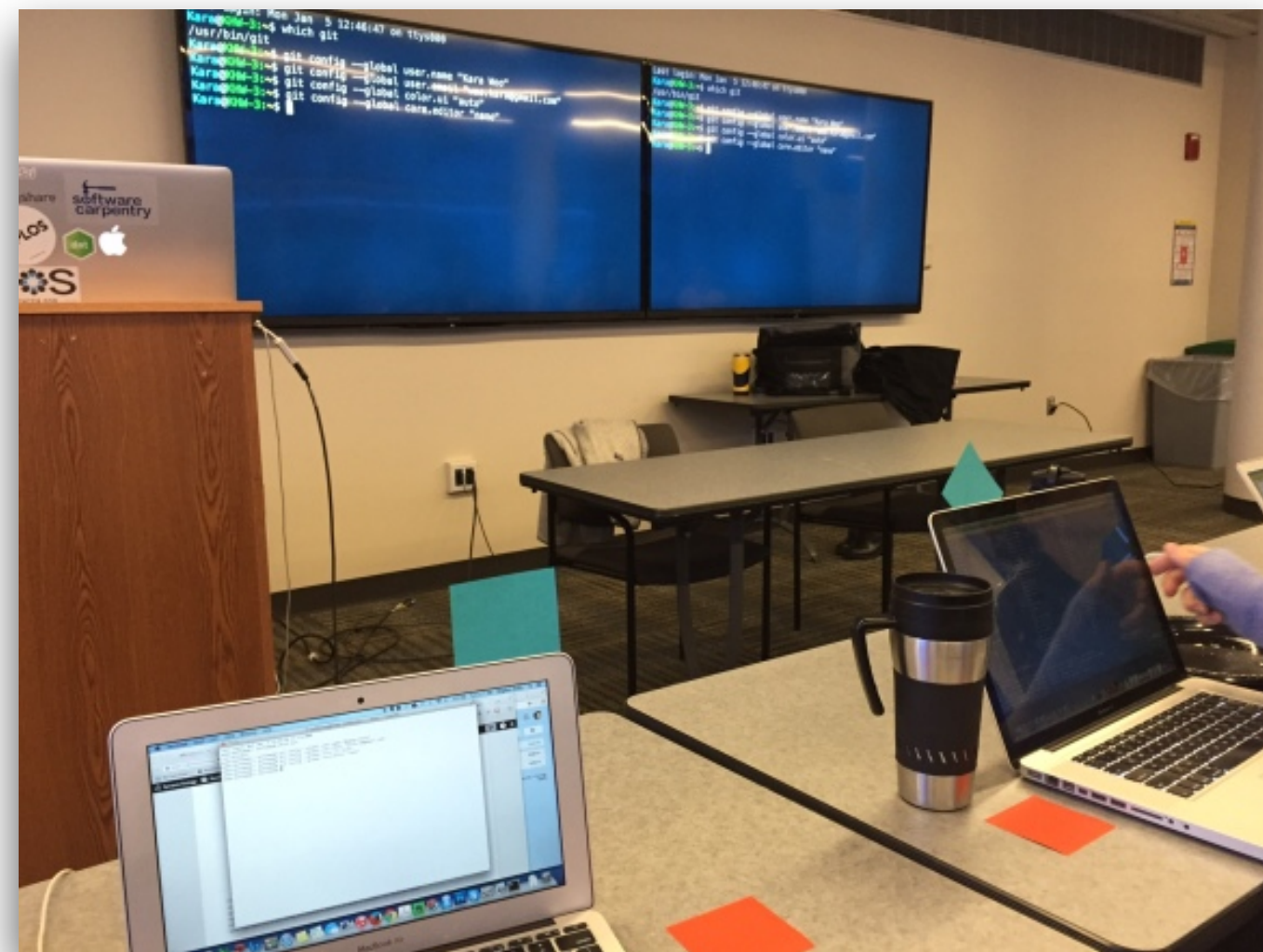
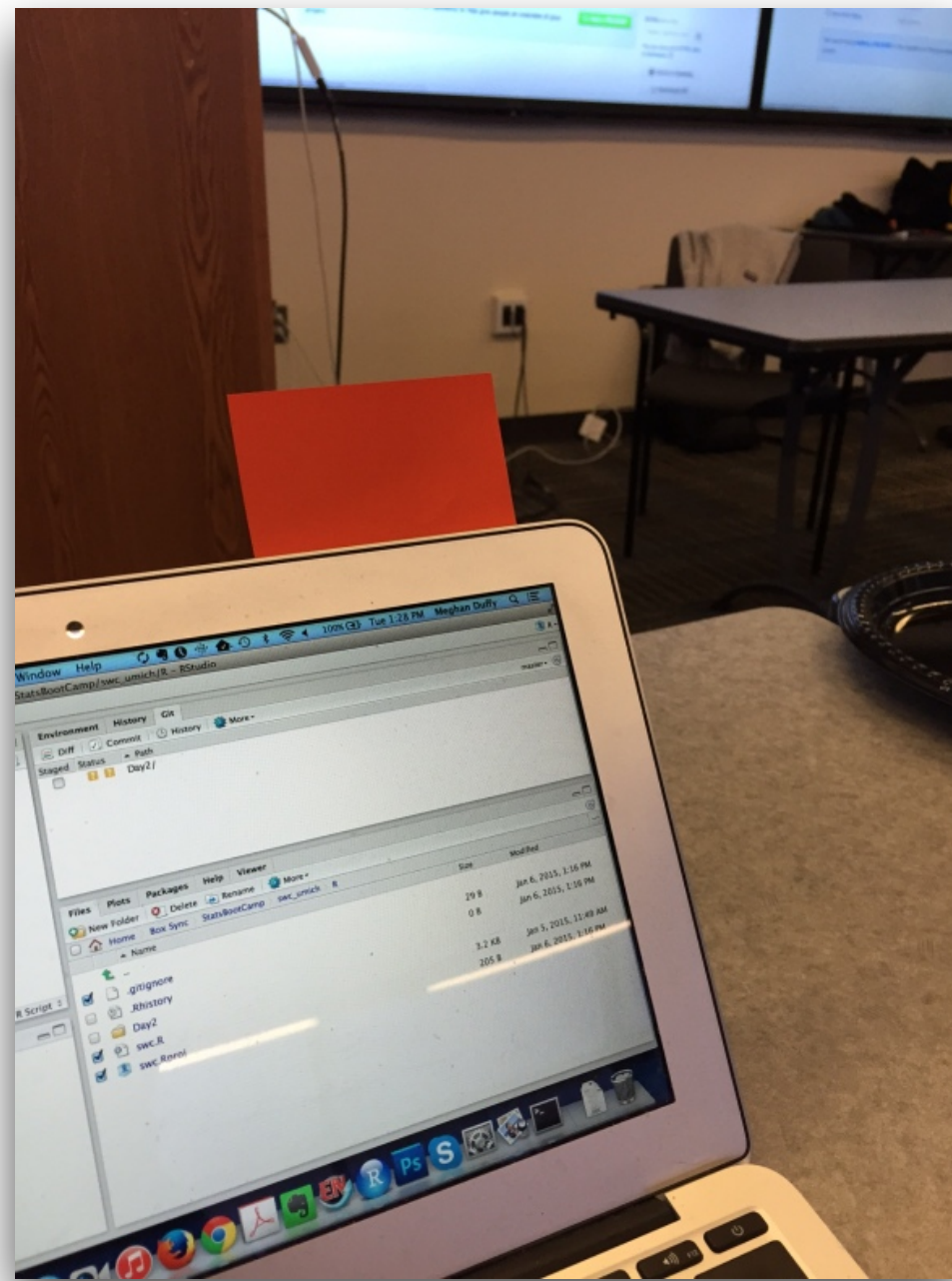
Sticky Notes

PINK

I'm stuck, please
send help

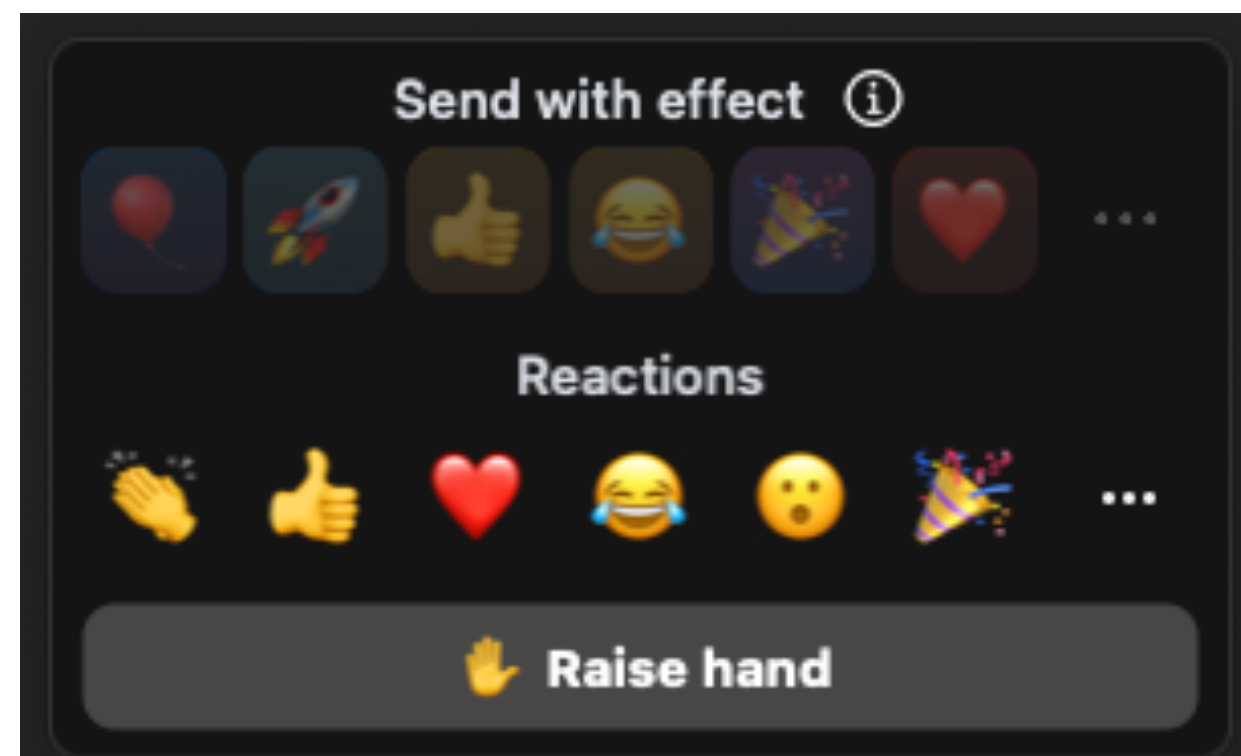
GREEN

I'm good, let's
move on

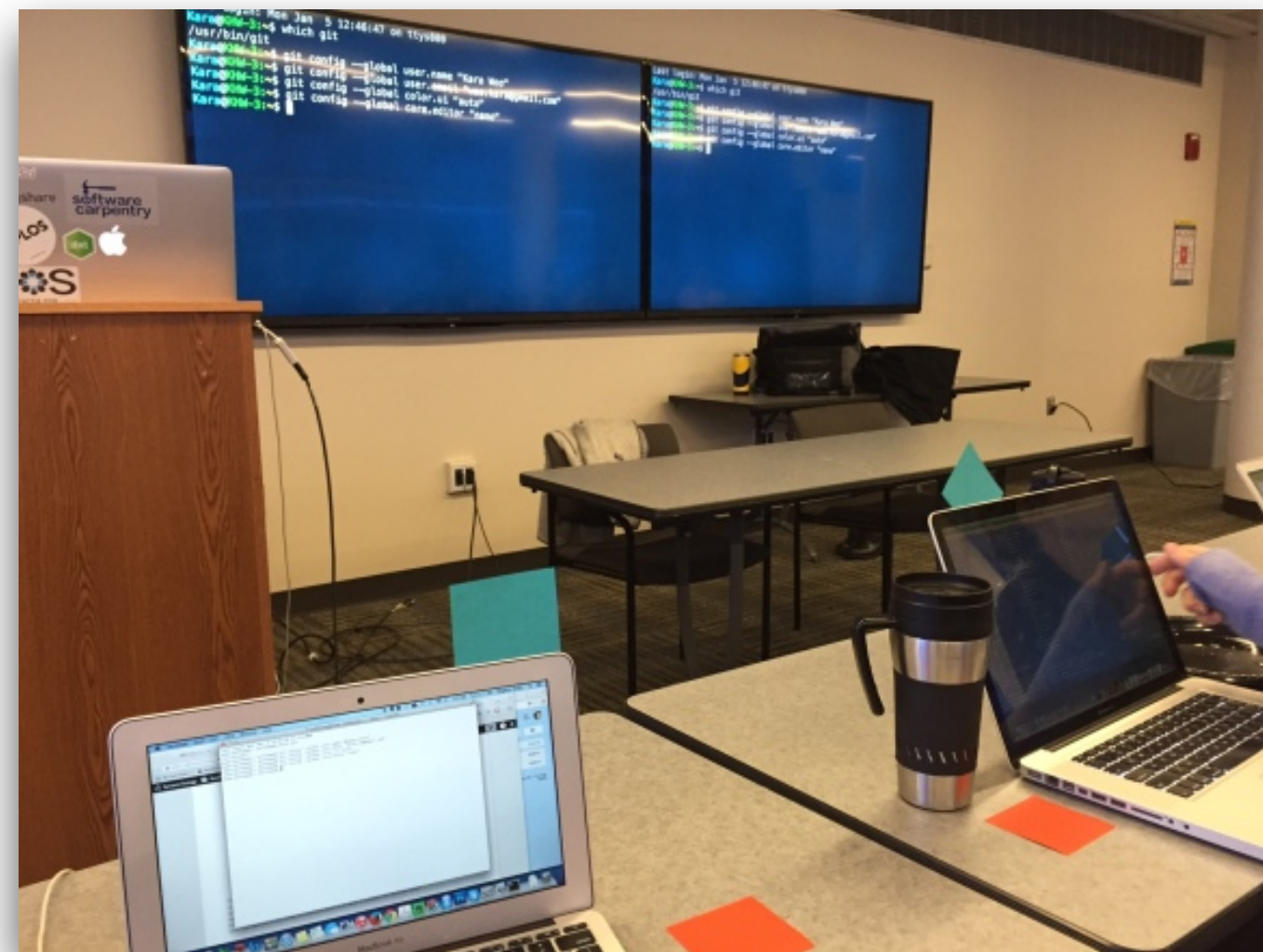
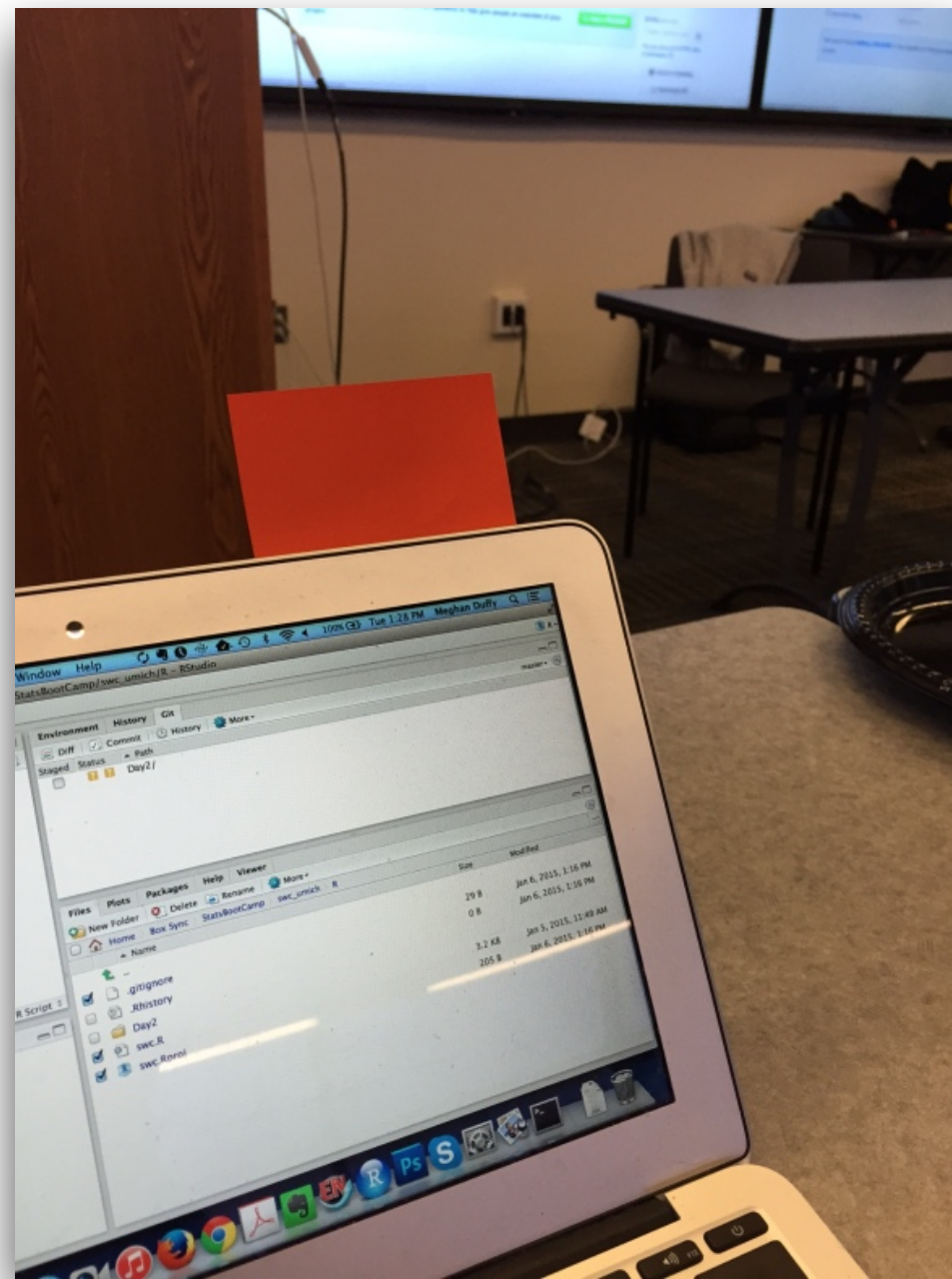


Sticky Notes

I'm stuck, please
send help



I'm good, let's
move on



Discussions / Q & A

- Zoom Chat

- GitHub Discussions:

<https://github.com/ateucher/pkg-dev-dfo-2024-11/discussions>

- Open the welcome discussion:
 - Introduce yourself:
 - Office/department, city/town
 - Why you're here
 - Share something you're good at

Resources

- Prework:

<https://andyteucher.ca/pkg-dev-dfo-2024-11/setup.html>

- Workshop website:

<https://andyteucher.ca/pkg-dev-dfo-2024-11/>

- Cheatsheet:

<https://rstudio.github.io/cheatsheets/html/package-development.html>

Day 1: Schedule and Learning Objectives

08:00 - 09:20	What is a Package? Package Structure and State	80 min
09:20 - 09:35	Break	15 min
09:35 - 10:55	Package Creation and Metadata	80 min
10:55 - 11:10	Break	15 min
11:10 - 12:30	Documentation	80 min

Day 2: Schedule and Learning Objectives

08:00 - 09:20	Testing & Dependencies	80 min
09:20 - 09:35	Break	15 min
09:35 - 10:55	Continuous Integration & Design Principles	80 min
10:55 - 11:10	Break	15 min
11:10 - 12:30	Design Principles cont'd & Website Creation	80 min

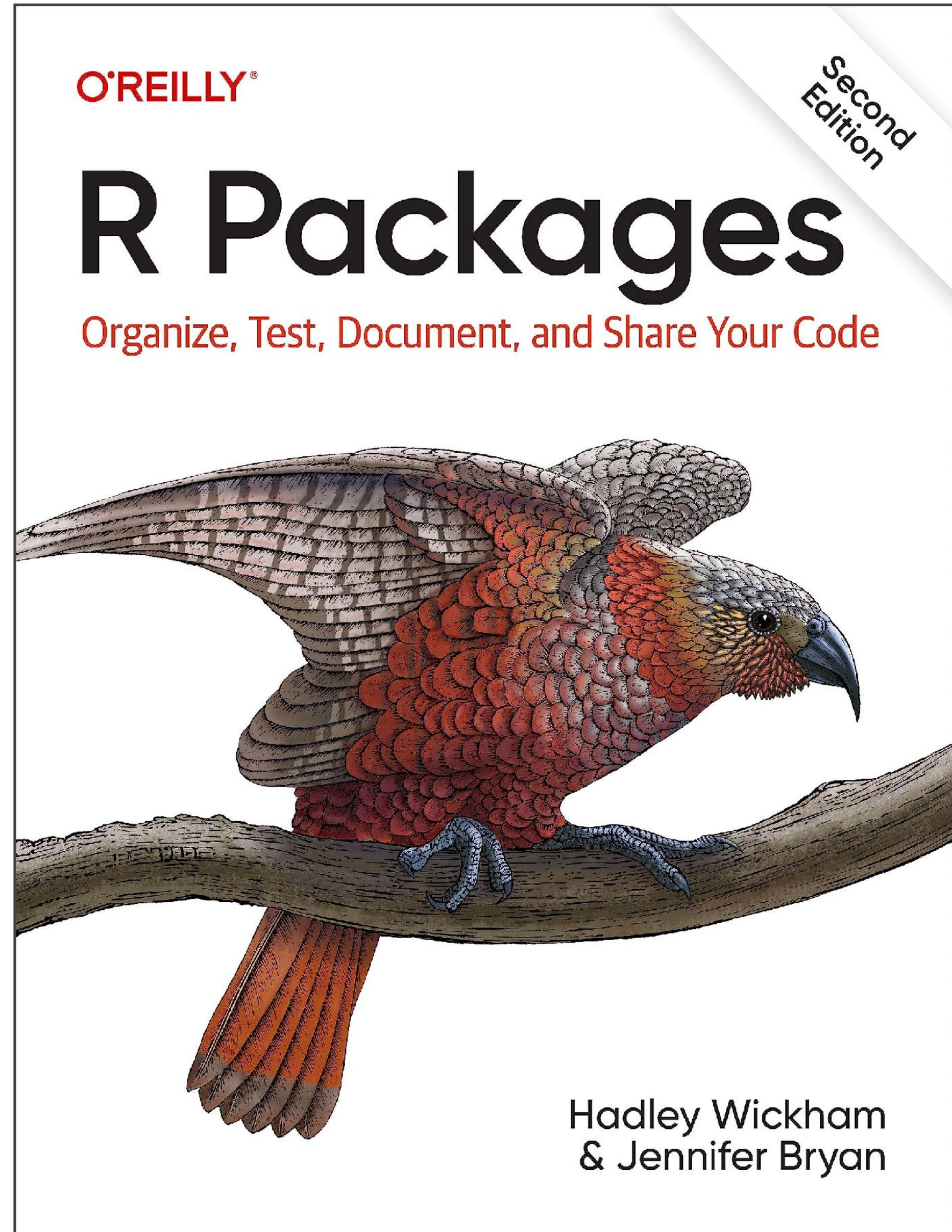
Day 3: Schedule and Learning Objectives

08:00 - 09:20	Using the Tidyverse in Your Package	80 min
09:20 - 09:35	Break	15 min
09:35 - 10:55	Communicating with your Users	80 min
10:55 - 11:10	Break	15 min
11:10 - 12:30	Distribution & General Package Discussion	80 min

R Packages (2e)

Hadley Wickham
Jenny Bryan

<https://r-pkgs.org>



Packages in a nutshell 🥜

Have you ever worked on a package before?

What do you feel the most confused or curious about?

 **Your Turn**

Why make a package?



- Easier to reuse functions you write
- A consistent framework which encourages you to better organize, document, and test your code
- This framework means you can use many standardized tools
- Easiest way to distribute code (and data)
 - To your team
 - To the world

Script vs Package

<https://r-pkgs.org/package-within.html>

Script

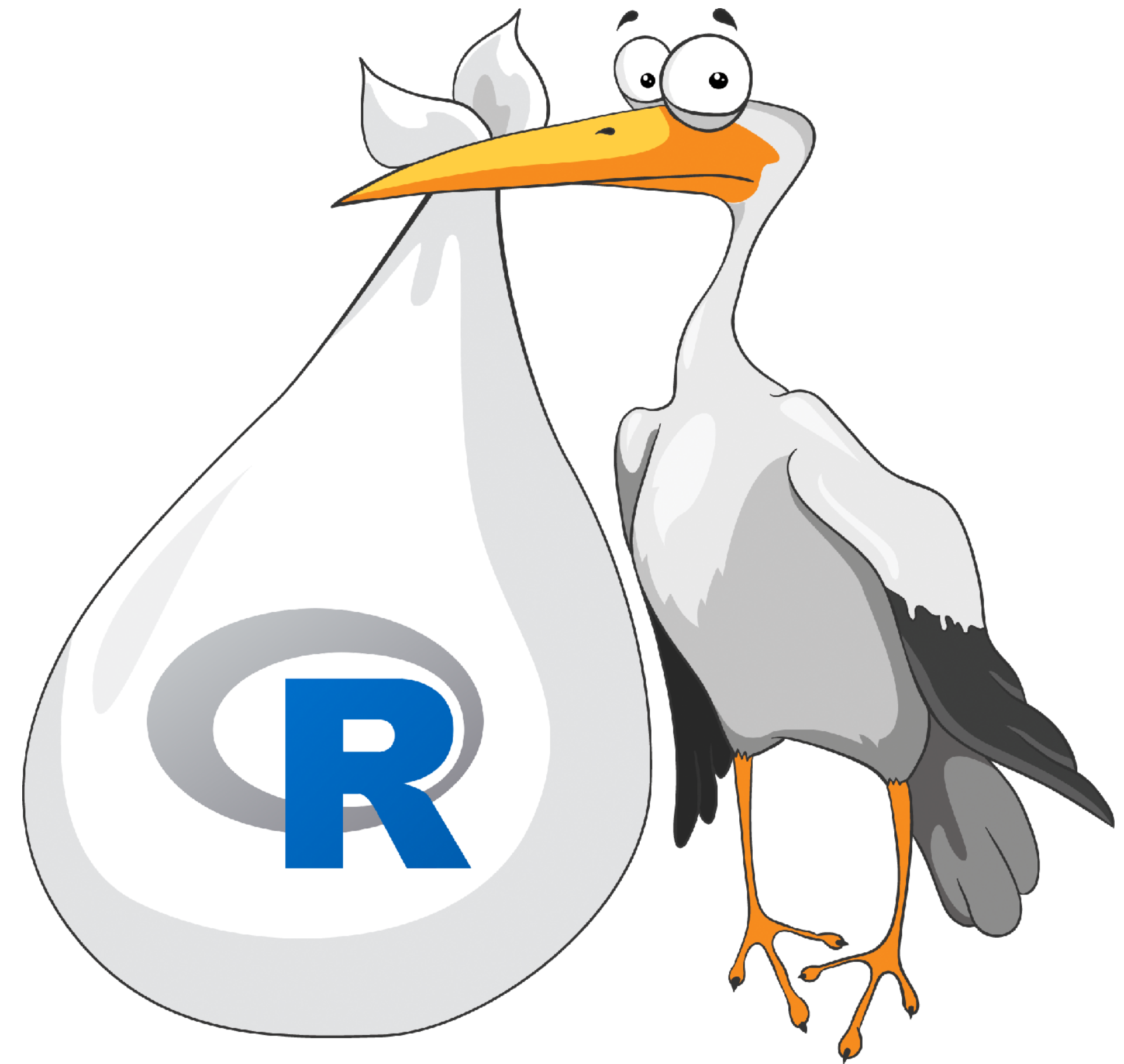
- Performs data analysis
- Collection of one or more **.R** files
- **library()** calls
- Documentation in # comments
- Run with **source()** or select+run

Package

- Reusable functions to use in analyses
- Defined by particular file organization
- Required packages in **DESCRIPTION**
- Documentation in **Roxygen** comments and “man” files
- Functions available when package attached

Where do packages come from?

- Discuss with your neighbour and put up a green sticky when you have:
 - Your favourite package
 - 2 **places** from which you install packages
 - 2 **functions** you can use to install packages
- Write them in the Chat



R Libraries - where do packages live?

- A **library** is a directory containing installed **packages**
- You have at least one library on your computer
- Common (and recommended) to have two libraries:
 1. A **system** library with **base** (14) and **recommended** (15) packages; installed with R.
 2. A **user** library with user-installed packages
- We use **library(pkg)** function to **attach** a package
- 7 base packages are always attached (**base, methods, utils, stats, grDevices, datasets, graphics**)

Your turn

Type `.libPaths()` to see your libraries

- How many libraries do you have?
- What are they? (Put them in the Chat)

 **Your Turn**

Package Structure and State

Five forms

Source

- Directory of files with specific structure
 - What you interact with as you build a package
-

Bundle

- Package compressed into a single file (tar.gz) via `devtools::build()` -> R CMD build
 - Vignettes are built and files listed in `.Rbuildignore` are left behind
-

Binary

- Platform-specific compressed file (.tgz, .zip)
 - Made with `devtools::build(binary = TRUE)` -> R CMD INSTALL --build
-

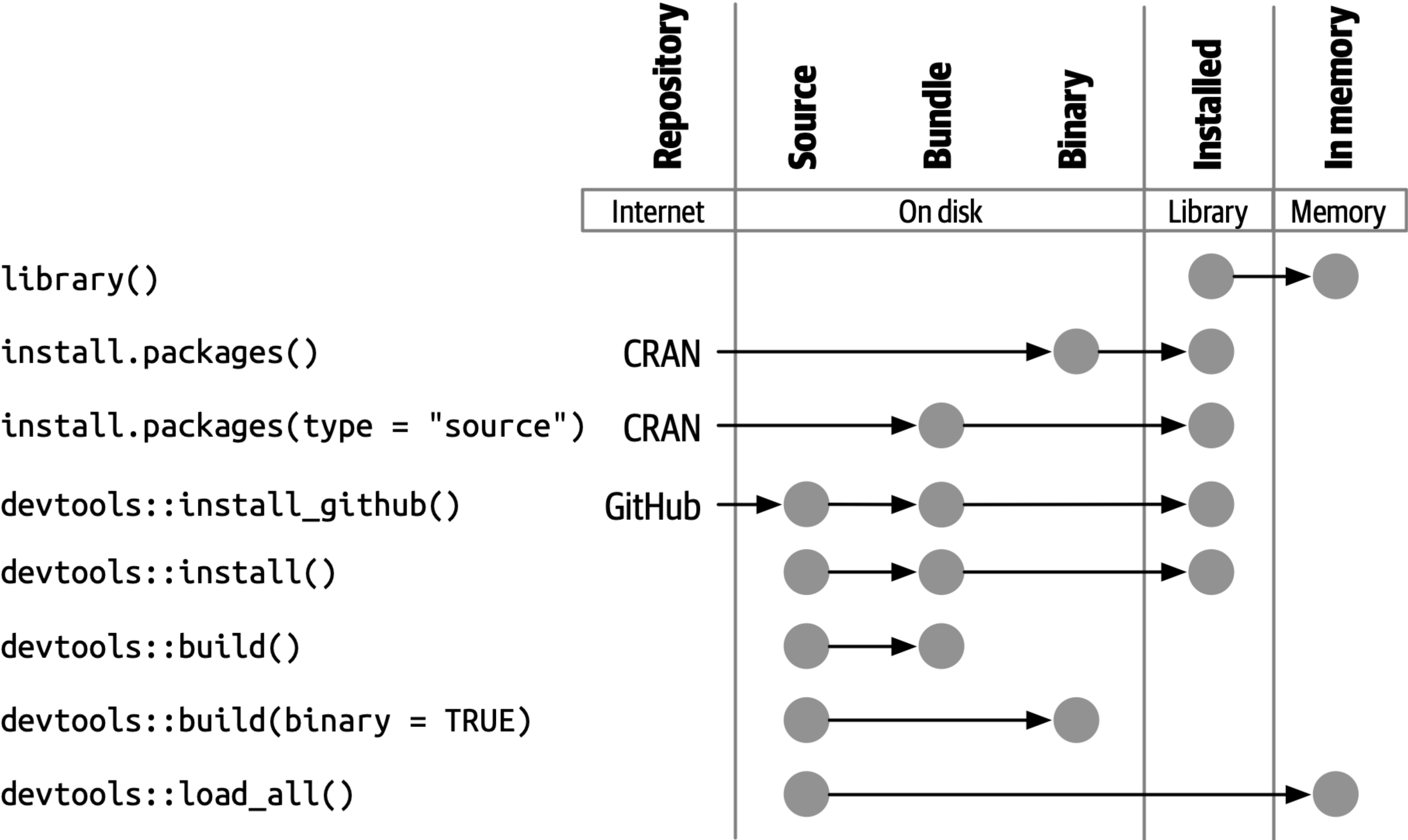
Installed

- Binary package decompressed into a user's library
 - `install.packages()`
-

In Memory

- Loaded and ready for use in an R session
- `library()`

Package Structure and State



Let's make a package together

We will:

- Create a simple package
- Use git to track our changes
- Push the code to a repository on GitHub
- Create tests for our functions
- Create documentation for our functions
- Create a package website
- Focus on workflows

We (probably) won't:

- Talk (much) about function writing
- Talk about how to include data in your package (even though it's possible and often helpful)

libminer

Sneak peak of our end goal on GitHub

- <https://github.com/ateucher/libminer.final>
- A package to explore our local R package libraries

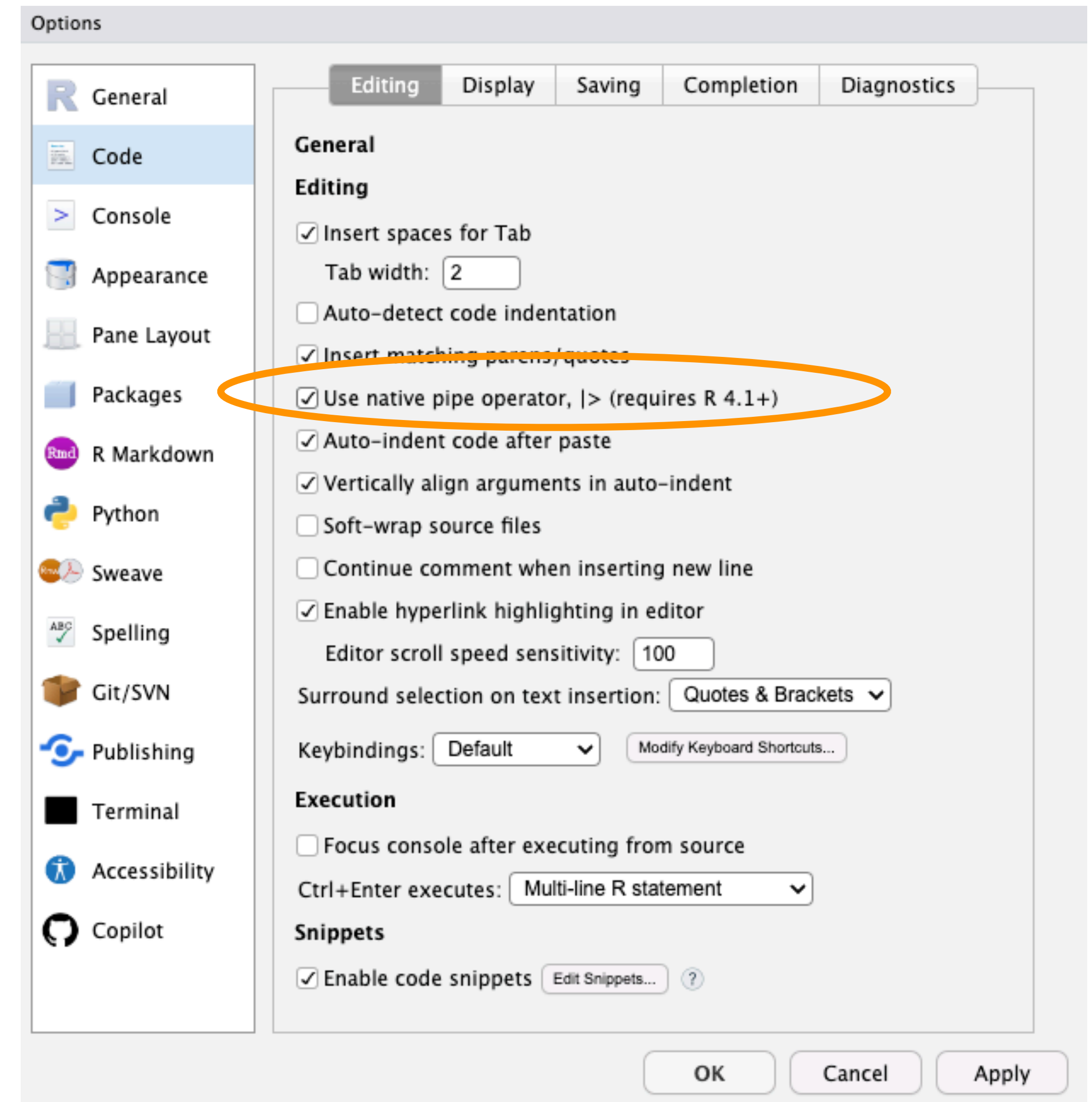
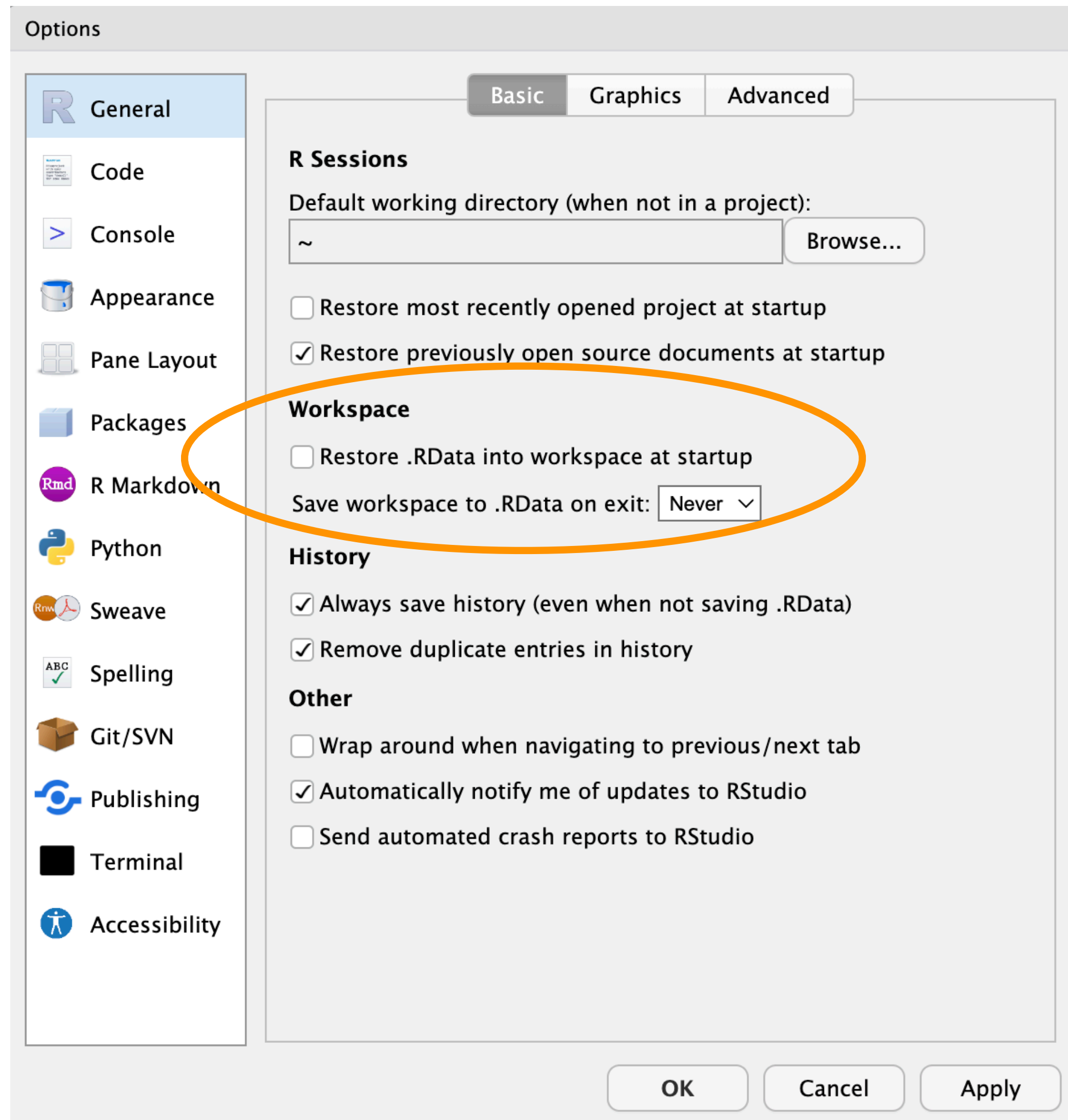


Get Ready



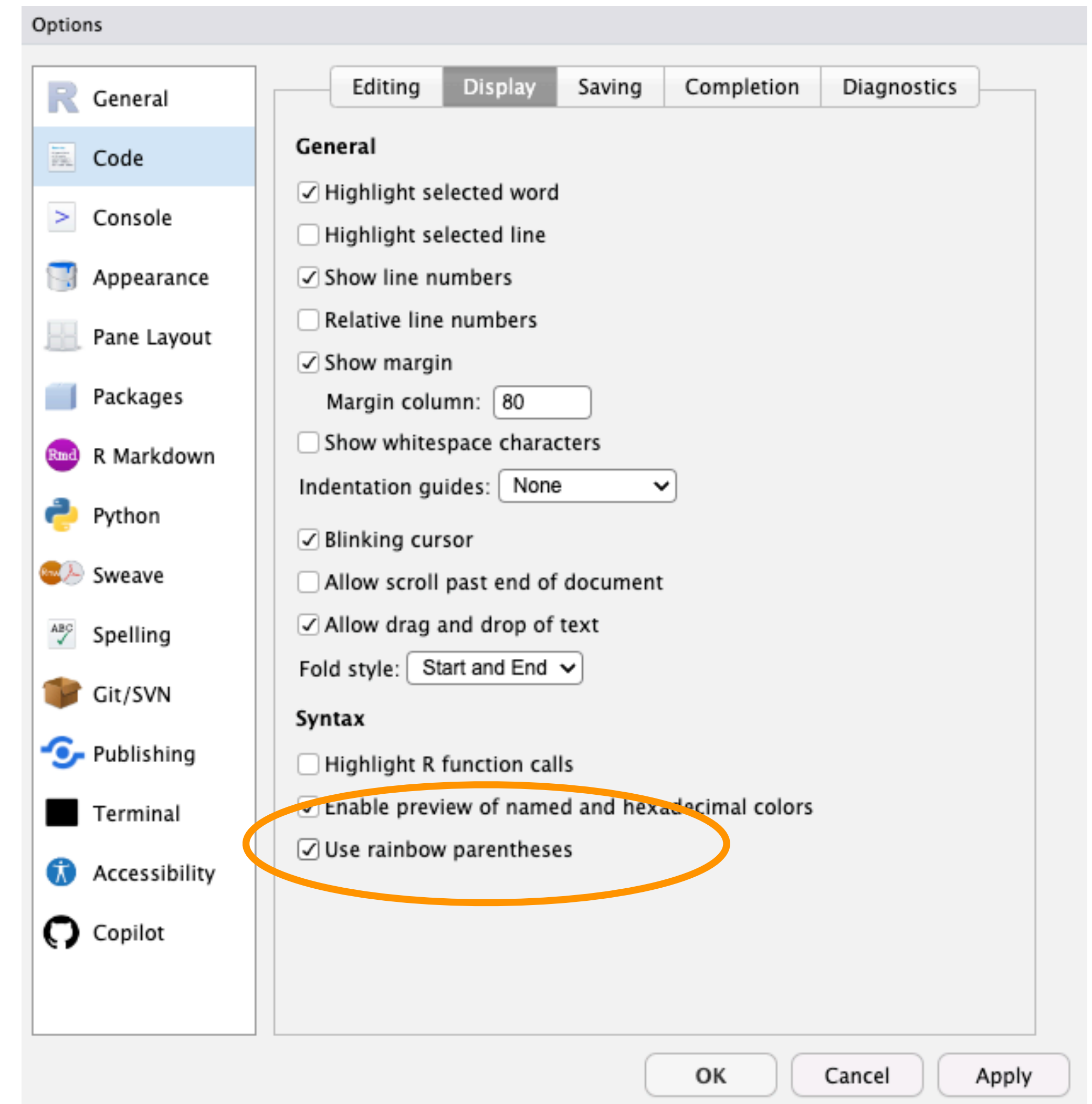
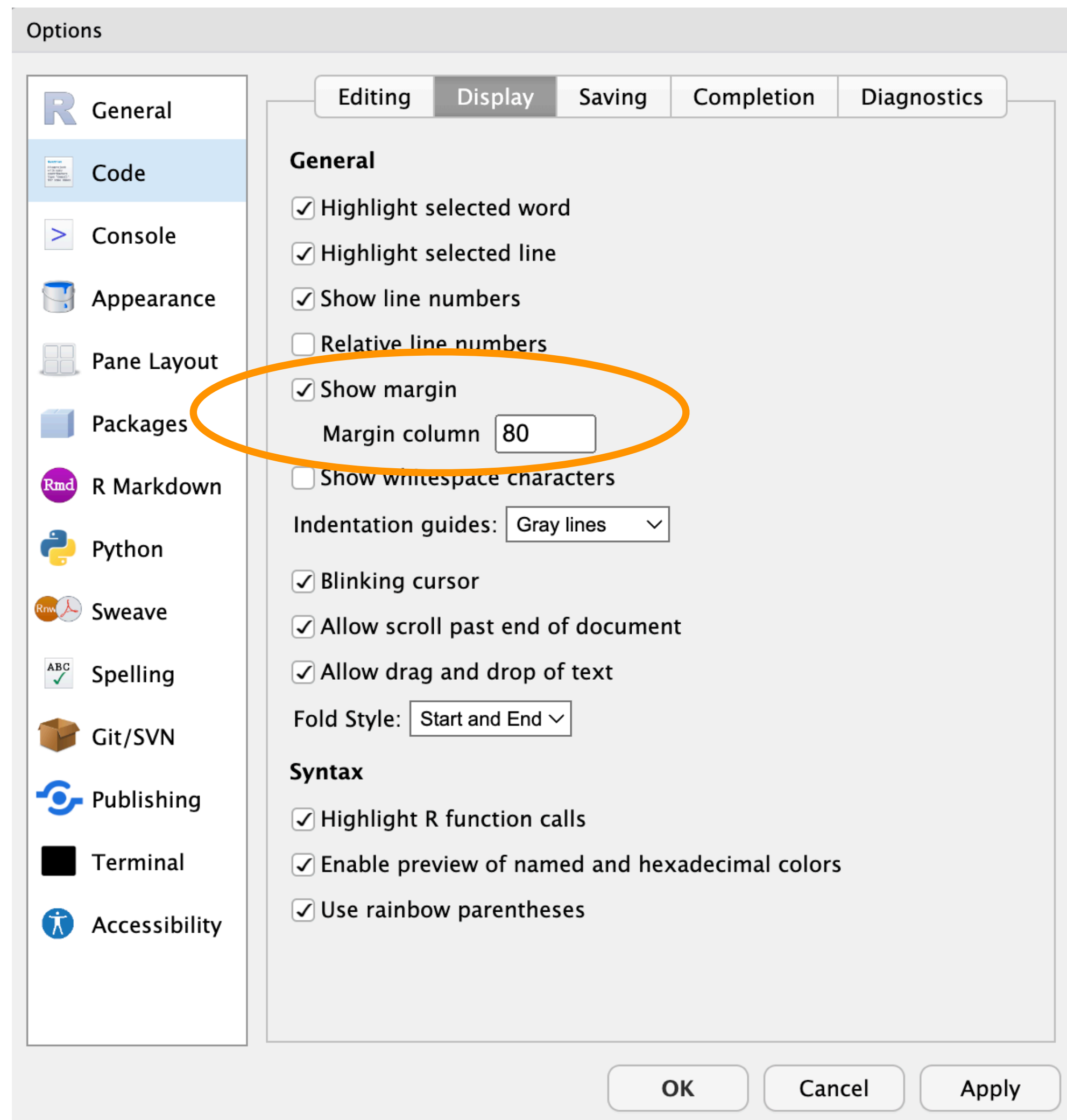
Configure RStudio

Tools > Global Options




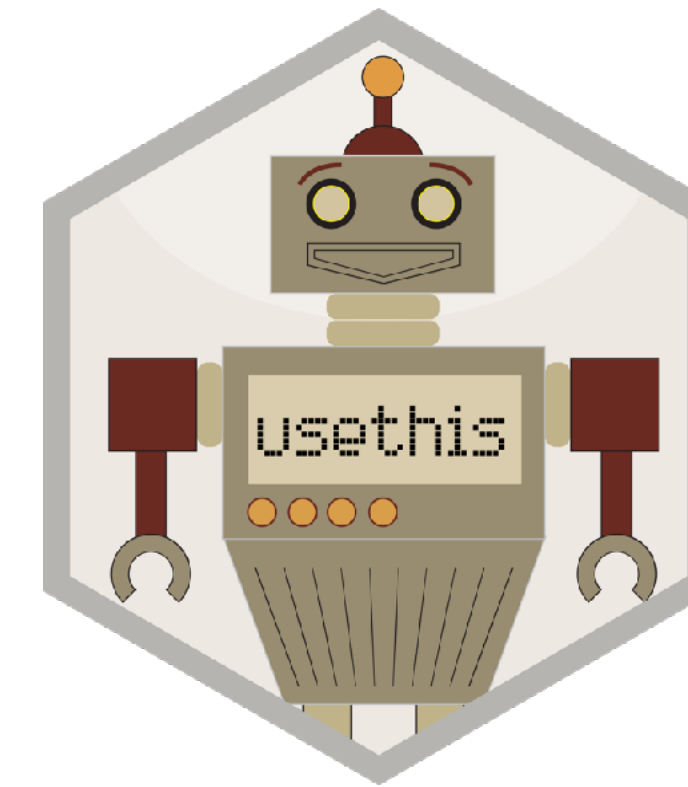
Configure RStudio

Tools > Global Options



Tools

- R >= 4.3.0
-  **Studio** (<https://posit.co/download/rstudio-desktop/>)
- Packages:



```
install.packages(  
  c("devtools", "roxygen2", "testthat", "knitr", "pkgdown")  
)
```



Check your setup

```
library(devtools)
# package development "situation report"
dev_sitrep()
# git/github "situation report"
git_sitrep()
```

 **Your Turn**

Break Time!



Create a package



Load devtools

```
library(devtools)
#> Loading required package: usethis

packageVersion("devtools")
#> [1] '2.4.5'
```

- Update if necessary!
- Provides a suite of functions to aid package development
- Loads **usethis**, the source of most functions we will be using

create_package()

```
create_package("~/Desktop/mypackage")
```

```
.Rbuildignore
```

```
.Rproj.user
```

```
.gitignore
```

```
DESCRIPTION
```

```
NAMESPACE
```

```
R
```

```
mypackage.Rproj
```

- Creates directory
 - Final part of path will be the package name
- Sets up basic package skeleton
- Opens a new RStudio project
- Activates "build" pane in RStudio

create_package()

```
create_package("~/Desktop/mypackage")
#> ✓ Creating '/Users/jane/Desktop/mypackage/'
#> ✓ Setting active project to '/Users/jane/Desktop/mypackage'
#> ✓ Creating 'R/'
#> ✓ Writing 'DESCRIPTION'
#> Package: mypackage
#> 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
#> ✓ Writing 'NAMESPACE'
#> ✓ Writing 'mypackage.Rproj'
#> ✓ Adding '^mypackage\\.Rproj$' to '.Rbuildignore'
#> ✓ Adding '.Rproj.user' to '.gitignore'
#> ✓ Adding '^\\.Rproj\\.user$' to '.Rbuildignore'
#> ✓ Setting active project to '<no active project>'
```

 **Your Turn**

use_git()

- `use_git_config(`
 - `user.name = "Jane Doe",`
 - `user.email = "jane@example.org"``)`
- `use_git()`
- Turns package directory into a git repository
- Commits your files (with a prompt)
- Restarts RStudio (with a prompt)
 - Activates "git" pane in RStudio

```
use_git()
#> ✓ Setting active project to
#>   '/Users/Jane/rrr/mypackage'
#> ✓ Adding '.Rhistory', '.Rdata',
#>   '.htrr-oauth', '.DS_Store',
#>   '.quarto' to '.gitignore'
#> There are 5 uncommitted files:
#> * '.gitignore'
#> * '.Rbuildignore'
#> * 'DESCRIPTION'
#> * 'metrify.Rproj'
#> * 'NAMESPACE'
#> Is it ok to commit these files?
#>
#> 1: Absolutely yes
#> 2: Not really
#> 3: Yeah, I don't know
```

 **Your Turn**

usethis::use_devtools()

Automatically load devtools when R starts

- Opens .Rprofile file
- Copies code to your clipboard
- Paste into .Rprofile
- Restart R

```
if (interactive()) {  
  # Load package dev packages:  
  suppressMessages(require("devtools"))  
}
```

 Ctrl+Shift+F10 (Windows &

 **Your Turn**

use_r()

Write your first function

- R code goes in **R/**
- Name the file after the function it defines

```
use_r("my-fun")
```

```
#> ✓ Setting active project to '/Users/jane/rrr/mypackage'
```

```
#> • Edit 'R/my-fun.R'
```

 **Your Turn**

Try your function in the new package

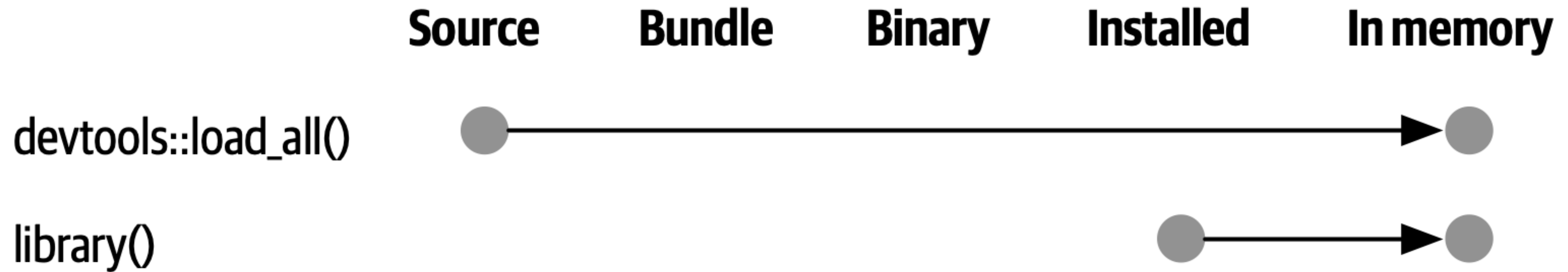
But how?

- `source("R/my-fun.R")`
- ~~Send function to console using RStudio (Ctrl/CMD+Return)~~
- `devtools::load_all()`

 Ctrl+Shift+L (Windows & Linux)

load_all()

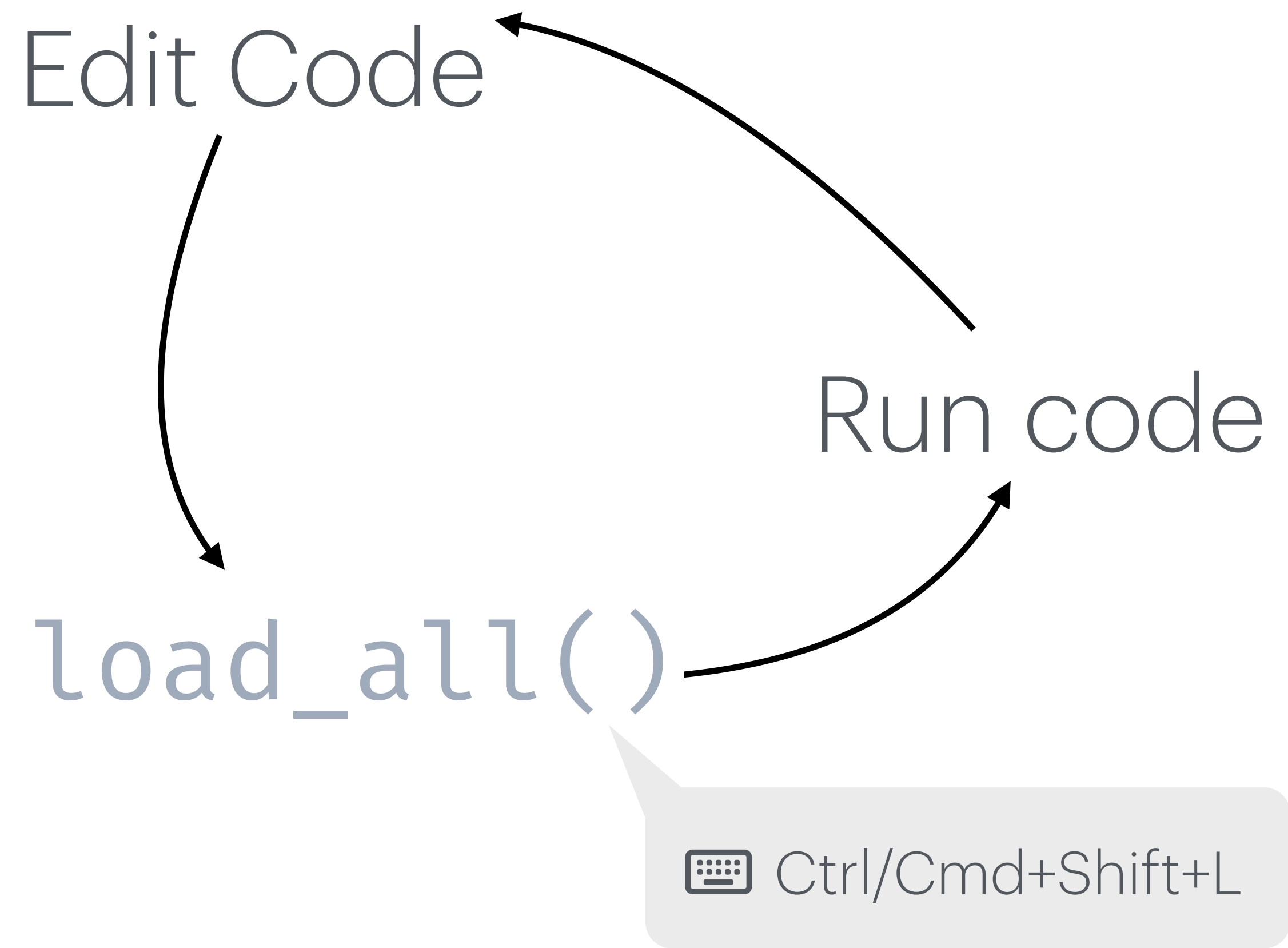
≈ **install.packages() + library()**



- Simulates building, installing, and attaching your package
- Makes all of the functions from your package immediately available to use
- Allows fast iteration of editing and test-driving your functions
- Good reflection of how users will interact with your package*

Try it out, and commit your
changes 

Workflow



check()

Run R CMD check from within R

```
check()

#> — R CMD check results —————
#> Duration: 3.1s
#>
#> > checking DESCRIPTION meta-information ... WARNING
#> Invalid license file pointers: LICENSE
#>
#> 0 errors ✓ | 1 warning ✘ | 0 notes ✓
```

- Reduce future pain by catching problems early*
- <https://github.com/wch/r-source/blob/97a5806aed3dceb174c/library/tools/R/check.R#L314>

 **Your Turn**

R CMD check

3 types of messages

- **ERRORs:** Severe problems - always fix.
- **WARNINGs:** Problems that you should fix, and must fix if you're planning to submit to CRAN.
- **NOTEs:** Mild problems or, in a few cases, just an observation.
 - When submitting to CRAN, try to eliminate all NOTEs.

Licenses

use*_license()

- Permissive:
 - **MIT:** simple and permissive.
 - **Apache 2.0:** MIT + provides patent protection.
- Copyleft:
 - Requires sharing of improvements.
 - **GPL (v2 or v3)**
 - **AGPL, LGPL** (v2.1 or v3)
- Creative commons licenses:
 - Appropriate for data packages.
 - **CC0:** dedicated to public domain.
 - **CC-BY:** Free to share and adapt, must give appropriate credit.

use_mit_license()

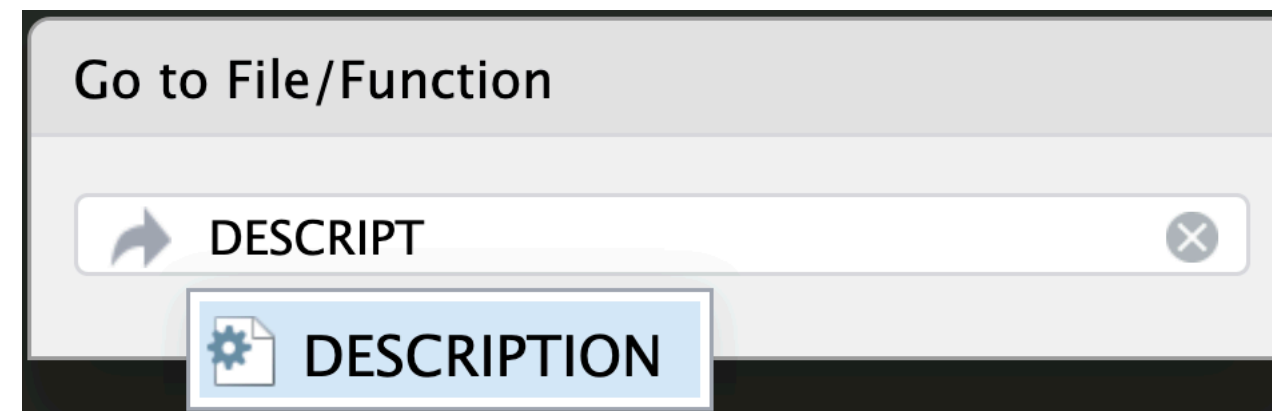
- ✓ Adding 'MIT + file LICENSE' to License
- ✓ Writing 'LICENSE'
- ✓ Writing 'LICENSE.md'
- ✓ Adding '^LICENSE\\.md\$' to '.Rbuildignore'

 **Your Turn**

The DESCRIPTION file

Package metadata

- Make yourself the author
 - Name & Email
 - Role
 - ORCID (optional)
- Write descriptive
 - **Title:**
 - **Description:**



 Ctrl+.

start typing DESCRIPTION

```
Package: mypackage
Title: What the Package Does (One Line, Title Case)
Version: 0.0.0.9000
Authors@R: person(
  "First", "Last", ,
  "first.last@example.com",
  role = c("aut", "cre"),
  comment = c(ORCID = "YOUR-ORCID-ID")
)
Description: What the package does
License: `use_mit_license()`, `use
  friends to pick a license
Encoding: UTF-8
Roxygen: list(markdown = TRUE)
RoxygenNote: 7.3.
```

 **Your Turn**

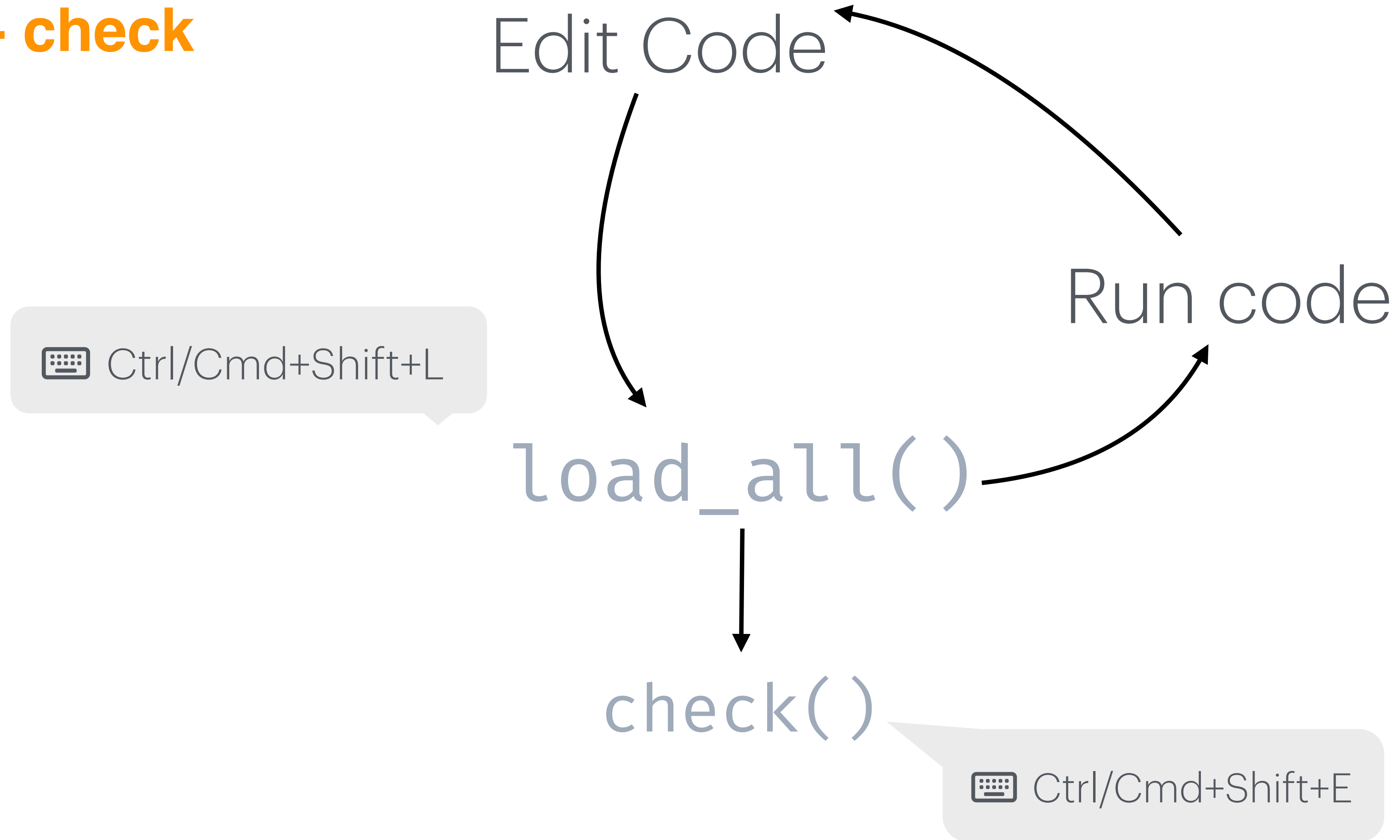
The DESCRIPTION file

Package metadata

- Take a look at the DESCRIPTION for ggplot2.
 - [CRAN Package page](#)
 - [DESCRIPTION on GitHub](#)
 - Note other Author roles:
 - 'cph' (copyright holder, often your employer)
 - 'fnd' (funder)

Workflow

Code + check



check() again

check()

#> *Documenting*

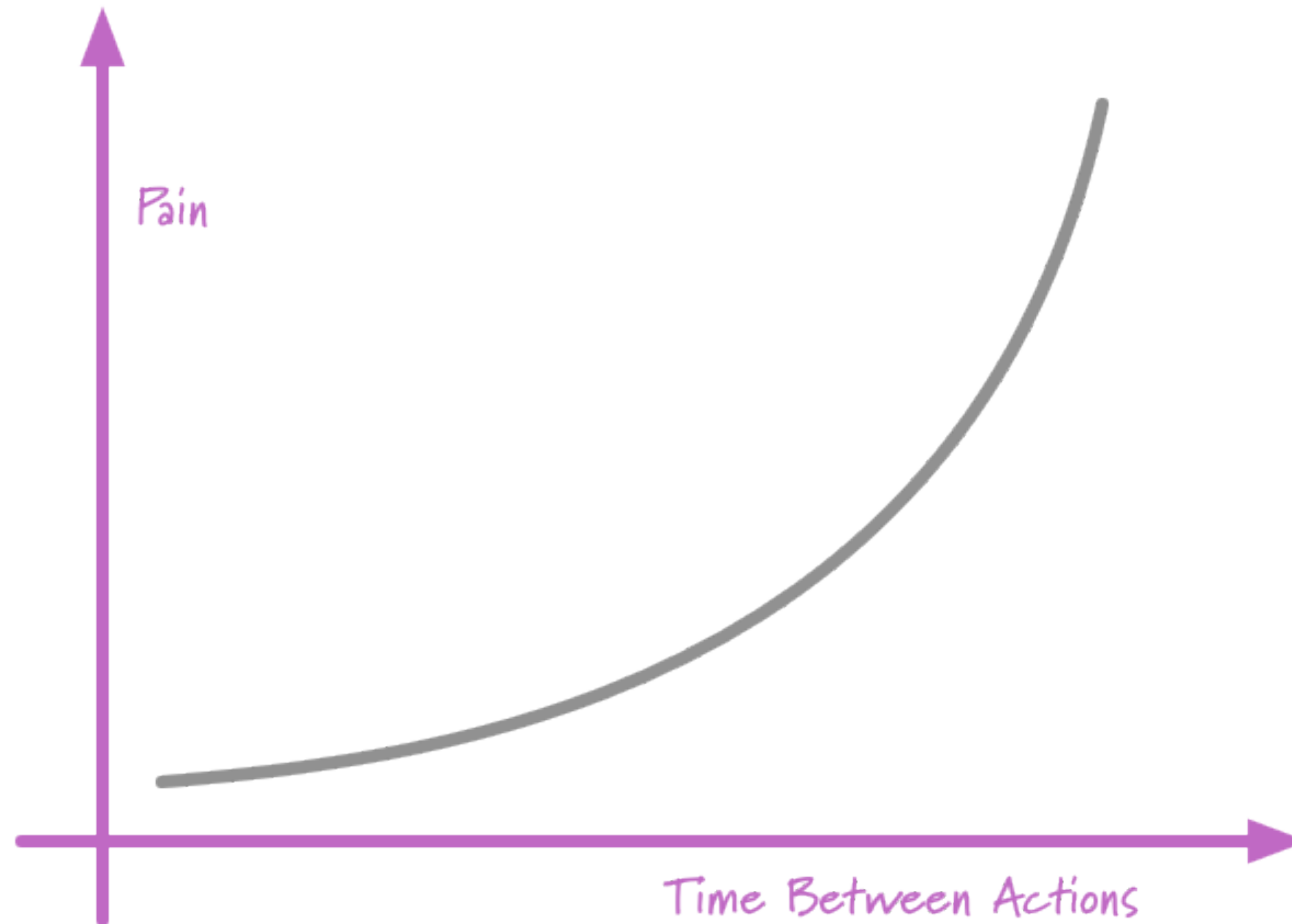
...
#> *Building*

...
#> *Checking*





"If it hurts, do it more often."



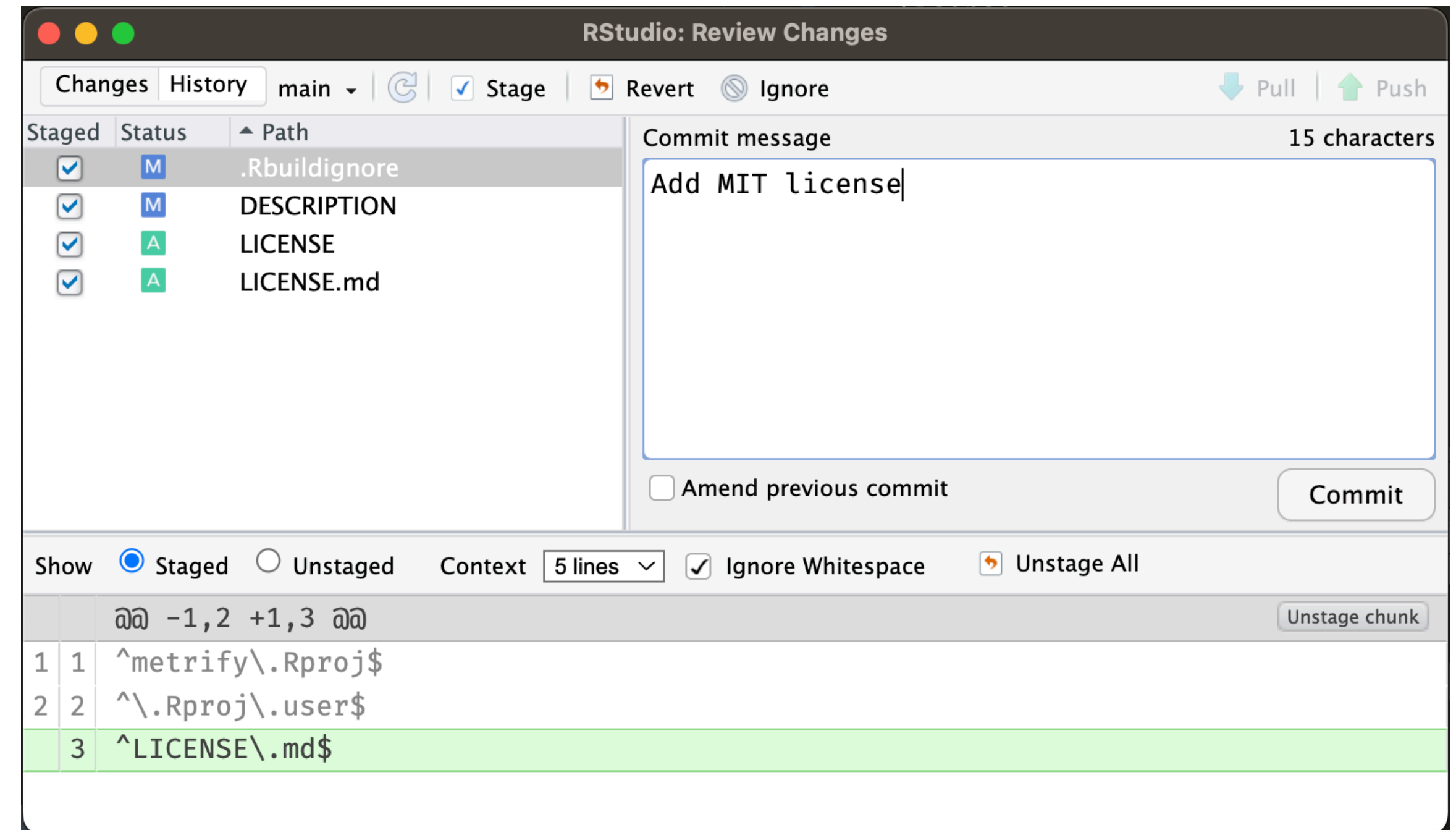
"If it hurts, do it more often."

- If `check()` goes from passing to failing after you've tinkered with one function or touched 10 lines of code, it's clear what to troubleshoot.
- If you've added 15 functions that all call each other and touched 100s of lines of code, bless your heart.
- Better to learn of a regrettable choice before you've built upon it for days or weeks.

Commit changes to git

```
$ git add DESCRIPTION \  
  LICENSE \  
  LICENSE.md \  
  .Rbuildignore
```

```
$ git commit -m "Add MIT license"
```



GitHub

Put your package code on GitHub

- Prerequisites:
 - GitHub account
 - Followed instructions to get GitHub PAT
 - <https://andyteucher.ca/pkg-dev-dfo-2024-11/setup.html#github>
 - Verify
 - `git_sitrep()`
 - Some things really are magic:
 - `use_github()`

 **Your Turn**

Avoid some pain of package setup: `edit_r_profile()`

And set default **DESCRIPTION** values

```
# Set usethis options:
options(
  usethis.description = list(
    "Authors@R" = utils::person(
      "Jane", "Doe",
      email = "jane@example.com",
      role = c("aut", "cre"),
      comment = c(ORCID = "0000-1111-2222-3333")
    )
  )
)
```

*<https://usethis.r-lib.org/articles/usethis-setup.html>

While you're in there...

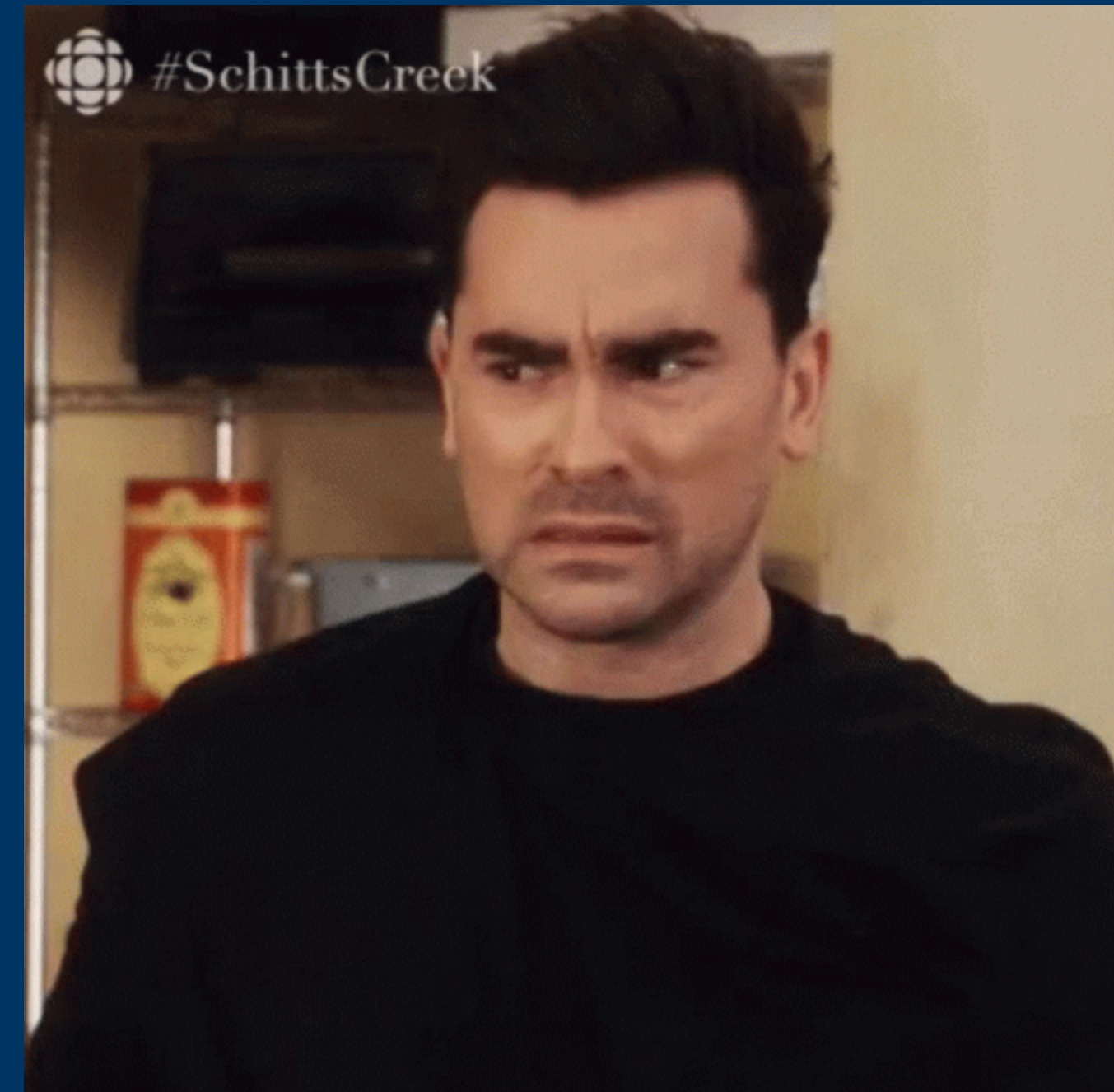
Set some other helpful defaults

```
options(  
  warnPartialMatchArgs = TRUE,  
  warnPartialMatchDollar = TRUE,  
  warnPartialMatchAttr = TRUE  
)
```

Documentation



Documentation

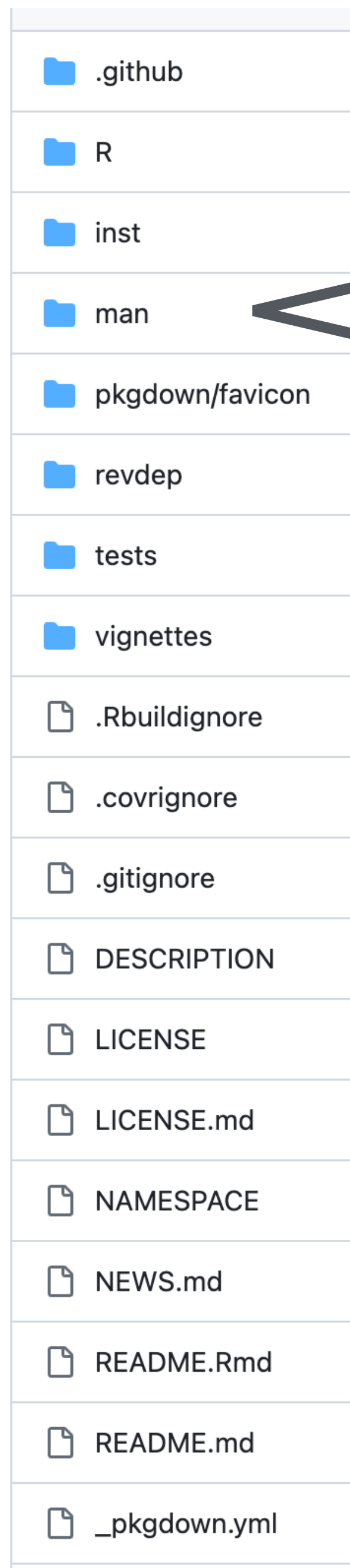




roxygen2

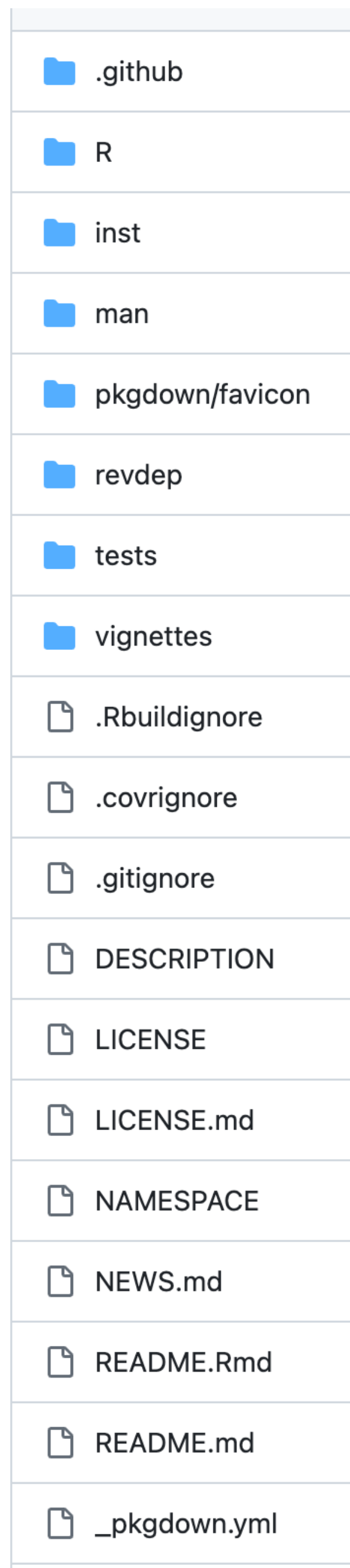


knitr



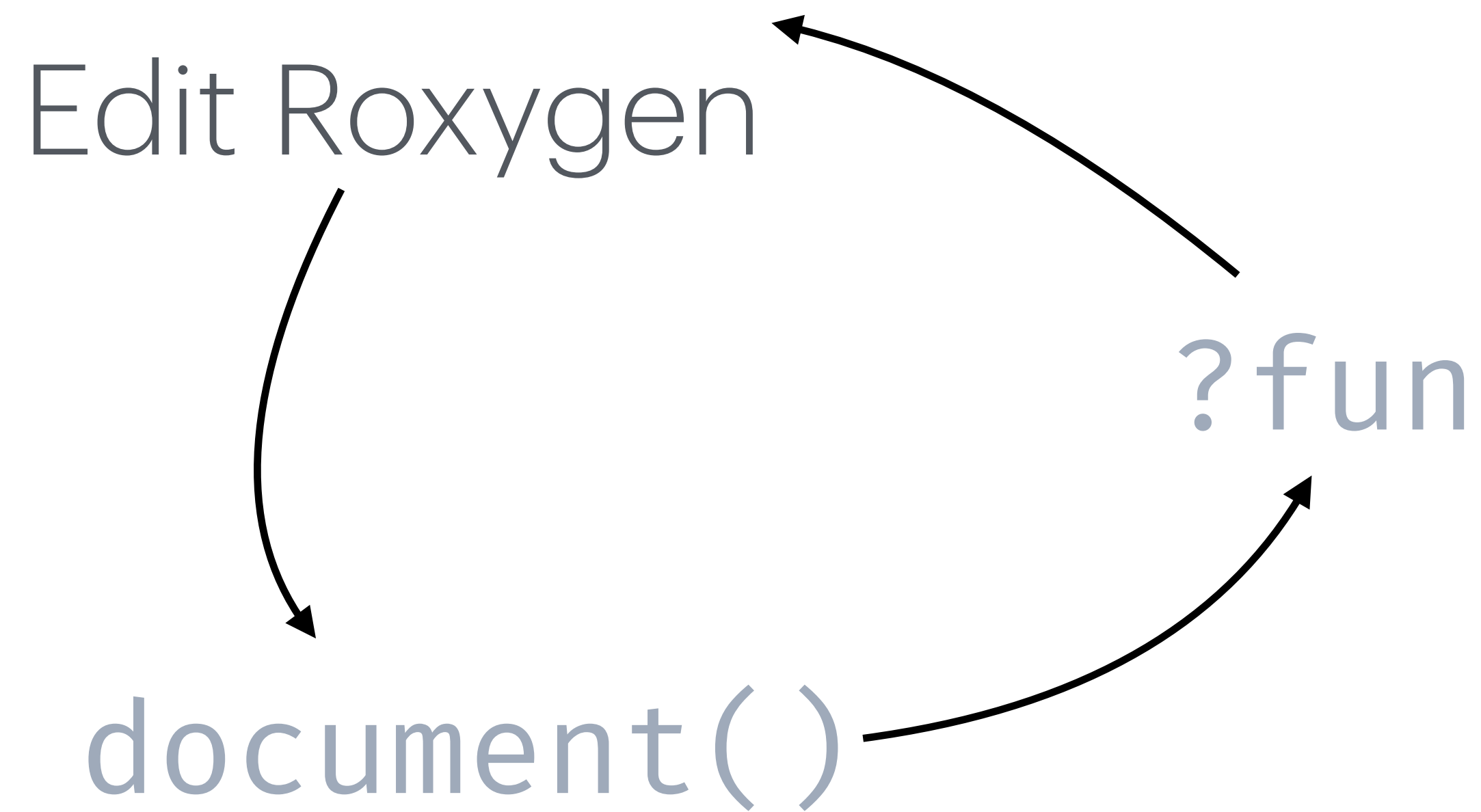
Help topics live here, in **.Rd** files

But you (mostly) don't write to **man/** with your bare hands.



Help topics are generated from the roxygen comments you write alongside your functions (or other objects), inside **R/**.

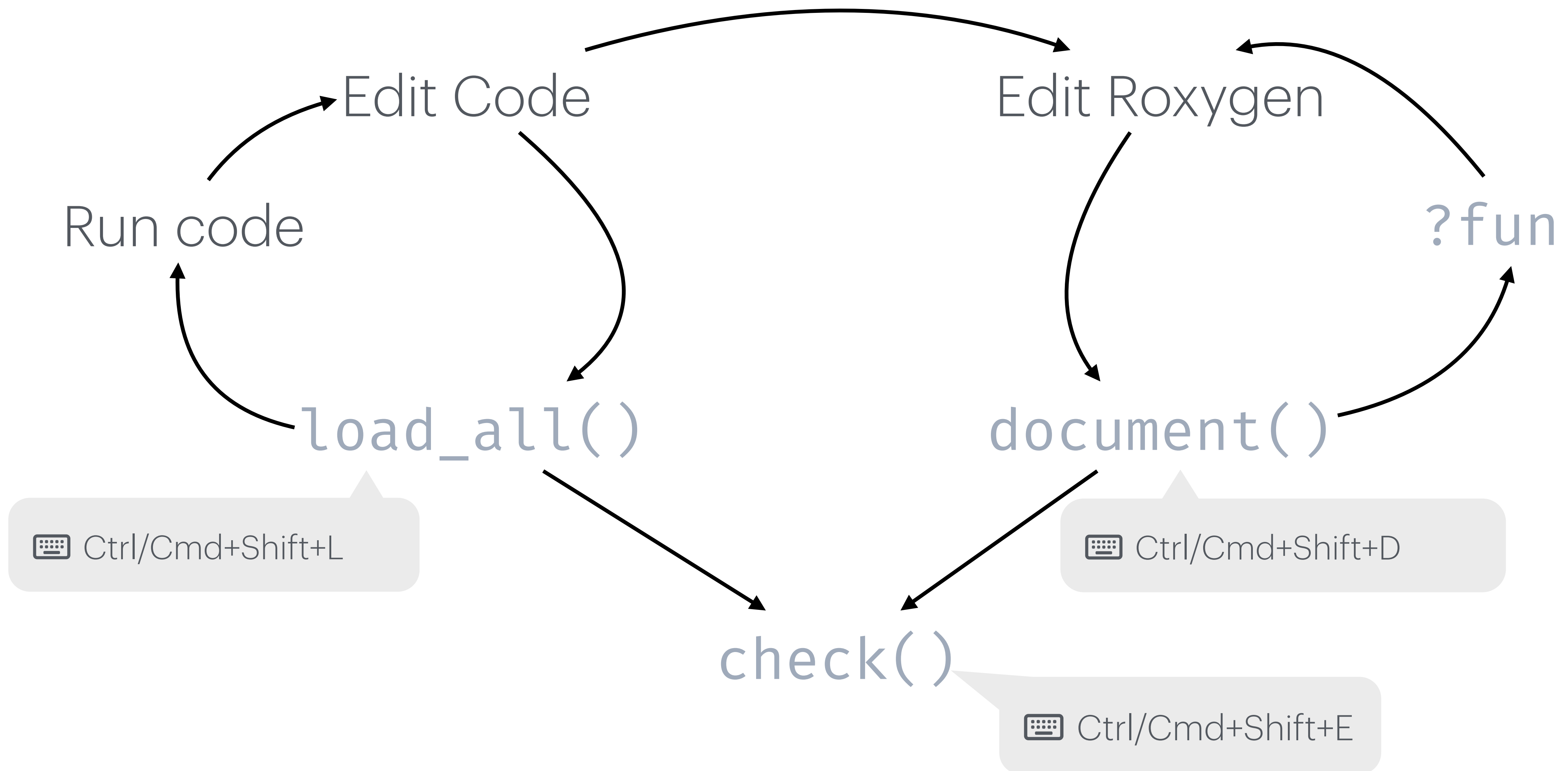
Documentation workflow

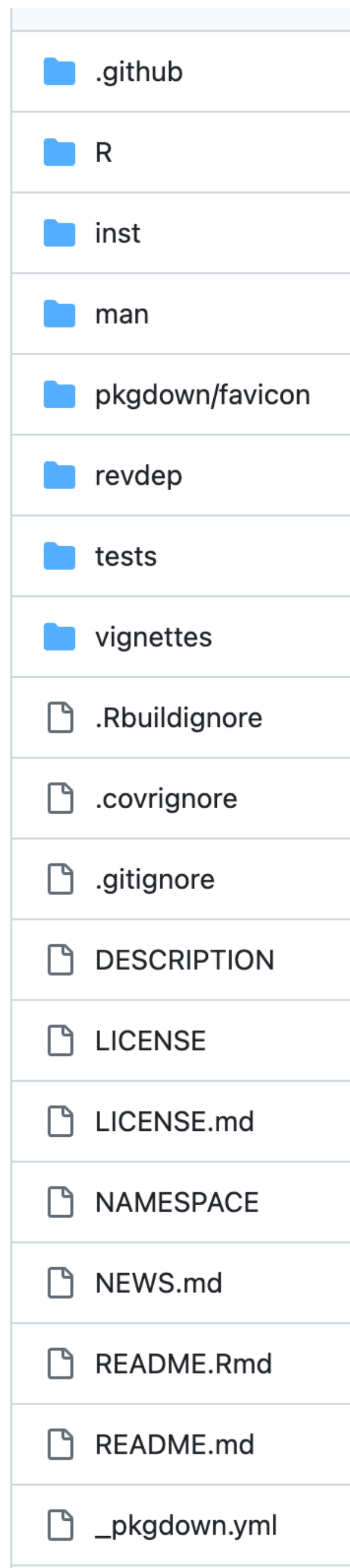


 Ctrl+Shift+D (Windows & Linux)

Workflow

Code + documentation + check



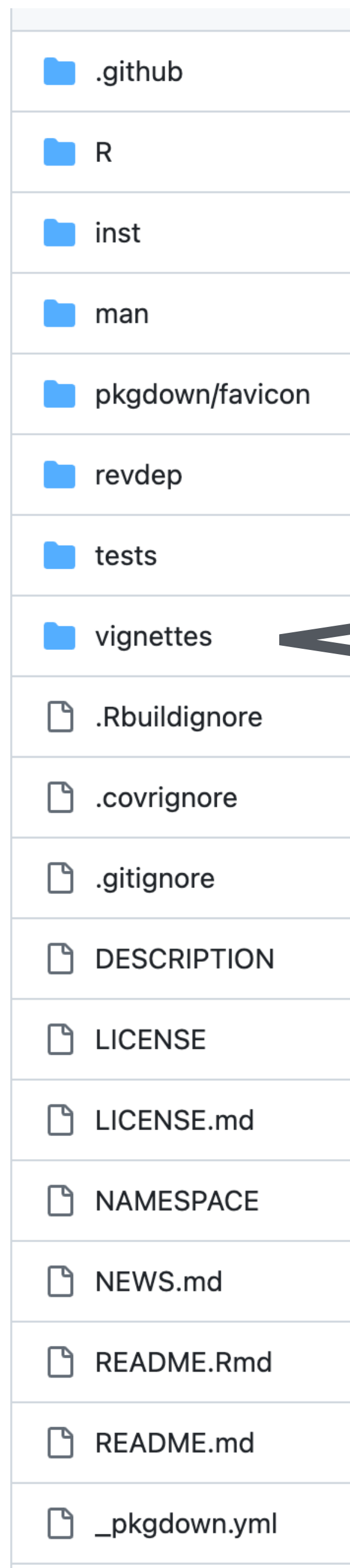


`README.md` is important:

- on CRAN
- on GitHub
- in the pkgdown site (it's the default `index.html`)

Ideally, `README.md` is generated from `README.Rmd`.

`build_readme()` is the best way to convert `README.Rmd` to `README.md`, because it actually installs the current dev package prior to rendering.



Vignettes are a great way to show truly authentic usage, combining multiple functions in your package to accomplish a realistic task.

If authentic usage involves moves that are impossible on CRAN, demonstrate that in an **article** instead. An article appears only in the pkgdown site.

- 📁 .github
- 📁 R
- 📁 inst
- 📁 man
- 📁 pkgdown/favicon
- 📁 revdep
- 📁 tests
- 📁 vignettes
- 📄 .Rbuildignore
- 📄 .covrignore
- 📄 .gitignore
- 📄 DESCRIPTION
- 📄 LICENSE
- 📄 LICENSE.md
- 📄 NAMESPACE
- 📄 NEWS.md
- 📄 README.Rmd
- 📄 README.md
- 📄 _pkgdown.yml

A user's experience with the pkgdown website is top of my mind when writing docs.

usethis 3.0.0 Setup Reference Articles ▾ News ▾

Search for

usethis

usethis is a workflow package: it automates repetitive tasks that arise during project setup and development, both for R packages and non-package projects.

Installation

Install the released version of usethis from CRAN:

```
install.packages("usethis")
```

LINKS

- [View on CRAN](#)
- [Browse source code](#)
- [Report a bug](#)

LICENSE

- [Full license](#)
- [MIT](#) + file [LICENSE](#)

COMMUNITY

- [Contributing guide](#)
- [Code of conduct](#)
- [Getting help](#)

CITATION

Items Needed for a help topic

- Insert a roxygen skeleton in RStudio with *Code > Insert Roxygen Skeleton*. Or do the equivalent "by hand".
- Complete the prepared fields
 - (Usually implicit) title and description
 - `@param` for function arguments
 - `@returns` for the return value
 - `@examples` for usage
 - `@export` to put in NAMESPACE
- Related workflow happiness: Cmd/Ctrl + Enter helps you develop your examples. Plays nicely with `load_all()` for staying synced with package source.

Typical roxygen comment

```
#' Remove duplicated strings
#'
#' `str_unique()` removes duplicated strings with additional control over
#' how duplication is measured.
#'
#' @param string Input vector. Either a character vector, or something
#'   coercible to one.
#' @param ... Other options used to control matching behavior between duplicate
#'   strings. Passed on to [stringi::stri_opts_collator()].
#' @returns A character vector, usually shorter than `string`.
#' @seealso [unique()], [stringi::stri_unique()] which this function wraps.
#' @examples
#' str_unique(c("a", "b", "c", "b", "a"))
#'
#' # Use ... to pass additional arguments to stri_unique()
#' str_unique(c("motley", "mötley", "pinguino", "pingüino"))
#' str_unique(c("motley", "mötley", "pinguino", "pingüino"), strength = 1)
#' @export
str_unique ← function(string, ... ) {
  ...
}
```

Title

Description

Parameters

What it returns

Examples

Hard and/or fiddly but worth it

- Links!
 - Other help topics in same package
 - Topics in other packages
 - Vignettes in same or other package
 - Other URLs
- Examples
 - Show authentic usage that hopefully does not ...
 - Create huge headaches in CI and on CRAN

Backticks to format as code. Typical way to refer to a function in its own topic. Use trailing parentheses.

```
#' It's obvious that `thisfunction()` is better than  
#' [otherpkg::otherfunction()] or even our own [olderfunction()].
```

```
#' Read more in our own `vignette("stuff")` or elsewhere in  
#' `vignette("things", package = "otherpkg")`.
```

```
#' It's obvious that `thisfunction()` is better than  
#' [otherpkg::otherfunction()] or even our own [olderfunction()].
```

Square brackets and parentheses for an auto-link to another function, in same package or other package.

See the (R)Markdown support article, Function links

```
#' Read elsewhere in
```

```
#' `vignette("things", package = "otherpkg")`.
```

```
#' It's obvious that `thisfunction()` is better than  
#' [otherpkg::otherfunction()] or even our own [olderfunction()].
```

This vignette syntax is auto-linked in pkgdown (and elsewhere) and gives working code, otherwise.

```
#' Read more in our own `vignette("stuff")` or elsewhere in  
#' `vignette("things", package = "otherpkg")`.
```

URL links

```
#' Use C++ via the cpp11 package
```

```
#'
```

```
#' Adds infrastructure needed to use the [cpp11](https://cpp11.r-lib.org)
```

```
#' package, a header-only R package that helps R package developers handle R
```

```
#' objects with C++ code: ...
```



<URL>

```
#' See <https://happygitwithr.com/common-remote-setups.html> for more about
```

```
#' GitHub remote configurations and, e.g., what we mean by the source repo. This
```

```
#' function works for the configurations `"ours"`, `"fork"`, and `"theirs"`.
```



[link text](URL)

URL links

- "The goal of `urlchecker` is to run the URL checks from R 4.1 in older versions of R and automatically update URLs as needed."
- Extracts the URL-checking logic inside R `CMD check` and exposes as `urlchecker::url_check()`.

```
library(urlchecker)
```

```
# `url_check()` will check all URLs in a package, as is done by CRAN when  
# submitting a package.
```

```
url_check("path/to/pkg")
```

```
# `url_update()` will check all URLs in a package, then update any 301  
# redirects automatically to their new location.
```

```
url_update("path/to/pkg")
```


What's tricky about examples?

- Tension between
 - Showing readable and realistic code
 - However, there's
 - no user, i.e. for interaction or credentials
 - side effects are forbidden
 - can't throw an error
- It's tempting to use `\dontrun{}` but you can get pushback from CRAN

@examplesIf is a mighty weapon

Put something that evaluates to TRUE/FALSE here.

```
#' @examplesIf rlang::is_interactive()  
#'# load/refresh existing credentials, if available  
#'# otherwise, go to browser for authentication and authorization  
#'# drive_auth()  
#'  
#'# see user associated with current token  
#'# drive_user()
```

@examples If is a mighty weapon

Examples

```
if (FALSE) { # rlang::is_interactive()
# load/refresh existing credentials, if available
# otherwise, go to browser for authentication and authorization
drive_auth()

# see user associated with current token
drive_user()
```

Examples should not change the world

- If at all possible, just don't do anything like this:
 - write a file
 - set an option
 - change working directory
- If you must change the world, put it back the way you found it, e.g.
 - write to the temp directory AND delete the file
 - restore the option or working directory to original state
- Sadly, withr functions don't work here. Neither does `on.exit()`.
- Sadly, nothing like `@examplesIf` is available to hide the machinery.

Use `try()` to show an error

```
#' @examples  
#'  
#'  
#'# Row sizes must be compatible when column-binding  
#'try(bind_cols(tibble(x = 1:3), tibble(y = 1:2)))
```



Put code that errors inside `try()`.

```
# Row sizes must be compatible when column-binding  
try(bind_cols(tibble(x = 1:3), tibble(y = 1:2)))  
#> Error in bind_cols(tibble(x = 1:3), tibble(y = 1:2)) :  
#> Can't recycle `..1` (size 3) to match `..2` (size 2).
```

Create roxygen comments

- Go to function definition
- Cursor in function definition
- Insert roxygen skeleton
- Complete the roxygen fields
 - `document()`
 - `?myfunction`
 - 🎉

 Ctrl+. (Start typing function name...)

 Cmd/Ctrl+Alt+Shift+R

 Cmd/Ctrl+Shift+D

 **Your Turn**

check() 

Commit your changes 

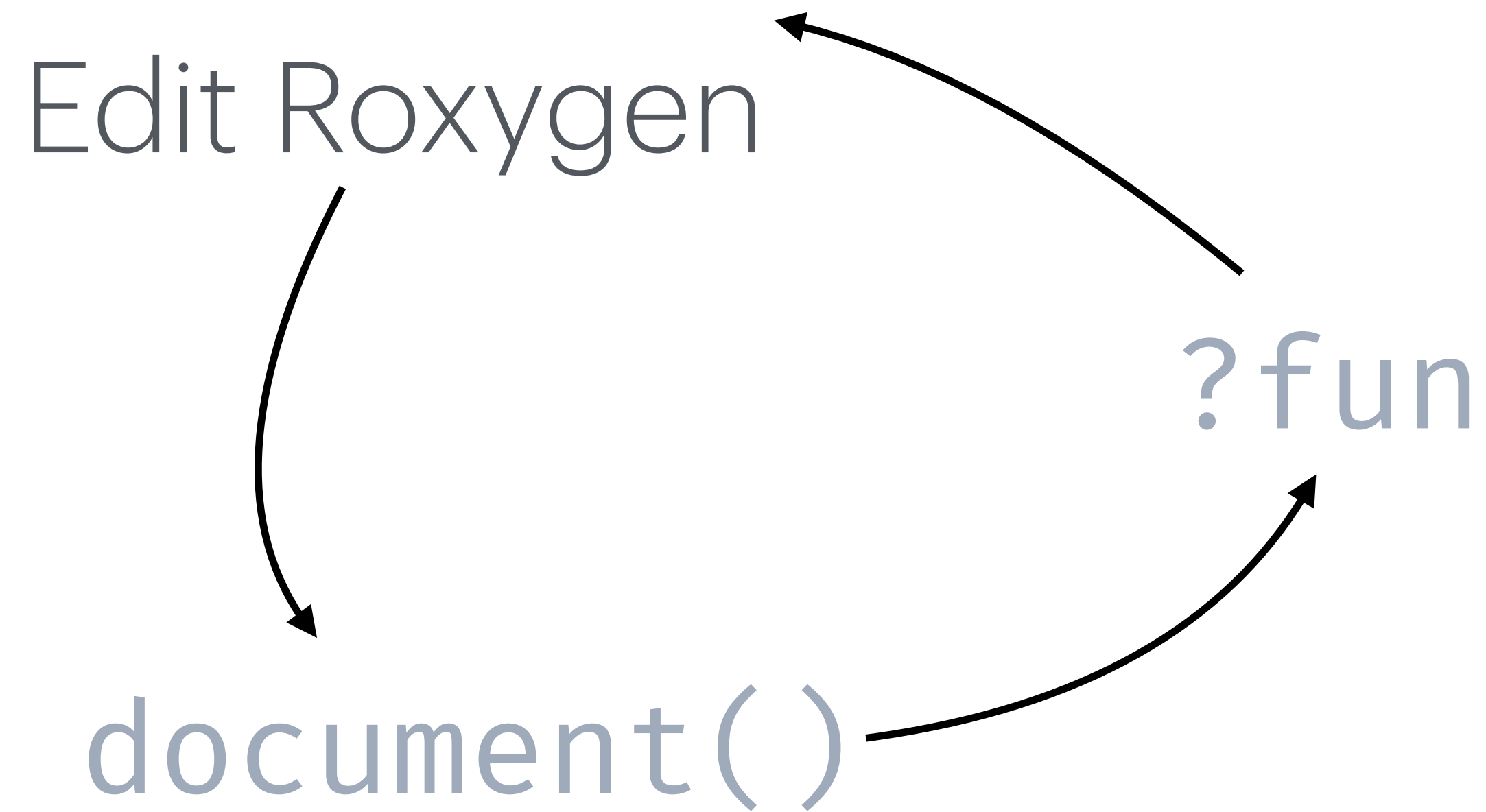
Push to GitHub 

NAMESPACE

An introduction

- Lists R objects that are:
 - **Exported** from your package to be used by package users
 - `export()`, `S3method()`, ...
 - **Imported** from another package to be used internally by your package
 - `import()`, `importFrom()`, ...
- `document()` updates the **NAMESPACE** file with directives from Roxygen comments in your R code.

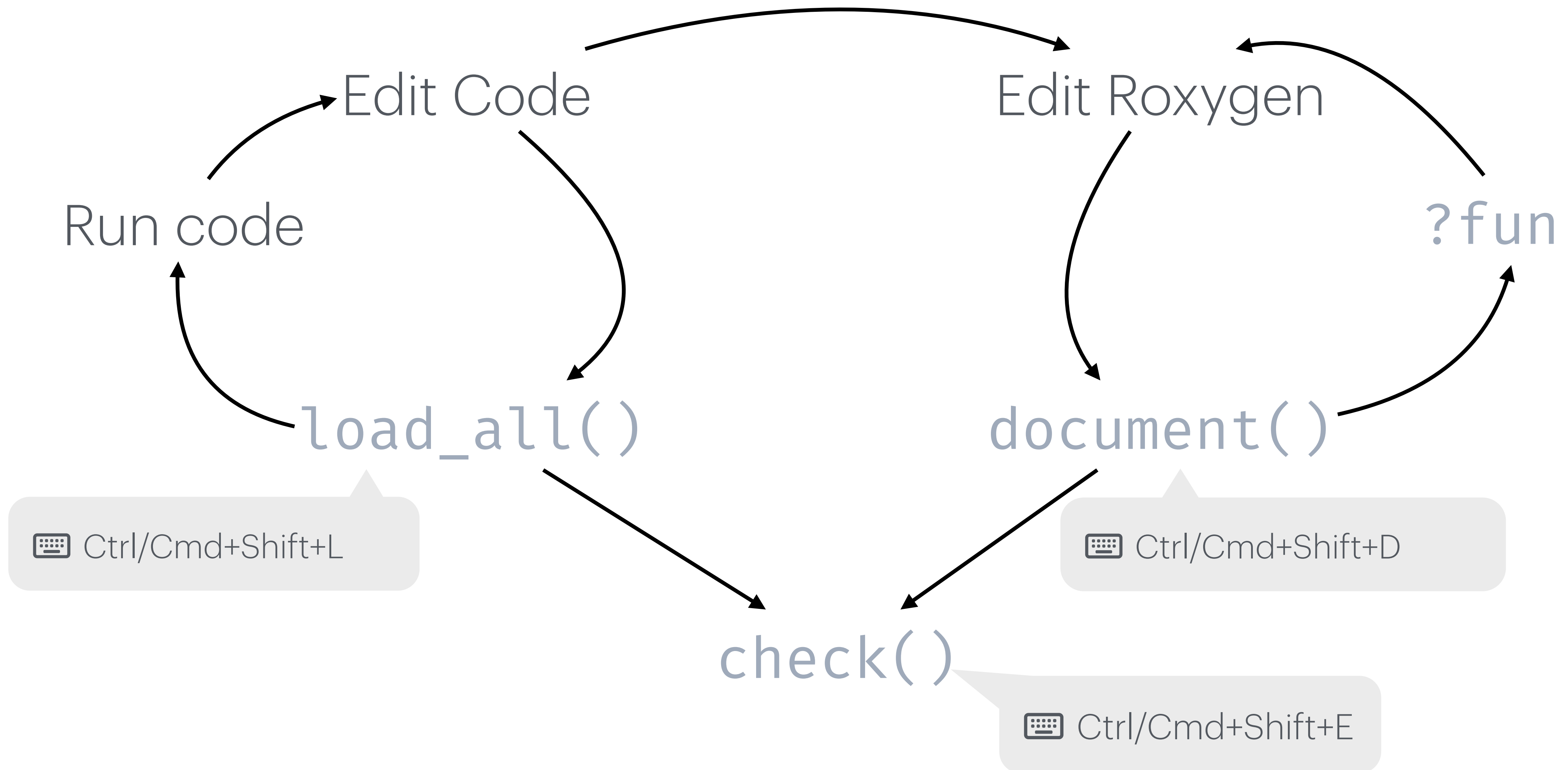
Documentation workflow



 Ctrl+Shift+D (Windows & Linux)

Workflow

Code + documentation + check



Package-level documentation

use_package_doc()

```
use_package_doc()
```

```
#> ✓ Writing 'R/mypackage-package.R'
```

```
#> • Modify 'R/mypackage-package.R'
```

```
document()
```

- Package-level help available via `?mypackage`
- Creates relevant `.Rd` file from `DESCRIPTION`
- A good place for roxygen dependency directives

 **Your Turn**

check() again

check()

#> Documenting

...
#> Building

...
#> Checking


install()

Install package to your library

- R CMD INSTALL

 Ctrl+Shift+B (Windows & Linux)

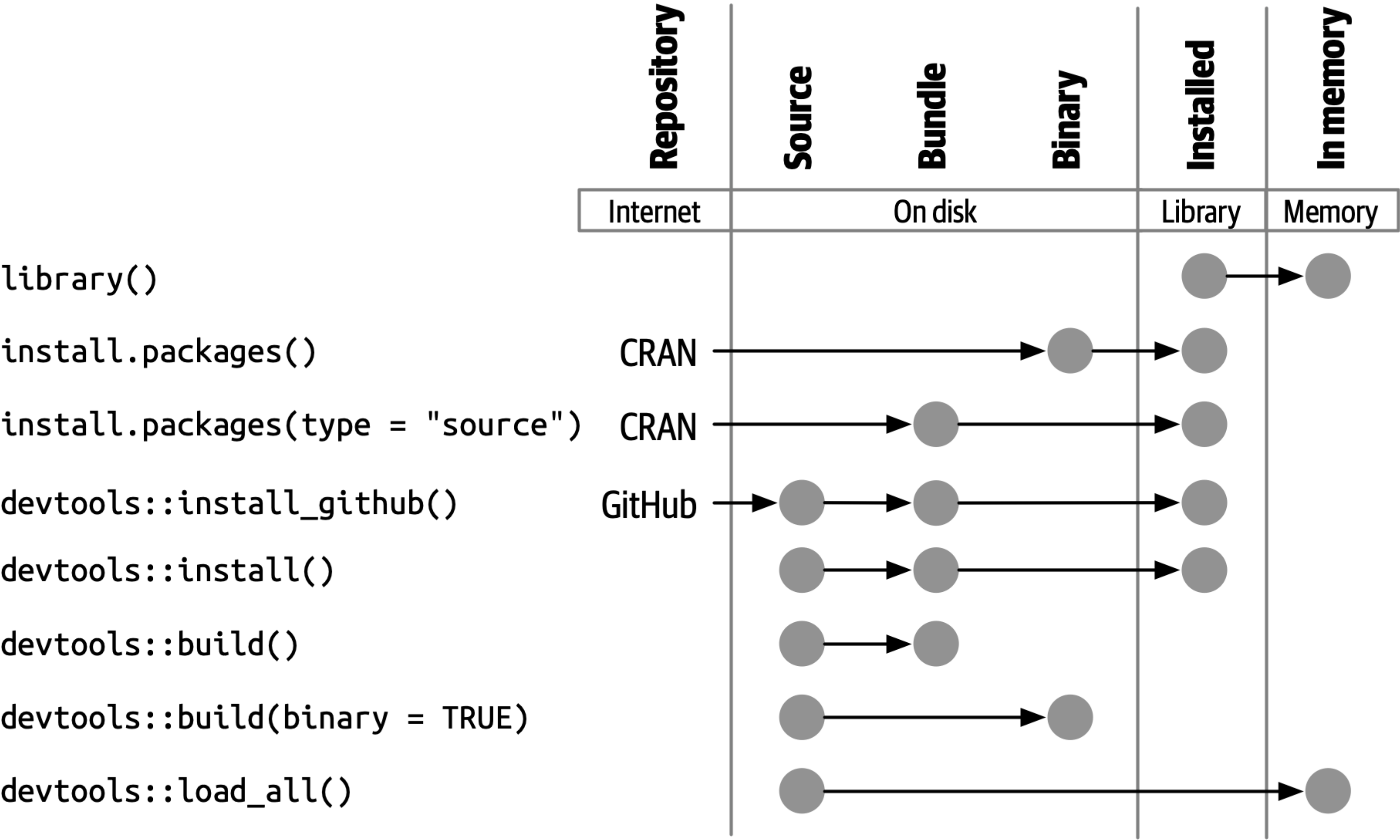
- Restart R

 Ctrl+Shift+F10 (Windows & Linux)

- Attach package with `library()` like any other package

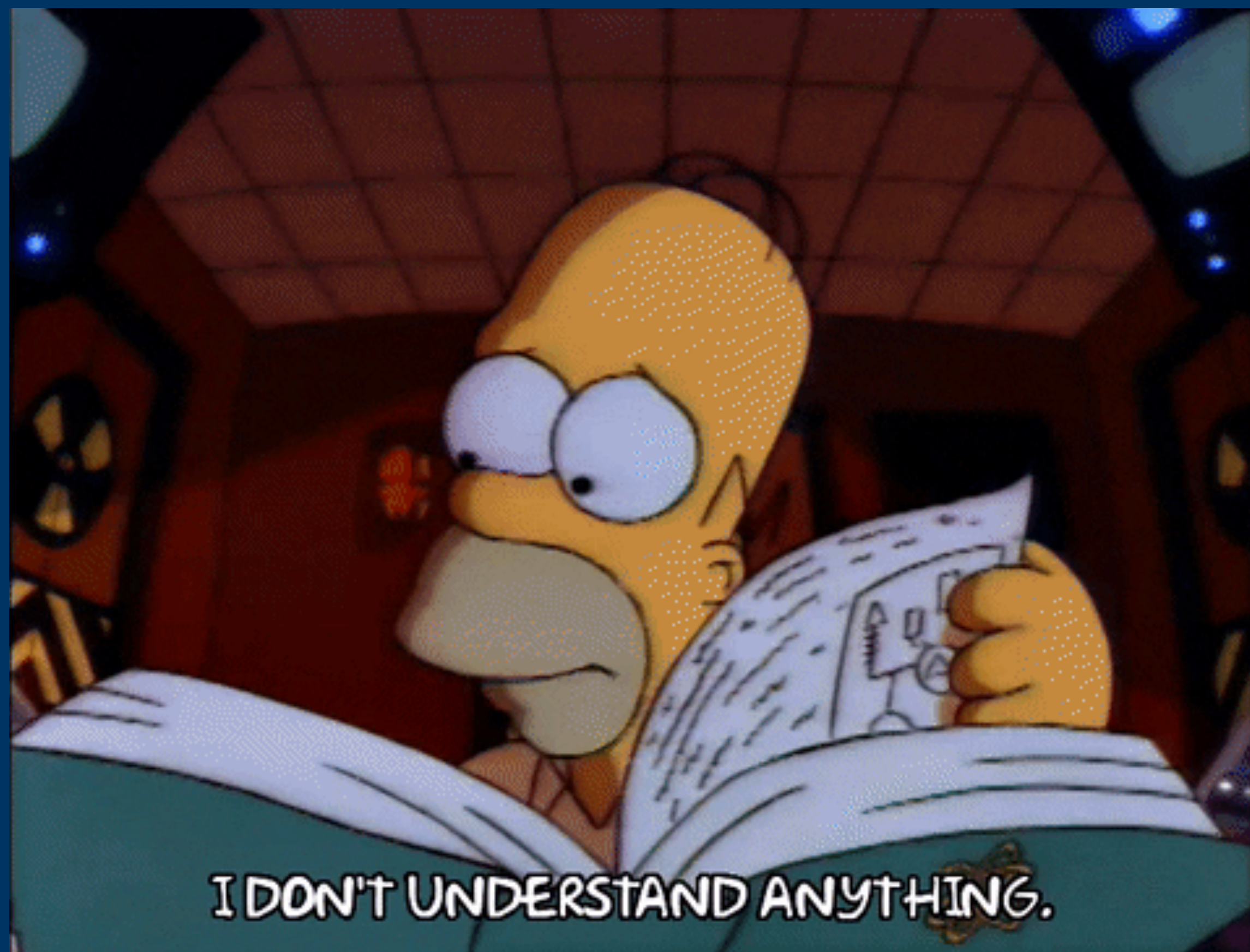
 Your Turn

Package Structure and State



Commit your changes 
Push to GitHub 

README



use_readme_rmd()

Generates README.md, your package's home page on GitHub

- The purpose of the package
- Installation instructions
- Example usage
- Contributing guide

```
use_readme_rmd()
```

```
#> ✓ Writing 'README.Rmd'
```

```
#> ✓ Adding '^README\\.Rmd$' to '.Rbuildignore'
```

```
#> • Update 'README.Rmd' to include installation instructions.
```

```
#> ✓ Writing '.git/hooks/pre-commit'
```

build_readme()

README.Rmd -> README.md

- Installs package to a temporary directory before rendering
- README.md renders on the front page of your GitHub repo

 **Your Turn**

Final check() and install()

You did it!

check()

```
#> — R CMD check results
```

```
#> Duration: 3.1s
```

```
#>
```

```
#> 0 errors ✓ | 0 warnings ✓ | 0 notes ✓
```

install()

```
#> — R CMD build —
```

```
#> checking for file '/Users/jane/rrr/mypackage/DESCRIPTION' ... ✓
```

```
#> preparing 'mypackage':
```

```
#> checking DESCRIPTION meta-information ... ✓
```

```
#> checking for LF line-endings in source and make files and shell scripts
```

```
#> checking for empty or unneeded directories
```

```
#> building 'mypackage_0.0.0.9000.tar.gz'
```

```
#> Running /usr/local/bin/R CMD INSTALL \
```

```
#> /tmp/RtmpK6WnOX/mypackage_0.0.0.9000.tar.gz --install-tests
```

```
#> * installing to library '/Users/jane/Library/R/arm64/4.3/library'
```

```
#> * installing *source* package 'mypackage' ...
```

```
#> ** using staged installation
```

```
#> ** help
```

```
#> *** installing help indices
```

```
#> ** building package indices
```

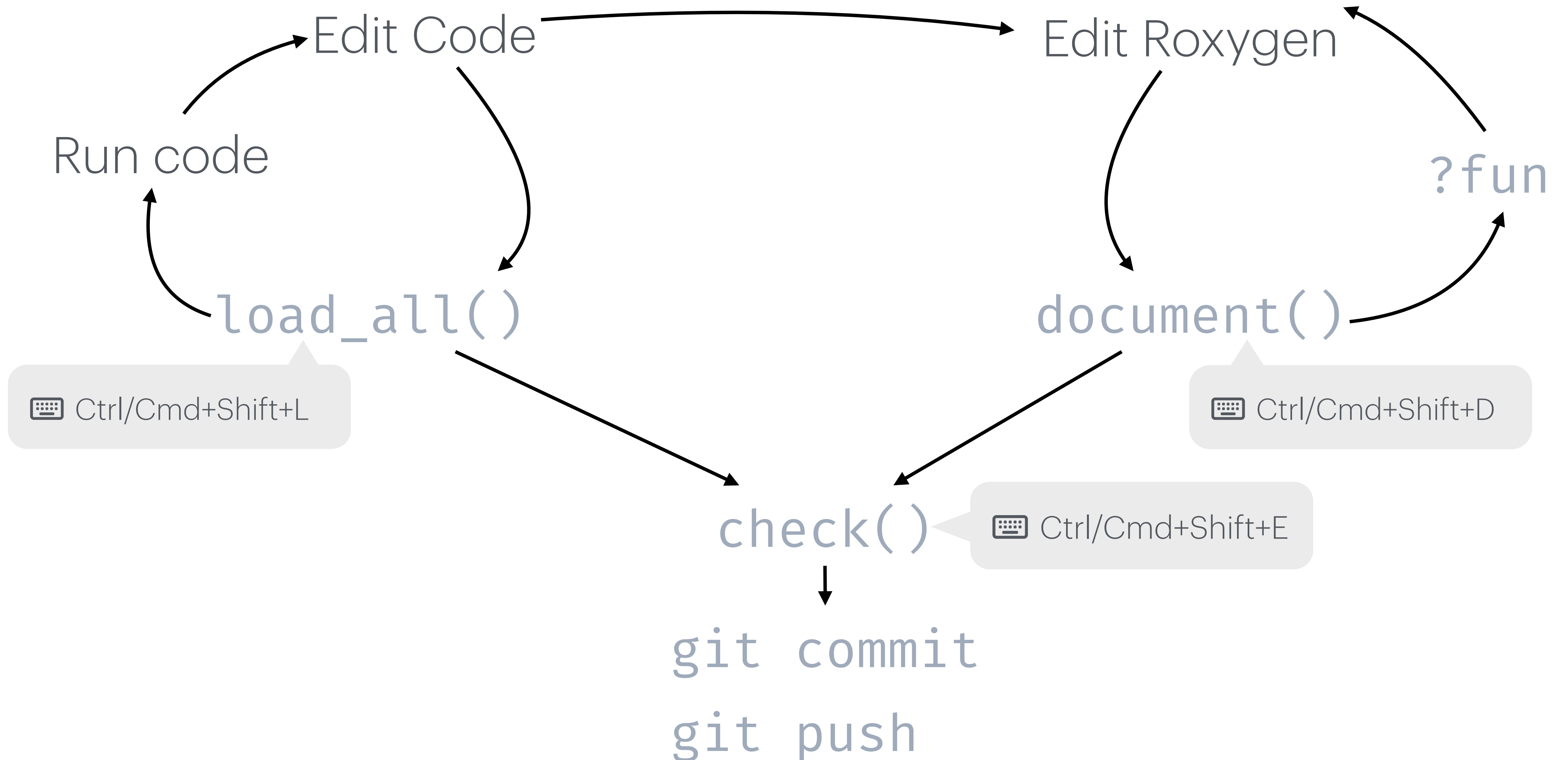
```
#> ** testing if installed package can be loaded from temporary location
```

```
#> ** testing if installed package can be loaded from final location
```

```
#> ** testing if installed package keeps a record of temporary installation path
```

```
#> * DONE (mypackage)
```

Review: Workflow



Commit your changes 
Push to GitHub 

Break Time!

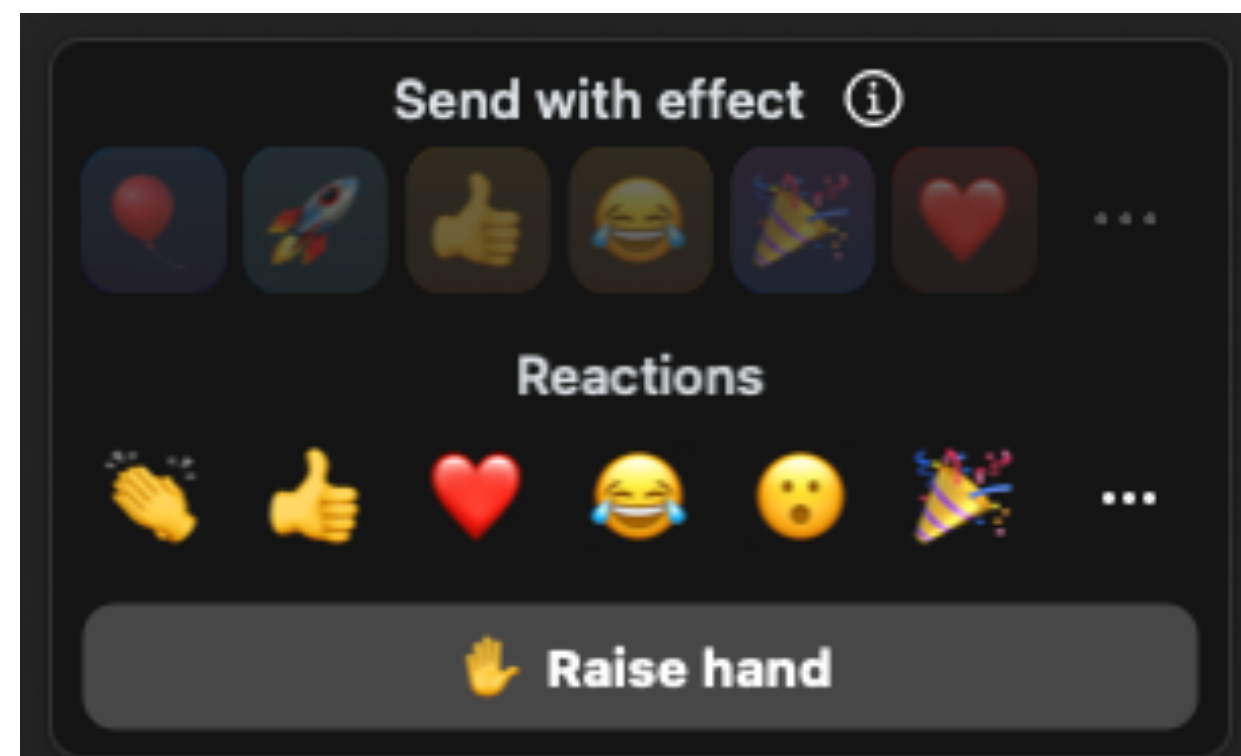


Day one complete!

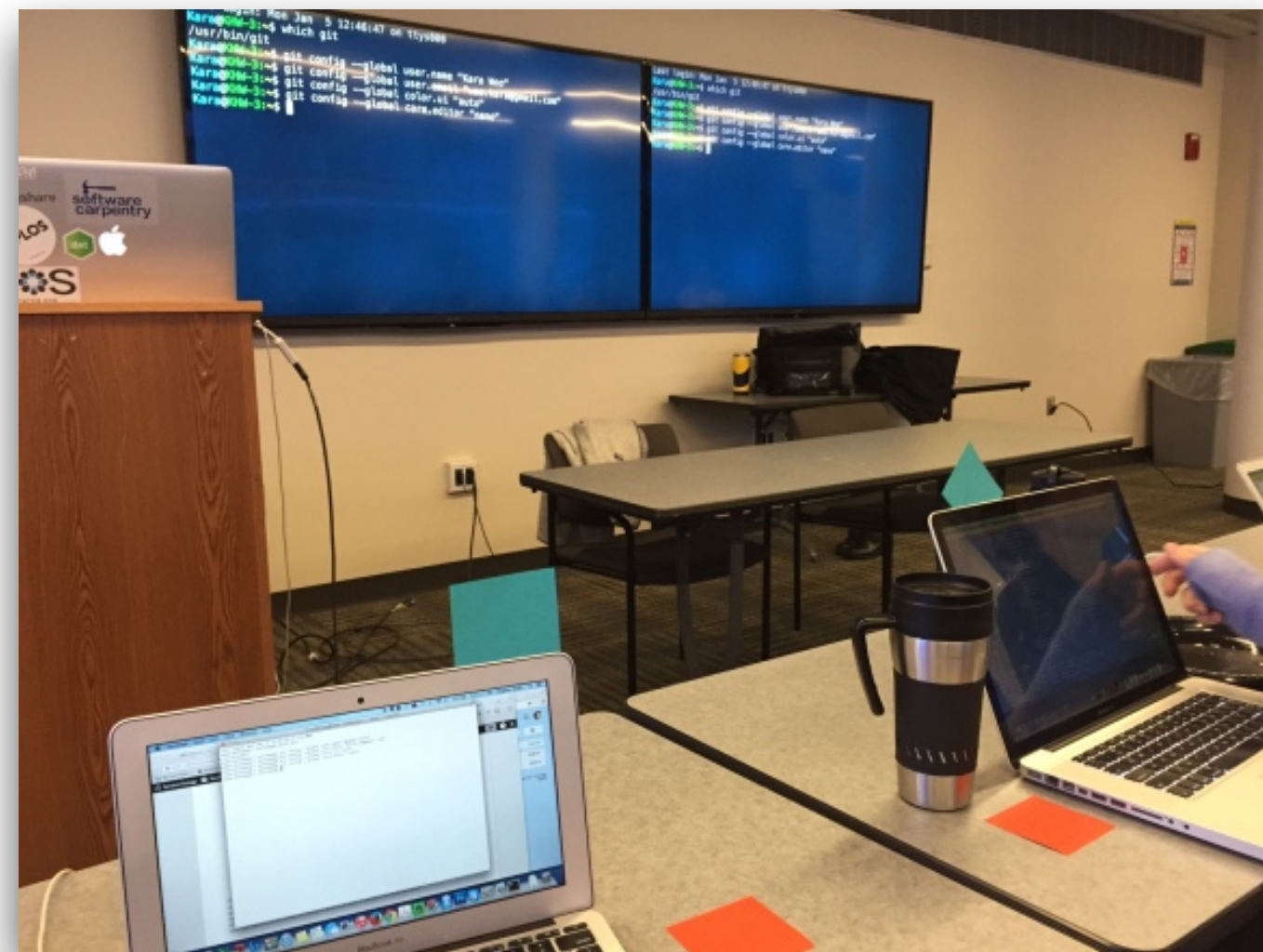
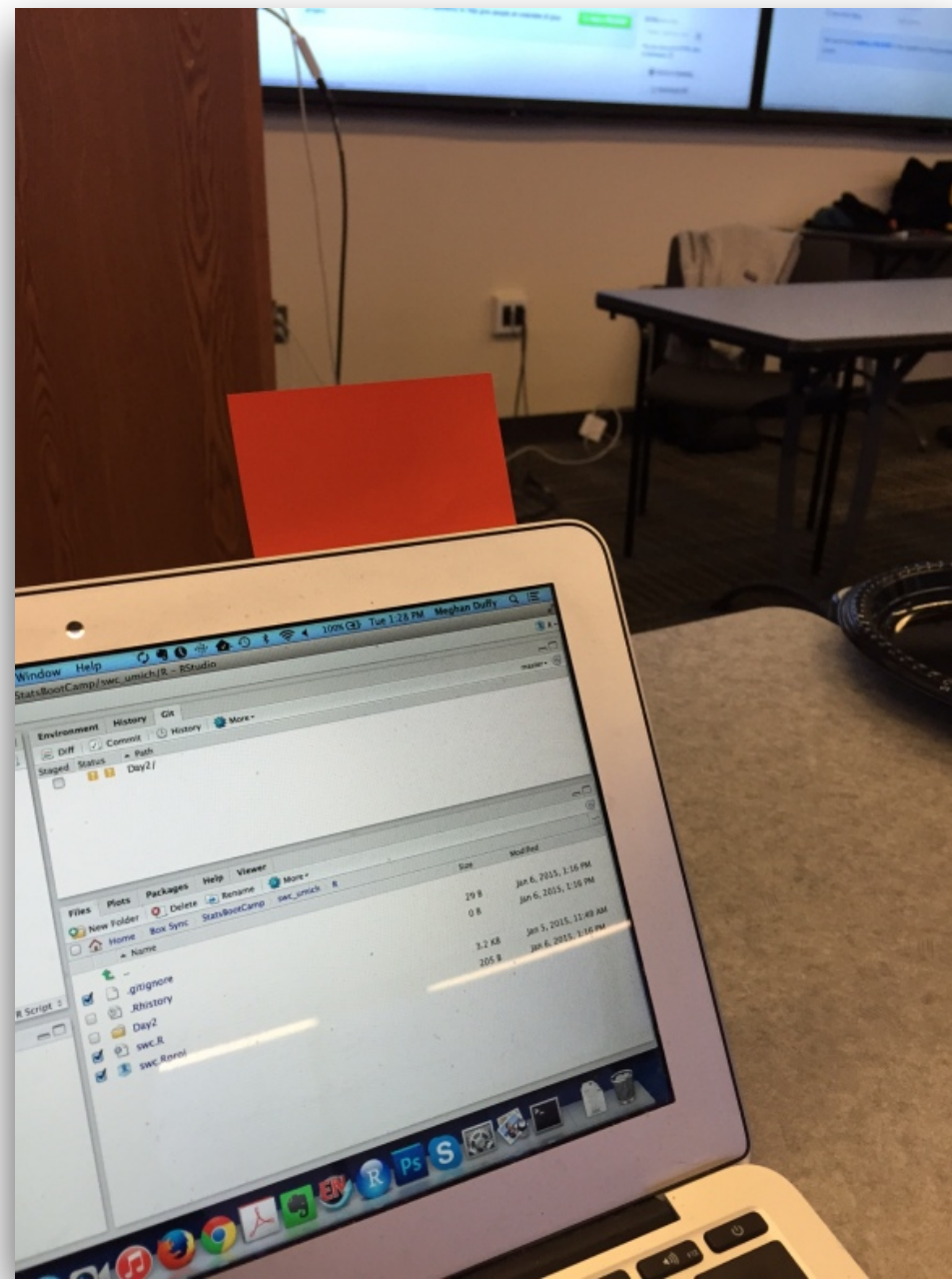
Welcome back to day two

Sticky Notes

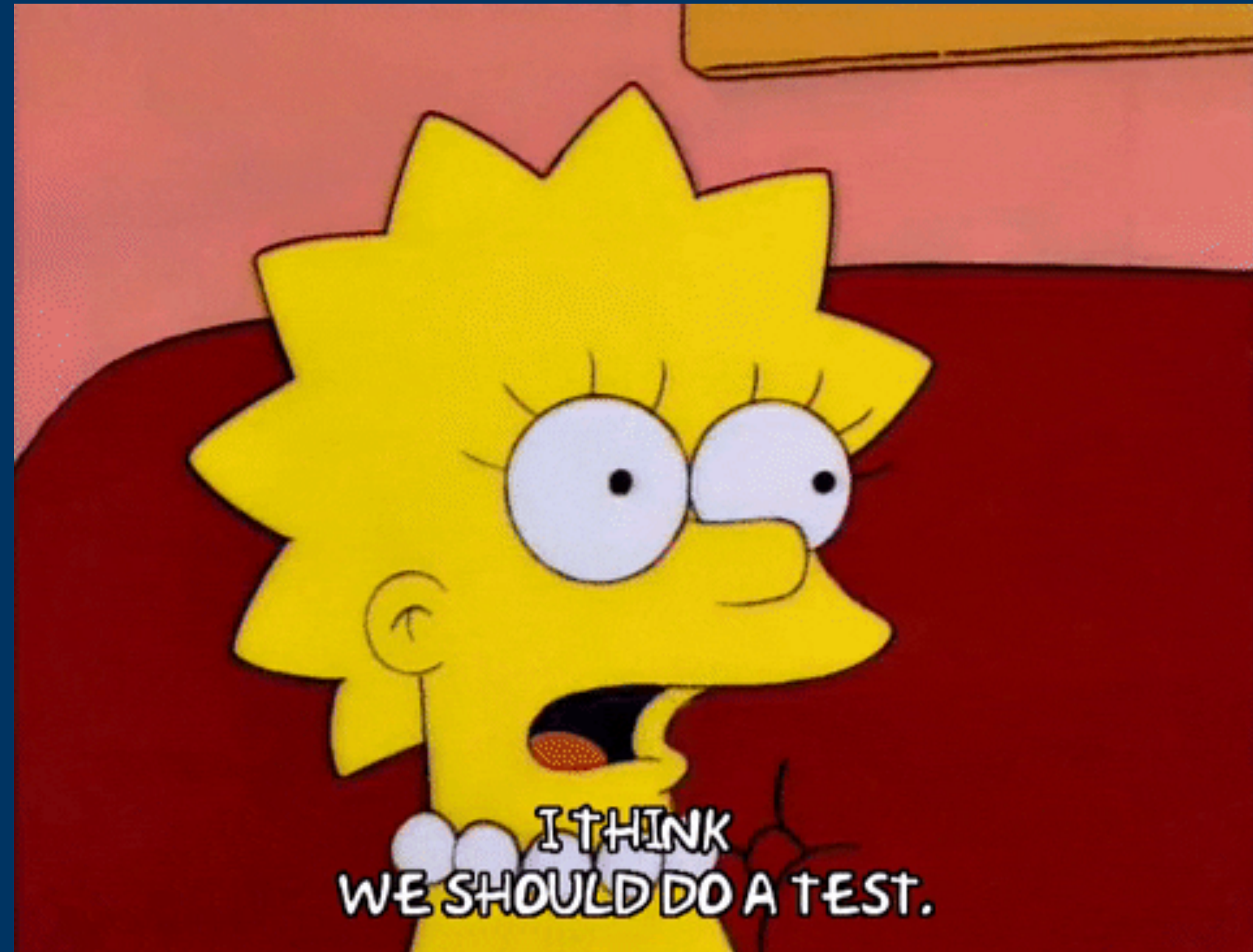
I'm stuck, please send help



I'm good, let's move on



Testing



Day 2: Schedule and Learning Objectives

08:00 - 09:20	Testing & Dependencies	80 min
09:20 - 09:35	Break	15 min
09:35 - 10:55	Continuous Integration & Design Principles	80 min
10:55 - 11:10	Break	15 min
11:10 - 12:30	Design Principles cont'd & Website Creation	80 min

Review: functions

Run once

- `create_package()`
- `use_git()`
- `use_mit_license()`
- `use_readme_rmd()`

Run periodically

- `use_r()`
- `build_readme()`

Run frequently

- `load_all()`

 Ctrl/Cmd+Shift+L

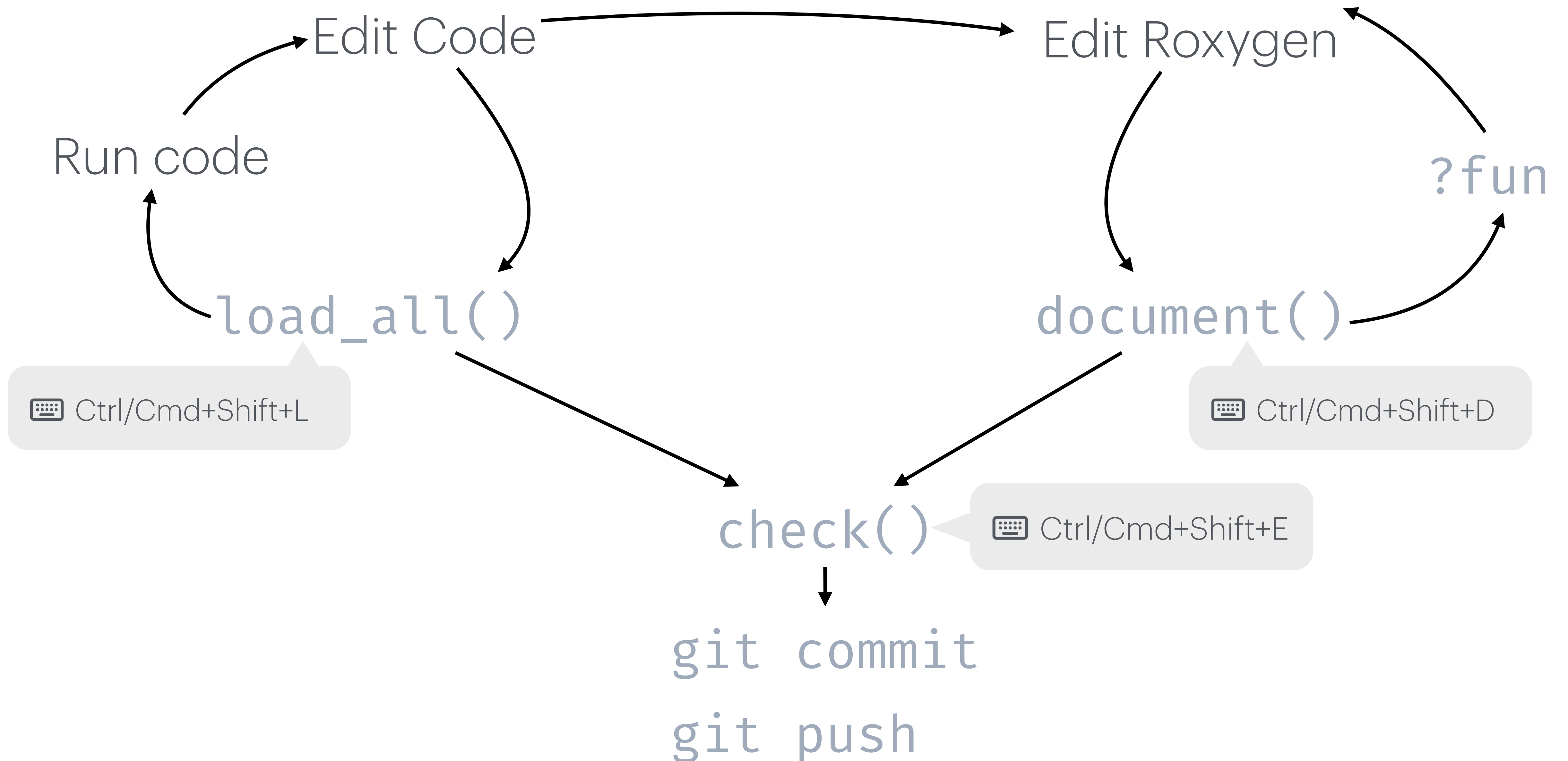
- `document()`

 Ctrl/Cmd+Shift+D

- `check()`

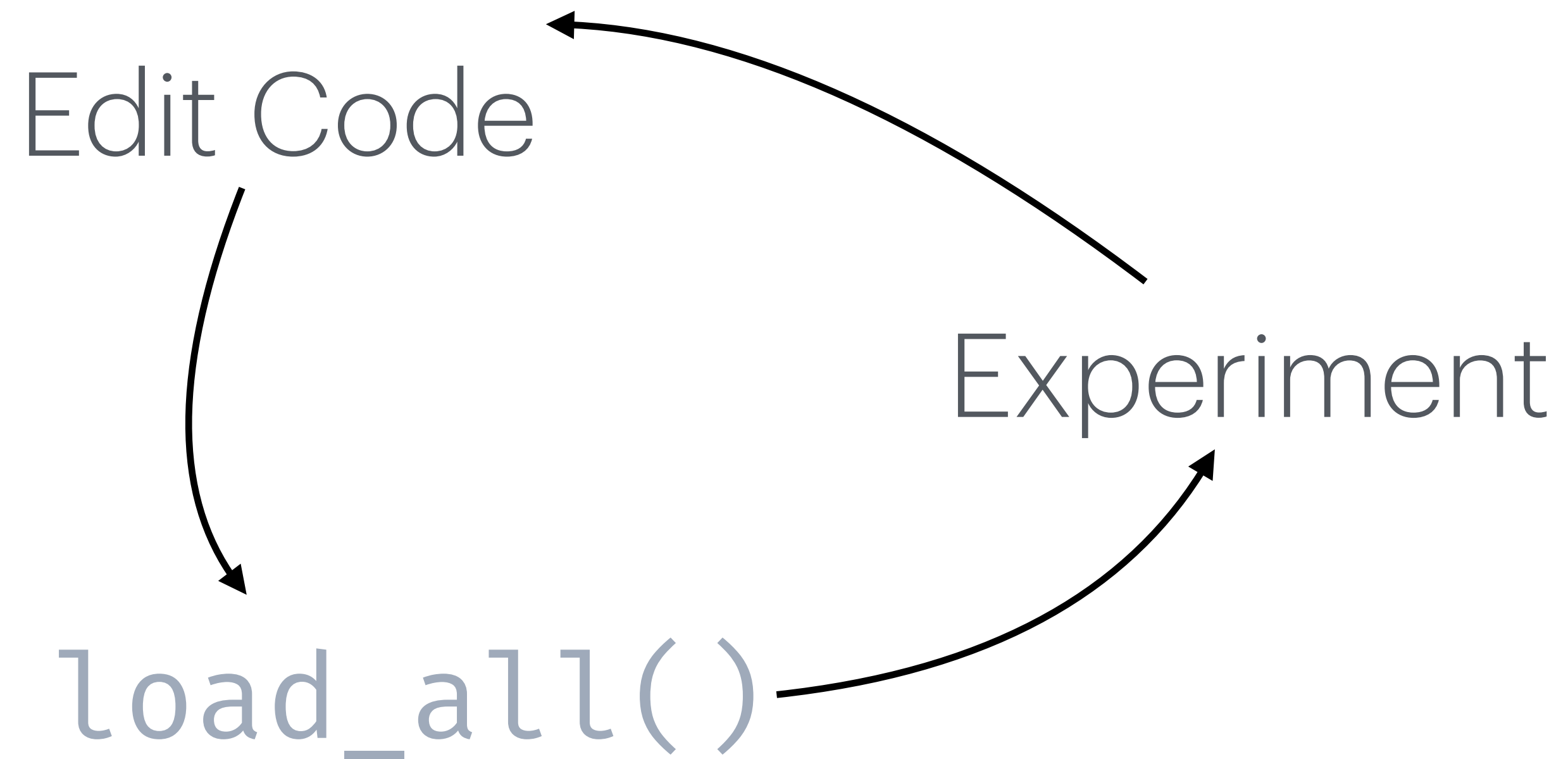
 Ctrl/Cmd+Shift+E

Review: workflow



Testing

Current workflow



Ctrl+Shift+L (Windows & Linux)

Automated Testing

Benefits

- Fewer bugs
- Better code structure
- Call to action when fixing bugs
- Robust (future-proof) code



use_testthat()

Set up formal testing of your package*

```
use_testthat()
```

```
#> ✓ Adding 'testthat' to Suggests field in DESCRIPTION
```

```
#> ✓ Adding '3' to Config/testthat/edition
```

```
#> ✓ Creating 'tests/testthat/'
```

```
#> ✓ Writing 'tests/testthat.R'
```

```
#> • Call `use_test()` to initialize a basic test file and open it for editing.
```

*Sorry, you still have to write the tests

use_test()

```
use_test('my-fun.R')*
```

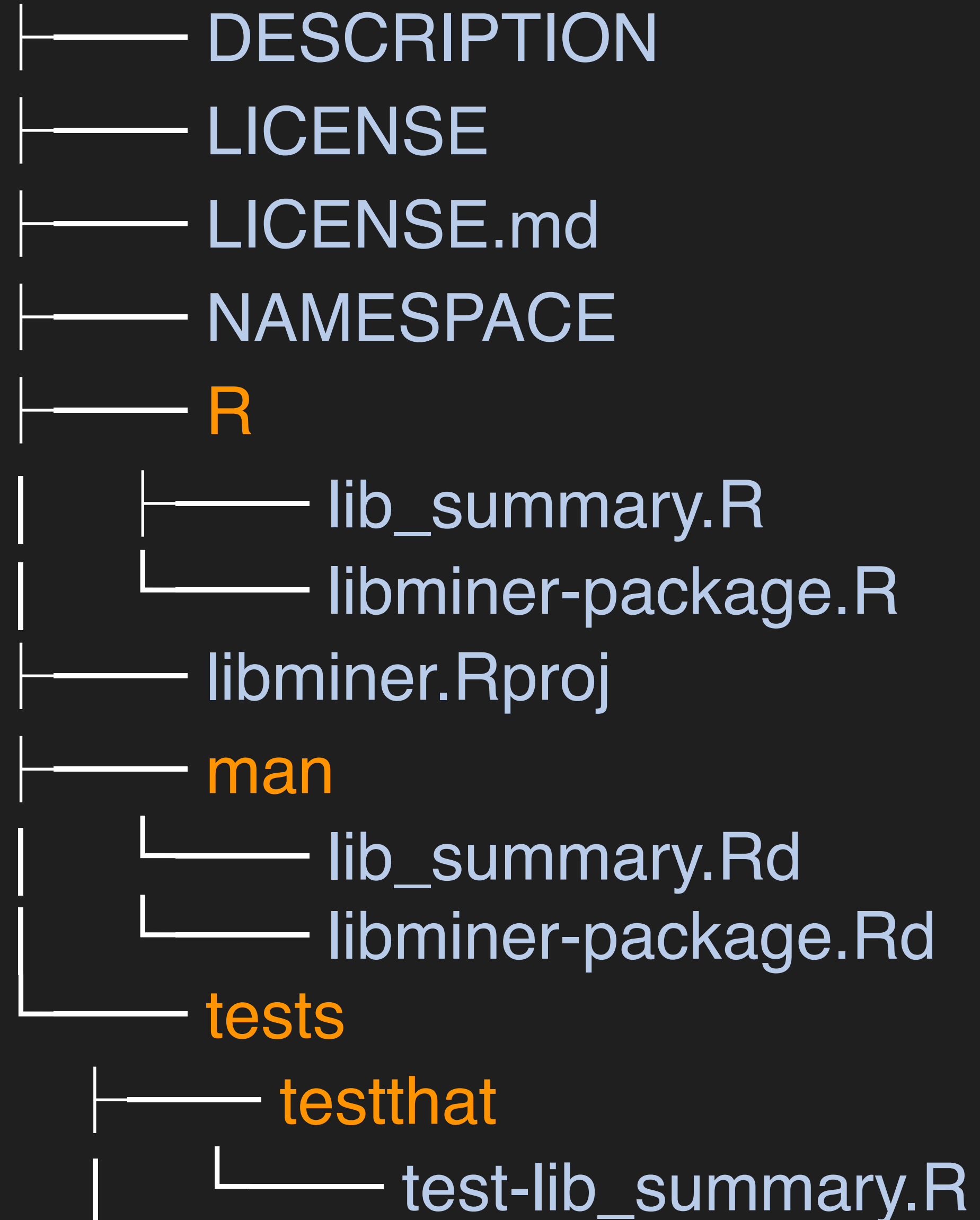
```
#> ✓ Writing 'tests/testthat/test-my-fun.R'
```

```
#> • Edit 'tests/testthat/test-my-fun.R'
```

*Omit file name when **'R/my-fun.R'** is active file

File structure

libminer



`use_test()` + `use_r()` vibe with file pairs

<code>R/a.R</code>	<code>tests/testthat/test-a.R</code>
<code>R/b.R</code>	<code>tests/testthat/test-b.R</code>
<code>R/c.R</code>	<code>tests/testthat/test-c.R</code>

Test structure

```
testthat("description of what you're testing", {  
  expect_equal([function output], [expected output])  
})
```

- **File:** one or more related tests
- **Test:** `test_that("...")`
 - Tests a unit of functionality (hence unit tests)
 - Contains one or more expectations
- **Expectation:** `expect_*(...)`
 - Tests a specific computation and compares it to an expected value

Expectations

- **Objects**

- `expect_equal()`, `expect_type()`, `expect_s3_class()` , ...

- **Vectors**

- `expect_length()`, `expect_lt()`, `expect_gte()`, `expect_true()`, `expect_contains()`, ...

- **Side-effects**

- `expect_error()`, `expect_silent()`, `expect_no_warning()`, `expect_output()`, ...

- **Snapshots**

- `expect_snapshot()`, `expect_snapshot_value()`, ...

test()

- Runs all tests in your test suite

```
test()
```

```
#> i Testing
```

```
#> ✓ I F W S OK I Context
```

```
#>
```

```
#> :  | 0 |
```

```
#> ✓  | 1 |
```

```
#>
```

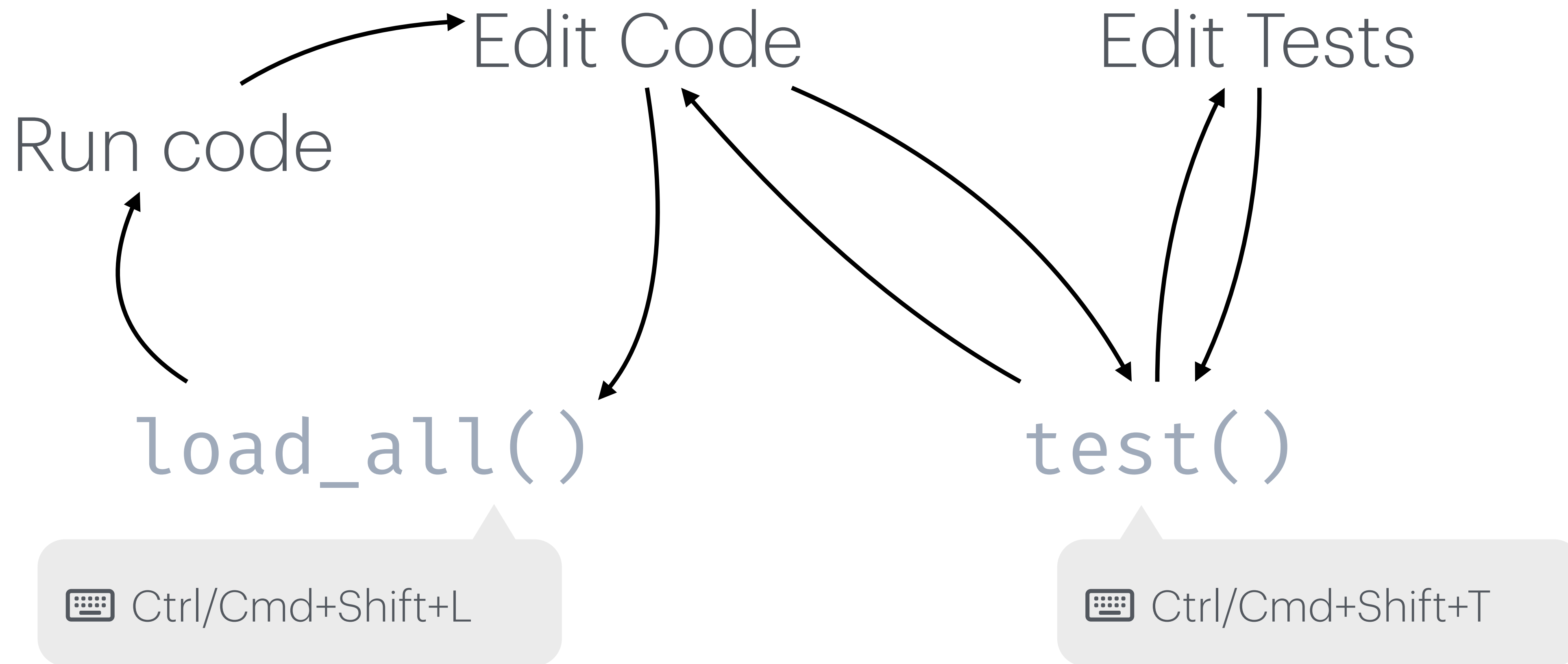
```
#> — Results
```

 Ctrl+Shift+T (Windows & Linux)

 **Your Turn**

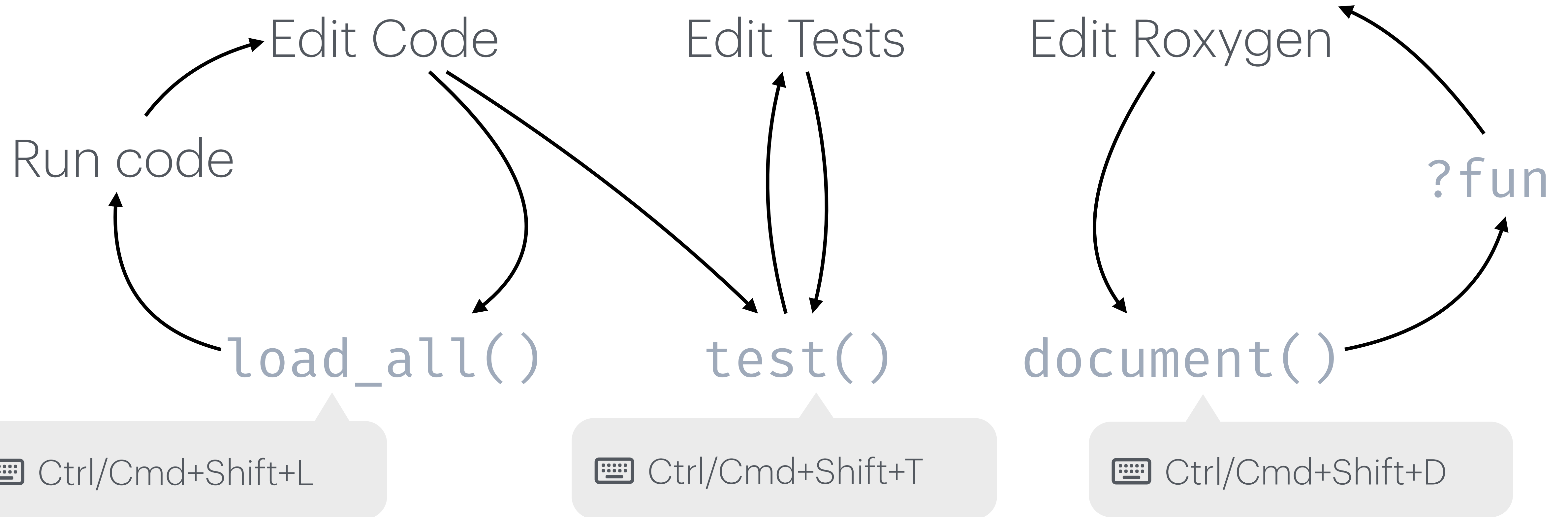
Updated workflow

Code + testing



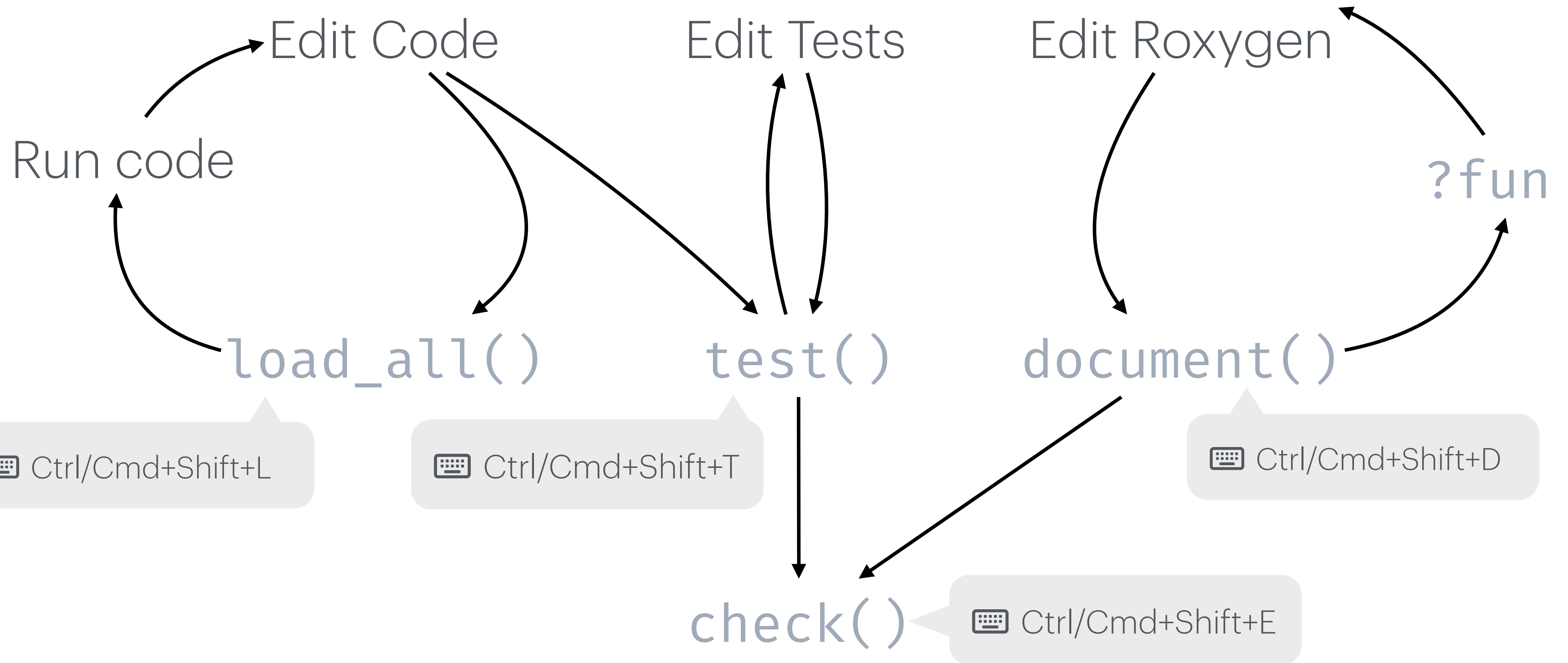
Workflow

Code + testing + documentation



Workflow

Code + testing + documentation + check



check() 

Commit your changes 

Push to GitHub 

testthat workflow patterns

load_all()

- testthat's workflow is designed around `load_all()`
- Makes entire package namespace available
- Attaches testthat
- Sources `tests/testthat/helper.R`



Functions that help with testing

Workflow: micro-iteration, interactive experimentation

- tweak the `foofy()` function and re-load it with `load_all()`
 - also attaches `testthat` and sources any test helpers
- interactively explore and refine expectations
`expect_equal(foofy(...), EXPECTED_FOOFY_OUTPUT)`
- and tests
`test_that("foofy does good things", { ... })`

Workflow: mezzo-iteration, whole test file

```
test_file("tests/testthat/test-foofy.R")
```

- In RStudio, with test or R file focused

```
test_active_file()
```

```
test_coverage_active_file()
```

*Consider binding these to Cmd + T, Cmd + R

Keyboard Shortcuts

Show: All Customized

[? Customizing Keyboard Shortcuts](#)

Name	Shortcut	Scope
Compare test results for Shiny application		Workbench
Record a test for Shiny		Workbench
Run shinytest Test		Workbench
Run tests for Shiny application		Workbench
Run testthat Tests		Workbench
Test Package	Shift+Cmd+T	Package Development
Calculate package test coverage		Addin
Calculate package test coverage		Addin
Format test_that test file		Addin
Initialize test_that()		Addin
Navigate To Test File		Addin
Report test coverage for a file		Addin
Report test coverage for a file	Cmd+R	Addin
Report test coverage for a package		Addin
Report test coverage for a package	Shift+Cmd+R	Addin
Run a test file		Addin
Run a test file	Cmd+T	Addin
View Latest Run		Addin



Workflow: macro-iteration, all files

test()

 Ctrl/Cmd+Shift+T

test_coverage()

 Ctrl/Cmd+Shift+R

check()

 Ctrl/Cmd+Shift+E

 **Your Turn**

Test suite design principles

- A test should be self-sufficient and self-contained.
- The interactive workflow is important.
- Obvious >>> DRY
- Don't let a nonstandard workflow "leak".

Test smell: top-level code that's outside `test_that()`

```
dat <- data.frame(x = c("a", "b", "c"), y = c(1, 2, 3))
```

```
skip_if(today_is_a_monday())
```

```
test_that("foofy() does this", {  
  expect_equal(foofy(dat), ...) })
```

```
dat2 <- data.frame(x = c("x", "y", "z"), y = c(4, 5, 6))
```

```
skip_on_os("windows")
```

```
test_that("foofy2() does that", {  
  expect_snapshot(foofy2(dat, dat2)) })
```

Deodorizing the previous example

```
test_that("foofy() does this", {  
  skip_if(today_is_a_monday())  
  
  dat ← data.frame(x = c("a", "b", "c"), y = c(1, 2, 3))  
  
  expect_equal(foofy(dat), ... )  
})
```

Move file-scope logic to a narrower scope (as done here) or a broader scope (test setup and/or test helpers).

```
test_that("foofy() does that", {  
  skip_if(today_is_a_monday())  
  skip_on_os("windows")  
  
  dat ← data.frame(x = c("a", "b", "c"), y = c(1, 2, 3))  
  dat2 ← data.frame(x = c("x", "y", "z"), y = c(4, 5, 6))  
  
  expect_s3_class(foofy(dat, dat2), "data.frame")  
})
```

It's OK to repeat yourself!

Leave the world the way you found it

```
circumference <- function(r) {2 * pi * r}
test_that("calculations work", {
  withr::local_options(digits = 3)
  expect_equal(circumference(r = 8), 50.3))
})
```

withr's `local_*()` functions are super useful for making changes that are scoped to a single `test_that()`.

Things to avoid in your tests

`library(somedependency)`

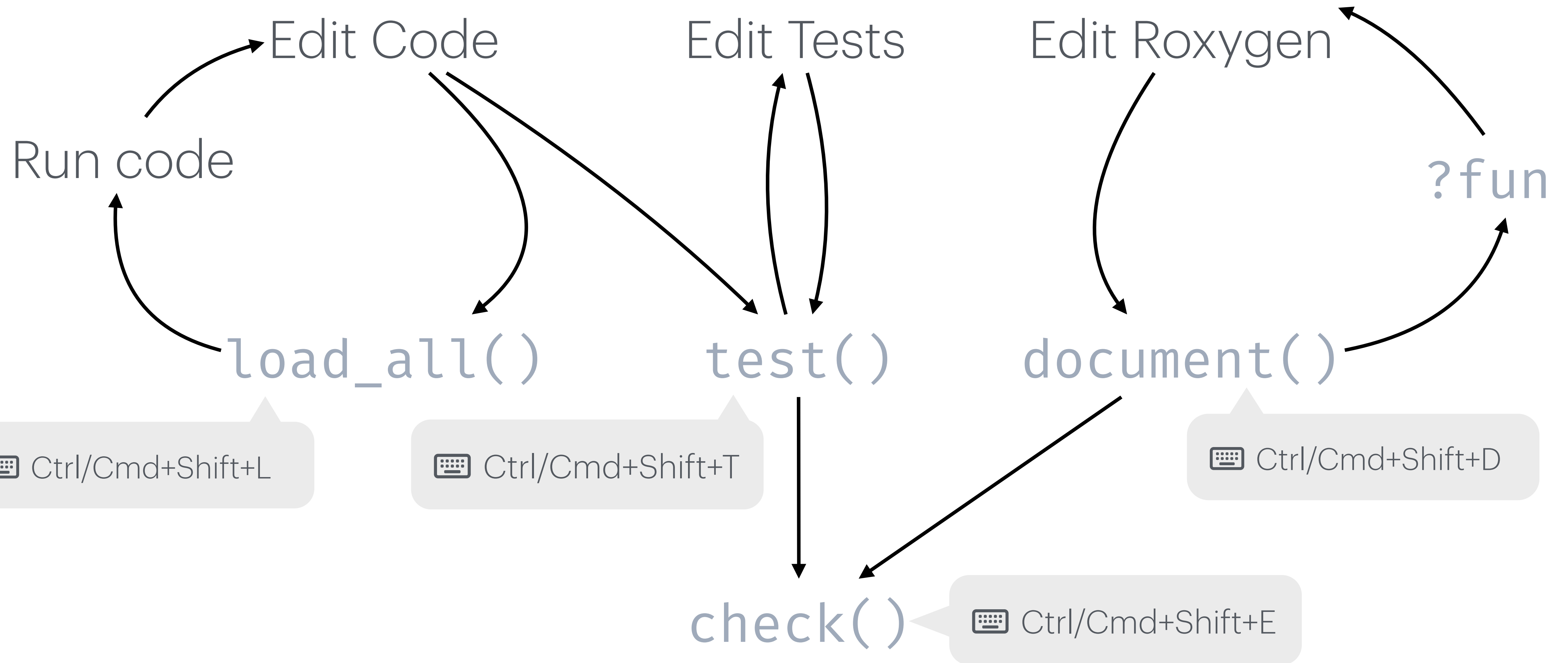
Please no! Access functions from your dependencies, in your tests, exactly as you do below R/.

`source("stuff-thats-handy-for-your-tests.R")`

Please no! Unexported helper functions and test helper files are a better mechanism for this.

Workflow

Code + testing + documentation + check



Dependencies



Use functions from another package inside your package

```
library("fs")
```

```
# use function
```


use_package()

Add a dependency

- Use functions from another package inside your package
- Dependencies must be declared
 - Even from included packages (`stats::sd()`, `tools::file_ext()` etc.)
- Never call `library(pkg)` in code below `R/!`

```
use_package("fs")
```

```
#> ✓ Adding 'fs' to Imports field in DESCRIPTION
```

```
#> • Refer to functions with `fs::fun()``
```

Listing dependencies in DESCRIPTION

Three options

- **Depends:**

- Ensures the package is installed with your package
- *Attaches the package when yours is attached*
- Rarely needed or recommended

- **Imports:**

- Ensures the package is installed with your package
- Most common location for dependencies

- **Suggests:**

- Does not ensure installation automatically
- Packages required for development (running tests, building vignettes, etc).
- Rarely used functionality (especially if the dependency is difficult to install)

devtools DESCRIPTION file:

<https://github.com/r-lib/devtools/blob/main/DESCRIPTION>

Imports: DESCRIPTION vs NAMESPACE

DESCRIPTION

- Lists packages that your package requires
- Ensures required packages are **installed** during package installation
- **Does not** import that package into your package's namespace
- Add via `use_package()` (or manually)

NAMESPACE

- **Imports** R objects from another package into your package's namespace
- **import** == Available to be used internally by your package
- Don't edit manually - use roxygen tags:
`#' @importFrom pkg fun`
`#' @import pkg`

3 ways to use functions from another package

1 - Call function with namespace qualifier

1. Add package to **DESCRIPTION** file in **Imports**
2. Call function like **package::fun()**

☑ Most common and recommended pattern

DESCRIPTION

```
Imports:  
  purrr
```

R/my-fun.R

```
#' @export  
myfun <- function(x) {  
  purrr::map(x, mean)  
}
```

NAMESPACE

```
export(myfun)
```

document()



3 ways to use functions from another package

2 - Import just the functions you want to use via `@importFrom` tag:

1. Add package to **DESCRIPTION** file in **Imports**
2. Use `@importFrom` roxygen tags
3. Call function like `fun()`

DESCRIPTION

```
Imports:  
purrr
```

R/my-fun.R

```
#' @importFrom purrr map  
#' @export  
myfun <- function(x) {  
  map(x, mean)  
}
```

NAMESPACE

```
importFrom(purrr, map)  
export(myfun)
```

document()



3 ways to use functions from another package

3 - Import the entire package via `@import` roxygen tag:

1. Add package to **DESCRIPTION** file in **Imports**
2. Use `@import` roxygen tag
3. Call functions like `fun()`

DESCRIPTION

```
Imports:  
  purrr
```

R/my-fun.R

```
#' @import purrr  
#' @export  
myfun <- function(x) {  
  y <- map(x, mean)  
  reduce(y, `+`)  
}
```

NAMESPACE

```
import(purrr)  
export(myfun)
```

document()

3 ways to use functions from another package

1. `package::fun()`
2. Import just the functions you want to use via `@importFrom` roxygen tag:

```
#' @importFrom pkg fun1 fun2
```

Adds to NAMESPACE:

```
importFrom(pkg, fun1)
```

```
importFrom(pkg, fun2)
```

*Shortcut: `usethis::use_import_from("pkg", "function")`

3. Import the entire package with `@import`:

```
#' @import pkg
```

Adds to NAMESPACE:

```
import(pkg)
```



Use your new dependency

Write a function using a function from the dependent package

- `use_package("fs")`
- Write/edit function using dependency: `pkg::fn()`
- Edit roxygen comments
- `document()`
 - Writes `man/*.Rd` files & regenerates `NAMESPACE`
- Update tests

 **Your Turn**

Let's add one more

`use_import_from("pkg", "function")`

- See:
 - DESCRIPTION
 - R/mypackage-packge.R (remember `use_package_doc()`?)
 - NAMESPACE
- Write/edit function using dependency: `fn()`
- `*test()`

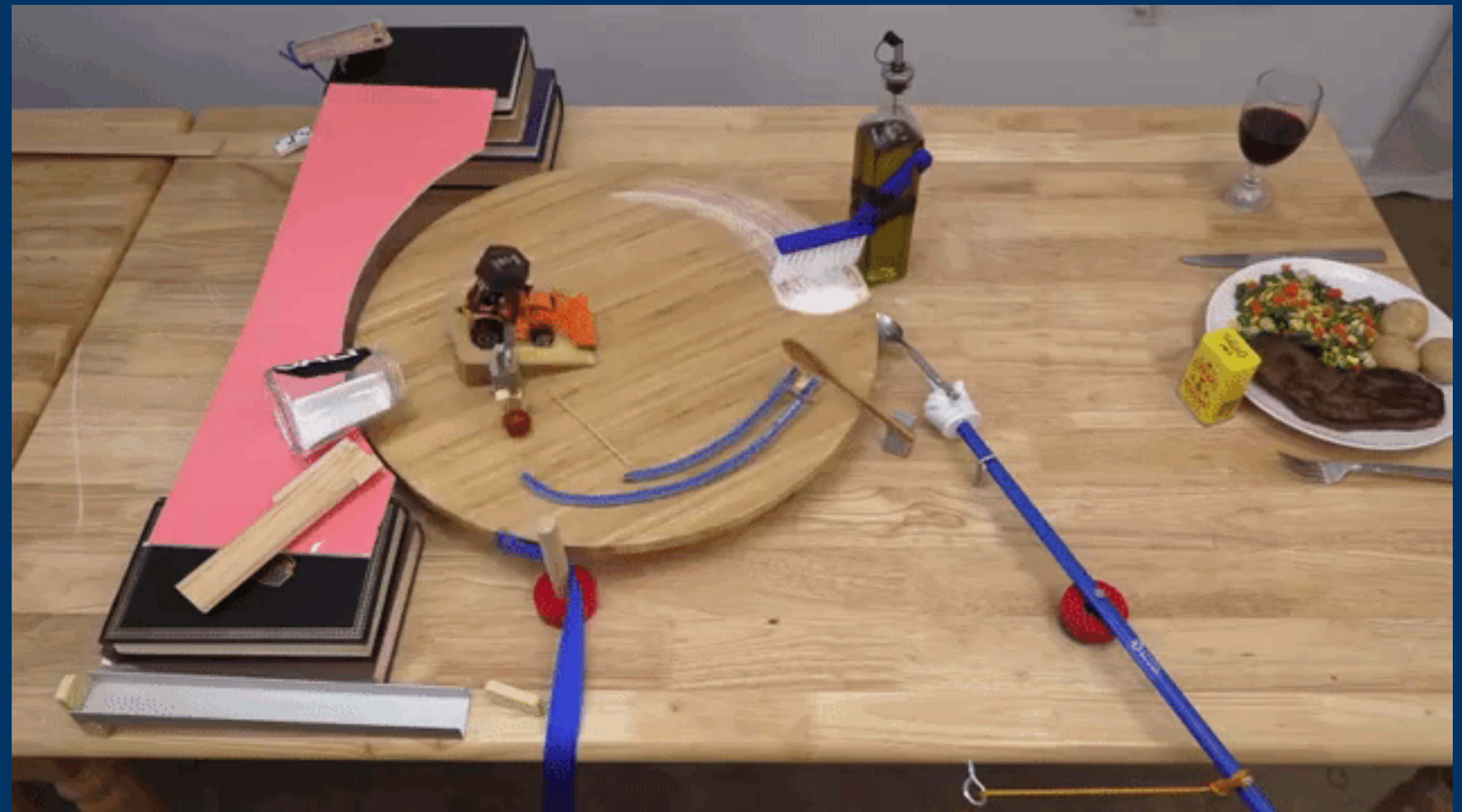
 Your Turn

check() 

Commit your changes 

Push to GitHub 

Continuous Integration



use_github_action()

github.com/r-lib/actions

- "check-standard": Runs **R CMD check** when you push
- "test-coverage": Compute test coverage and report at codecov.io
 - (Requires an account and API key)
- "pr-commands": Enables automatic documentation and styling of code via special PR comments
- And more...

<https://github.com/r-lib/actions/tree/v2-branch/examples>

 **Your Turn**

Break Time!

Design Principles

Review: functions

Run once

- `create_package()`
- `use_git()`
- `use_mit_license()`
- `use_testthat()`
- `use_readme_rmd()`

Run periodically

- `use_r()`
- `use_test()`
- `use_package()`
- `rename_files()`
- `build_readme()`
- `use_github_action()`

Run frequently

- `load_all()`

 Ctrl/Cmd+Shift+L

- `document()`

 Ctrl/Cmd+Shift+D

- `test()`

 Ctrl/Cmd+Shift+T

- `check()`

 Ctrl/Cmd+Shift+E

Learning Objectives

- At the end of this section you will be able to:
 - Understand core principles about **function design**
 - Apply the **DRY principle** to avoid redundancy and improve maintainability in your R code
 - Design functions with **default arguments, input validation, and helper functions** for greater clarity and flexibility
 - Enhance **code readability and consistency** by following structured style guidelines and effective naming conventions

Why Design?

Focus on core principles to make R code cleaner and easier to manage

Tidyverse guiding design principles

- Programming is a task performed by **humans**
- Reduce your cognitive load with the **consistent** design (as much as possible)
- Make your functions and systems **composable**
- Think about others who are not like us to create an **inclusive** space

```
cor(1:10, 2:11, "pairwise.complete.obs", "spearman")
```

```
cor(1:10, 2:11, use = "pairwise.complete.obs", method = "spearman")
```

Naming things

Some general guidelines

- Use verbs to ascribe as action
- Use consistent style
- Use singular (because English is weird)
- Consider short prefixes to unify package functions
- Don't be afraid to be verbose

```
## bad
```

```
gse(date = "1977-05-25")
```

```
## good
```

```
get_salmon_escapement(date = "1977-05-25")
```

Pure functions versus Side effects

- A pure function:
 - produces the same output for the same input, with no impact on other parts of the program
 - only change is what is returned
 - Easier to test and maintain
- Function with side effects
 - A function that interacts with the outside environment

Pure functions

```
add ← function(x, y) {  
  x + y  
}  
add(2, 3)  
#> [1] 5
```

Functions with Side effects

```
library(readr)
read_csv(readr_example("mtcars.csv"))
#> # A tibble: 32 × 11
#>   mpg   cyl  disp    hp  drat    wt   qsec    vs  am  gear  carb
#>   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#> 1  21     6  160   110  3.9   2.62  16.5    0    1     4     4
#> 2  21     6  160   110  3.9   2.88  17.0    0    1     4     4
#> 3  22.8    4  108    93  3.85  2.32  18.6    1    1     4     1
#> 4  21.4    6  258   110  3.08  3.22  19.4    1    0     3     1
#> 5  18.7    8  360   175  3.15  3.44  17.0    0    0     3     2
#> 6  18.1    6  225   105  2.76  3.46  20.2    1    0     3     1
#> 7  14.3    8  360   245  3.21  3.57  15.8    0    0     3     4
```

The DRY Principle: Don't Repeat Yourself

What is DRY?

- Minimize redundancy by centralizing logic
- Benefits
 - Consistency
 - Easier debugging
 - Maintenance

The DRY Principle: Don't Repeat Yourself

Identifying Redundant Code

- Signs of redundant code
 - Repeated data transformations
 - Hardcoded values
- Think MODULAR
- Look for patterns

The DRY Principle: Don't Repeat Yourself

Refactor to use functions

```
# Without DRY
se_age ← sd(df$age) / sqrt(length(df$age))
se_income ← sd(df$income) / sqrt(length(df$income))

# With DRY
se ← function(x) sd(x) / sqrt(length(x))

# Now we can reuse the function for any column
se_age ← se(df$age)
se_income ← se(df$income)
```


Helper Functions

- Breaking (**decompose**) up Large Functions
 - Limitations of lengthy, complex functions
 - How small, focused functions increase readability and testing ease
- Can make help to DRY but also has value to itself
- When and How to Use Helper Functions
 - Define helper functions to handle subtasks within a larger function

Helper Functions

Example

```
sum_of_squares ← function(a, b) {  
  (a * a) + (b * b)  
}
```

```
sum_of_squares(3, 5)
```

```
#> [1] 34
```

Helper Functions

Example

```
square ← function(x) {  
  x * x  
}  
  
sum_of_squares ← function(a, b) {  
  square(a) + square(b)  
}  
  
sum_of_squares(3, 5)  
#> [1] 34
```

Helper Functions

libminer

```
lib_summary ← function(sizes = FALSE) {  
  pkgs ← utils::installed.packages()  
  pkg_tbl ← table(pkgs[, "LibPath"])  
  pkg_df ← as.data.frame(pkg_tbl, stringsAsFactors = FALSE)  
  names(pkg_df) ← c("Library", "n_packages")  
  
  if (sizes) {  
    pkg_df$lib_size ← map_dbl(  
      pkg_df$Library,  
      ~ sum(fs::file_size(fs::dir_ls(.x, recurse = TRUE))),  
    )  
  }  
  pkg_df  
}
```

 **Your Turn**

check() 

Commit your changes 

Push to GitHub 

Arguments

- What Are Function Arguments?
 - Values passed to a function that specify how it should operate
 - Allow customization and flexibility in function behaviour
- Types of Arguments
 - Required arguments: Must be provided when the function is called
 - Default arguments: Have preset values; can be overridden
 - Variable arguments: Use ... to accept a flexible number of inputs

Default Argument Values

- Why Use Default Arguments?
 - Adds flexibility by setting typical values that simplify function calls.
 - Allow functions to handle common cases without needing all arguments specified every time.
 - Useful in cases where arguments have default values that cover frequent use-cases

Arguments

Refactor to use functions

```
greet ← function(name, greeting = "Hello") {  
  paste(greeting, name)  
}
```

```
# Usage
```

```
greet(name = "Andy")
```

```
#> [1] "Hello Andy"
```

```
greet(name = "Andy", greeting = "Hi")
```

```
#> [1] "Hi Andy"
```


Default Argument Values

Setting Defaults

- Use defaults for parameters that often remain the same, making it easier to call the function without redundant inputs.
- Avoid defaults that might hide potential issues (e.g., setting a default that could lead to silent errors if overlooked).
- Document defaults clearly, so it's clear what will happen if arguments are left unspecified.

```
#' @param name a person's name.  
#' @param greeting the greeting you wish to choose. defaults to  
"Hello"
```

Input Checking and Validation

Why Input Checking Matters

- Prevents Unexpected Errors: Ensures that the function receives valid inputs, reducing the likelihood of errors.
- Improves Code Robustness: Increases confidence in code behaviour
- Helps make failures intentional by catching errors early in the process, making debugging easier by validating assumptions

Tools for Input Checking

- Packages: `assertthat`, `check`, `validate`
- Built-in functions: `stop`, `is.numeric`, `is.character`

Input Checking and Validation

An example

```
calculate_area ← function(length, width) {  
  area ← length * width  
  area)  
}  
calculate_area(4, 2)  
#> [1] 8
```

Input Checking and Validation

An example - bad inputs!

```
calculate_area(-5, 2)
```

```
#> [1] -10
```

```
calculate_area('5', 2)
```

```
#> Error in length * width: non-numeric argument to binary operator
```

Input Checking and Validation

An example - bad inputs!

```
calculate_area ← function(length, width) {  
  if (length ≤ 0 || width ≤ 0) {  
    stop("Length and width must be positive numbers.", call. = FALSE)  
  }  
  area ← length * width  
  area  
}  
calculate_area(-5, 2)  
#> Error: Length and width must be positive numbers.
```

Input Checking and Validation

An example - bad inputs!

```
calculate_area('5', 2)
```

```
#> Error in length * width: non-numeric argument to binary operator
```

 **Your Turn**

Quickly back to helper functions

```
validate_inputs ← function(length, width) {  
  if (length ≤ 0 || width ≤ 0) {  
    stop("Length and width must be positive numbers.", call. = FALSE)  
  }  
  if (!is.numeric(length) || !is.numeric(width)) {  
    stop("Length and width must be a number.", call. = FALSE)  
  }  
}
```

Quickly back to helper functions

```
calculate_area ← function(length, width) {  
  validate_inputs(length, width)  
  area ← length * width  
  area  
}  
  
calculate_area(-5, 2)  
#> Error: Length and width must be positive numbers.  
  
calculate_area('5', 2)  
#> Error: Length and width must be a number.
```

 **Your Turn**

libminer::Input Checking and Validation

```
lib_summary ← function(sizes = FALSE) {  
  pkg_df ← lib()  
  pkg_df ← table(pkg_df$LibPath)  
  names(pkg_df) ← c("Library", "n_packages")  
  
  if (sizes) {  
    pkg_df ← calculate_sizes(pkg_df)  
  }  
  pkg_df  
}
```

Code Style of Readability

Why should you care about code readability?

Facilitates Teamwork

- Clear and consistent code makes it easier for team members to understand each other's work, enhancing collaboration.

Enhances Maintainability

- Readable code simplifies debugging and updates, making it easier to maintain and extend functionality.

Reduces cognitive load

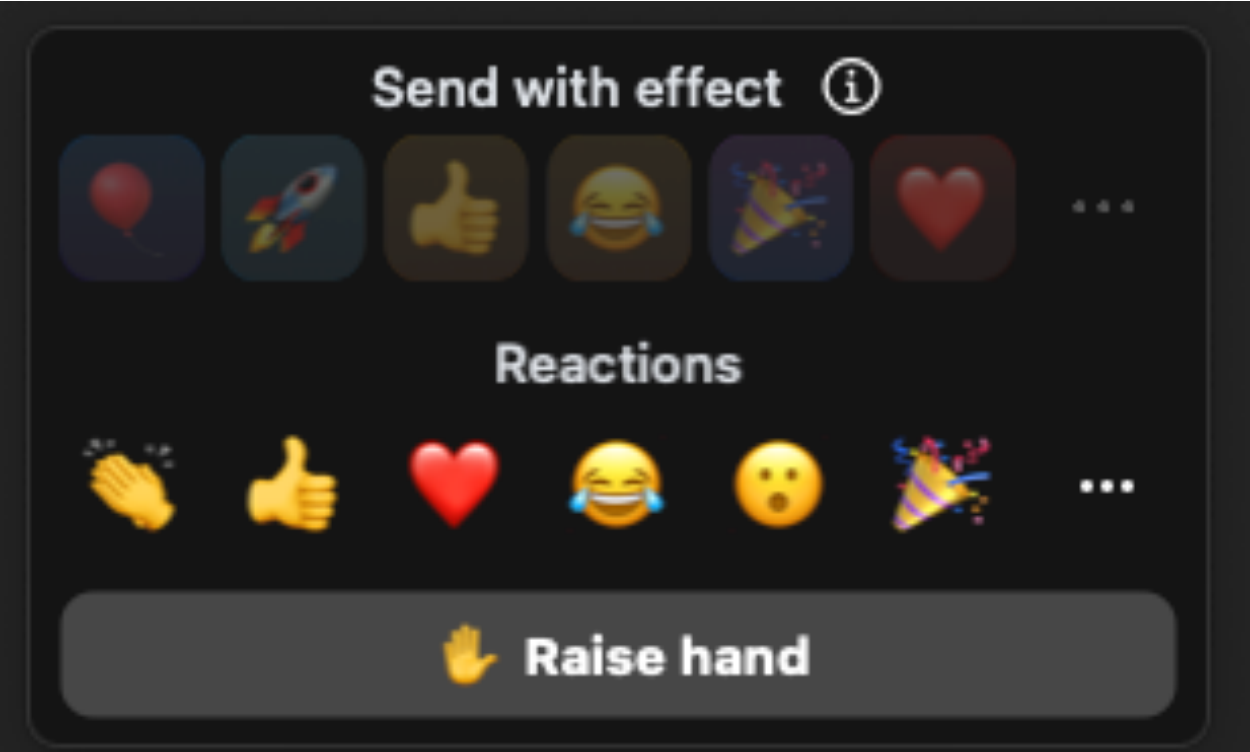
- Well-structured code reduces the time spent reading and understanding code, allowing developers to focus on problem-solving.

Day two complete!

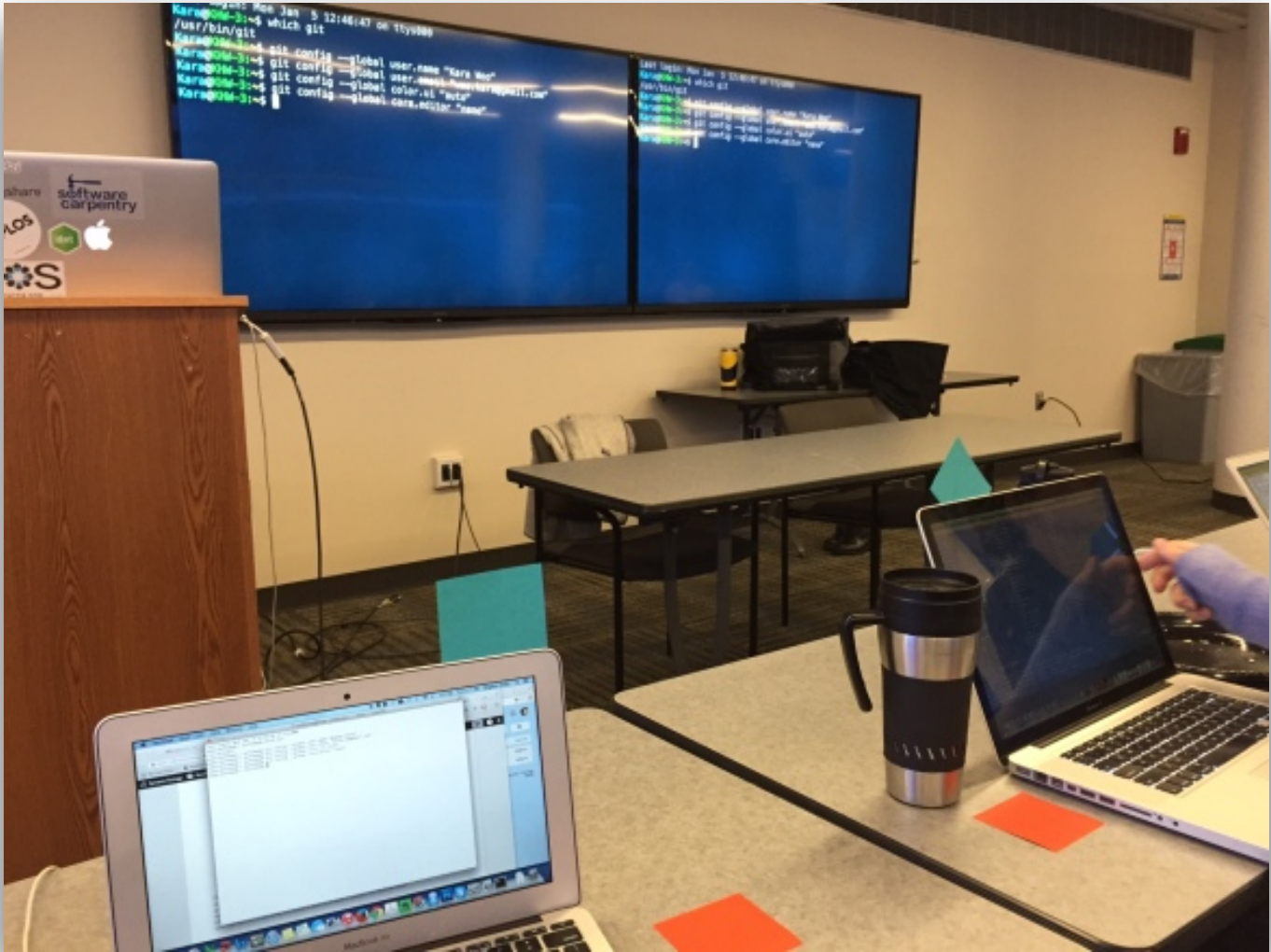
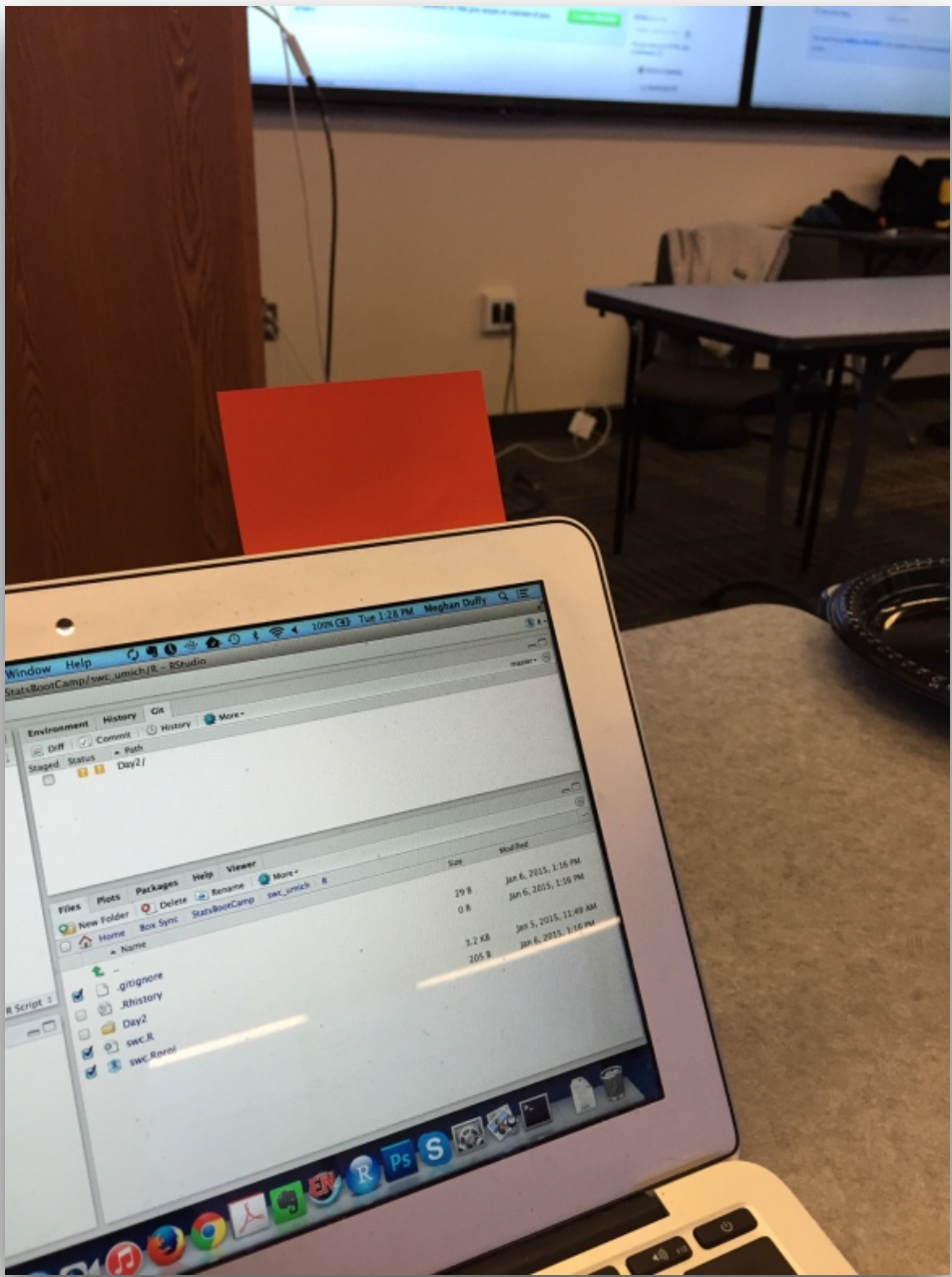
Welcome back to day three

Sticky Notes

I'm stuck, please send help



I'm good, let's move on



Day 3: Schedule and Learning Objectives

2024-11-06		
08:00 - 09:20	Website Creation & Using the Tidyverse in Your Package	80 min
09:20 - 09:35	Break	15 min
09:35 - 10:55	Using the Tidyverse in Your Package part 2	80 min
10:55 - 11:10	Break	15 min
11:10 - 12:30	Communicating with your Users & Distribution & General Discussion	80 min

Website Creation and Long Form Documentation

Why make a website?

- Professional Documentation
 - Offers a polished interface that enhances the credibility of your package.
- Improved Accessibility
 - Makes documentation easily accessible online, allowing potential users to quickly find information, tutorials, and examples.
- Enhanced Discoverability
 - Increases visibility of your package, helping users easily learn about its features and functionalities.
- It's easy!

use_pkgdown_github_pages()

pkgdown.r-lib.org/

- Automatically creates a website using pkgdown:
 - Function documentation
 - Dataset documentation
 - Vignettes (and articles)
 - README and NEWS
- Sets up a GitHub Action to deploy on GitHub Pages



 **Your Turn**

Vignettes

Long-form documentation

- `use_vignette()` is a good way to initiate a vignette
- `use_article()` initiates a article that only exists in the pkgdown website
- Today these functions create an `.Rmd` document
- pkgdown just gained experimental support for Quarto (`.qmd`) and usethis will adapt soon
- A vignette or article focuses on how to solve a target problem, versus documenting a specific function

Vignettes

Long-form documentation

- Vignettes can only use packages you formally depend on, i.e. they appear in **Imports** or **Suggests**
- Help topics live as **.Rd** files in a source package, but there's no equivalent for rendered vignettes
 - Creates some inherent awkwardness in vignette workflow
- Similar to examples, **R CMD check** treats vignettes like big, weird tests.
 - Creates tension between showing realistic usage and ensuring the code can be evaluated, 100% of the time and quickly.
 - Challenges: wrapping an external service, long-running code, etc.

Eval chunk option

- `eval` chunk option controls whether code is evaluated
- Similar to `@examplesIf`, you can call a function to determine if it's safe to evaluate a chunk
- `eval = requireNamespace("somedependency")`
- `eval = !identical(Sys.getenv("SOME_THING_YOU_NEED"), "")`
- `eval = file.exists("credentials-you-need")`
- You can use `eval` local to individual chunks or set it globally in a setup chunk.

```
```{r setup, include = FALSE}
can_decrypt ← gargle:::secret_can_decrypt("googlesheets4")
knitr::opts_chunk$set(
 collapse = TRUE,
 comment = "#>",
 error = TRUE,
 eval = can_decrypt
)
```

```{r eval = !can_decrypt, echo = FALSE, comment = NA}
message("No token available. Code chunks will not be evaluated.")
```

```{r index-auth, include = FALSE}
googlesheets4:::gs4_auth_docs()
```
```

Using the tidyverse in your
package

Learning Objectives

- At the end of this section you will be able to:
 - build functions that call tidyverse functions, using bare-name arguments:
 - designed for interactive use, i.e. like `dplyr::filter()`
 - understand terms: **data masking**, **tidy select**, **dynamic dots**

Tidy Evaluation

Motivation

- Tidy eval (and non-standard evaluation generally in R) exists so that we can refer to data columns directly using bare names.
 - `dplyr::filter(mtcars, cyl = 4)`

VS

- `mtcars[mtcars$cyl = 4,]`
- It makes things easier if you are working interactively (it allows *indirection*).
- It makes things more interesting if you are writing functions.

Welcome to *more interesting*.

Families of Tidy-eval Functions

- **data-masking:** compute on values within a dataset:

```
mtcars |>
```

```
  dplyr::mutate(wt_kg = wt * 1000 / 2.2)
```

- **tidy-select:** specify columns within a dataset

```
mtcars |>
```

```
  dplyr::select(starts_with("w"))
```

Families of Tidy-eval Functions

- **data-masking:** evaluate variables in context of data frame:
`arrange()`, `count()`, `filter()`, `group_by()`, `mutate()`,
and `summarize()`
- **tidy-select:** specify columns within a data frame
`across()`, `relocate()`, `rename()`, `select()`, and `pull()`

Data Masking

Two types of variables

- `mtcars`, `starwars` - datasets, variables available in the environment
 - **environment-variables**
- `cyl`, `mpg`, `homeworld`, and `species` - columns in datasets, variables available in data
 - **data-variables**

Data Masking

Evaluate variables in the context of a data frame

```
starwars[starwars$homeworld == "Naboo" & starwars$species == "Human", ]
```

```
starwars |>
```

```
  filter(homeworld == "Naboo", species == "Human")
```

```
mtcars |>
```

```
  group_by(cyl) |>
```

```
  summarise(n = n(), mpg = mean(mpg))
```

Data Masking

Identify env-variables vs data-variables

```
starwars[starwars$homeworld == "Naboo" & starwars$species == "Human", ]
```

```
starwars |>
```

```
  filter(homeworld == "Naboo", species == "Human")
```

```
mtcars |>
```

```
  group_by(cyl) |>
```

```
  summarise(n = n(), mpg = mean(mpg))
```

Programming with data variables

```
var_summary <- function(data, var) {  
  data |>  
    summarise(  
      min = min(var),  
      max = max(var)  
    )  
}
```

```
mtcars |>  
  group_by(cyl) |>  
  var_summary(mpg)  
#> Error in `summarise()`:  
#> i In argument: `min = min(var)`.  
#> i In group 1: `cyl = 4`.  
#> Caused by error:  
#> ! object 'mpg' not found
```

Programming with data variables

{{ embrace }} data variables as function arguments

```
var_summary <- function(data, var) {  
  data |>  
    summarise(  
      min = min({{ var }}),  
      max = max({{ var }})  
    )  
}
```

```
mtcars |>  
  group_by(cyl) |>  
  var_summary(mpg)  
#> # A tibble: 3 × 3  
#>   cyl   min   max  
#>   <dbl> <dbl> <dbl>  
#> 1     4  21.4  33.9  
#> 2     6  17.8  21.4  
#> 3     8  10.4  19.2
```

Programming with data variables

Use `.data$[[]]` for function arguments passed as characters

```
var_summary <- function(data, var) {  
  data |>  
    summarise(  
      min = min(.data[[var]]),  
      max = max(.data[[var]])  
    )  
}
```

```
mtcars |>  
  group_by(cyl) |>  
  var_summary("mpg")  
#> # A tibble: 3 × 3  
#>   cyl   min   max  
#>   <dbl> <dbl> <dbl>  
#> 1     4  21.4  33.9  
#> 2     6  17.8  21.4  
#> 3     8  10.4  19.2
```


Programming with data variables

Use `.data$` pronoun for variables you know about

```
big_cars_summary <- function(var) {  
  mtcars |>  
  filter(.data$cyl >= 6) |>  
  group_by(.data$cyl) |>  
  summarise(  
    n = n(),  
    mean = mean({{ var }}),  
  )  
}
```

```
big_cars_summary(displ)  
#> # A tibble: 2 × 3  
#>   cyl      n  mean  
#>   <dbl> <int> <dbl>  
#> 1     6     7  183.  
#> 2     8    14  353.
```

Write a function that summarizes one or more variables in **starwars** by any grouping variable

```
library(dplyr)
head(starwars)
#> # A tibble: 6 × 14
#>   name          height  mass hair_color skin_color eye_color birth_year sex  gender
#>   <chr>         <int> <dbl> <chr>         <chr>    <chr>      <dbl> <chr> <chr>
#> 1 Luke Sky...    172    77 blond         fair     blue        19    male masculi...
#> 2 C-3PO          167    75 <NA>          gold     yellow      112   none masculi...
#> 3 R2-D2          96     32 <NA>          white, bl... red        33    none masculi...
#> 4 Darth Va...   202   136 none          white     yellow      41.9  male masculi...
#> 5 Leia Org...   150    49 brown         light    brown        19    fema... femin...
#> 6 Owen Lars     178   120 brown, gr... light     blue        52    male masculi...
#> # i 5 more variables: homeworld <chr>, species <chr>, films <list>,
#> #   vehicles <list>, starships <list>
```

* Do this in a file that is not inside **R/**

* Add path to **.Rbuildignore**



Your Turn

Write a function that summarizes one or more variables in **starwars** by any grouping variable

```
height_sum <- function(data, group_var) {  
  data |>  
    dplyr::group_by({{ group_var }}) |>  
    dplyr::summarise(  
      n = dplyr::n(),  
      mean_height = mean(.data$height)  
    )  
}
```

```
height_sum(starwars, hair_color)
```

Modify that function to take > 1 grouping variables

- Hint: '...' can get passed through unaltered.

 **Your Turn**

Modify that function to take > 1 grouping variables

```
height_sum <- function(data, ...) {  
  data |>  
    dplyr::group_by(...) |>  
    dplyr::summarise(  
      n = dplyr::n(),  
      mean_height = mean(.data$height)  
    )  
}
```

```
height_sum(starwars, hair_color, eye_color)
```

Data Masking - Summary

- dplyr: `arrange()`, `filter()`, `group_by()`, `mutate()`, and `summarise()`;
- ggplot2: `aes()`
- Data-variable is passed as a **function argument** x:
 - `arrange(df, {{ x }})`
- You (as developer) know the **data-variable** is x:
 - `arrange(df, .data$x)`
- Name of data-variable is a **character string** in an **env-variable** x:
 - `arrange(df, .data[[x]])`
- Remember `...` can be passed through - no special treatment required.

Name injection with dynamic dots

Dynamically create new variable names

- Generate names programmatically by using `:=` instead of `=`
- Two ways:
 - From a variable containing a string using `{glue}` syntax
 - From a data variable using `{{embrace}}` syntax
- To use in your package:
`usethis::use_import_from("rlang", ":=")`

Name injection with dynamic dots

Dynamically create new variable names

```
var_summary <- function(data, var, new_col_name) {  
  data |>  
    summarise(  
      "{new_col_name}" := min({{ var }})  
    )  
}
```

```
mtcars |>  
  group_by(cyl) |>  
  var_summary(mpg, new_col_name = "min_mpg")  
#> # A tibble: 3 × 2  
#>   cyl min_mpg  
#>   <dbl> <dbl>  
#> 1     4    21.4  
#> 2     6    17.8  
#> 3     8    10.4
```


Name injection with dynamic dots

Dynamically create new variable names

```
var_summary <- function(data, var) {  
  data |>  
    summarise(  
      "{{var}}_min" := min({{ var }})  
    )  
}
```

```
mtcars |>  
  group_by(cyl) |>  
  var_summary(mpg)  
#> # A tibble: 3 × 2  
#>   cyl mpg_min  
#>   <dbl> <dbl>  
#> 1     4    21.4  
#> 2     6    17.8  
#> 3     8    10.4
```

Write a function that summarizes **starwars**, and dynamically creates new column names

```
library(dplyr)
head(starwars)
#> # A tibble: 6 × 14
#>   name          height  mass hair_color skin_color eye_color birth_year sex  gender
#>   <chr>         <int> <dbl> <chr>      <chr>      <chr>      <dbl> <chr> <chr>
#> 1 Luke Sky...    172    77 blond      fair        blue        19    male masculi...
#> 2 C-3PO          167    75 <NA>      gold        yellow       112   none masculi...
#> 3 R2-D2           96    32 <NA>      white, bl... red         33    none masculi...
#> 4 Darth Va...   202   136 none      white       yellow       41.9  male masculi...
#> 5 Leia Org...   150    49 brown     light       brown        19    fema... femini...
#> 6 Owen Lars     178   120 brown, gr... light       blue         52    male masculi...
#> # i 5 more variables: homeworld <chr>, species <chr>, films <list>,
#> #   vehicles <list>, starships <list>
```

* Do this in a file that is not inside **R/**

* Add path to **.Rbuildignore**



Your Turn

Write a function that summarizes **starwars**, and dynamically creates new column names

```
dynamic_sum <- function(data, group_var, sum_var) {  
  data |>  
    dplyr::group_by({{ group_var }}) |>  
    dplyr::summarise(  
      n = dplyr::n(),  
      "mean_{{sum_var}}" := mean({{ sum_var }})  
    )  
}
```

```
dynamic_sum(starwars, hair_color, mass)
```

Tidy Selection

Select columns in a data frame

- Using bare names, or tidyselect helpers:
 - `select(df, 1)`: selects the first column
 - `select(df, last_col())` selects the last column.
 - `select(df, c(a, b, c))` selects columns `a`, `b`, and `c`.
 - `select(df, starts_with("a"))` selects all columns whose name starts with “a”
 - `select(df, ends_with("z"))` selects all columns whose name ends with “z”.
 - `select(df, where(is.numeric))` selects all numeric columns.
- Using character vectors:
 - `select(df, all_of(c("a", "b", "c")))` strictly selects all specified
 - `select(df, any_of(c("a", "b", "c")))` selects specified columns if they exist

dplyr::across()

- Used inside *data-masking* verbs, e.g., `summarize()`
- Specify:
 - Which columns to consider, using *tidy-select*
 - What functions to apply, if any

```
mtcars |>
  dplyr::group_by(cyl) |>
  dplyr::summarise(
    dplyr::across(c("mpg", "disp"), mean)
  )
#> # A tibble: 3 × 3
#>   cyl   mpg  disp
#>   <dbl> <dbl> <dbl>
#> 1     4  26.7  105.
#> 2     6  19.7  183.
#> 3     8  15.1  353.
```

all_of(); any_of()

- Used to disambiguate external vector (env-var) of column names
- `all_of()` is strict

```
cols <- c("mpg", "disp", "xyz")
mtcars |>
  dplyr::group_by(cyl) |>
  dplyr::summarise(
    dplyr::across(dplyr::all_of(cols), mean)
  )
#> Error in `dplyr::summarise()`:
#> i In argument: `dplyr::across(dplyr::all_of(cols), mean)`.
#> ! Can't subset elements that don't exist.
#> ✘ Element `xyz` doesn't exist.
```

all_of(); any_of()

- Used to disambiguate external vector (env-var) of column names
- `any_of()` is permissive

```
cols <- c("mpg", "disp", "xyz")
mtcars |>
  dplyr::group_by(cyl) |>
  dplyr::summarise(
    dplyr::across(dplyr::any_of(cols), mean)
  )
#> # A tibble: 3 × 3
#>   cyl   mpg  disp
#>   <dbl> <dbl> <dbl>
#> 1     4  26.7  105.
#> 2     6  19.7  183.
#> 3     8  15.1  353.
```

Tidyversify libminer

- Use:
 - tidyverse functions
 - Data Masking with `{{ }}`
 - Data Masking with `.data`
 - ...
 - `across()` and `any_of()`
- Update tests
- Update documentation

 **Our turn**

Break Time!

Communicating with your user

Signaling Conditions

Three conditions

- **Errors**

- most severe
- no way for a function to continue and execution must stop.

- **Warnings**

- in between errors and message
- something has gone wrong or is unexpected but the function has executed

- **Messages**

- mildest
- Inform users that some action has been performed on their behalf.

Signaling Conditions

Three conditions

- **Errors**

- `stop("You provided an invalid input")`

- **Warnings**

- `warning("Unknown value converted to NA")`

- **Messages**

- `message("Reticulating splines... this may take a while")`

Conditions are better than `cat()`

- They inherently indicate the severity of the message
- They can be handled in special ways with things like `try()`, `tryCatch()`, etc.
- Warnings and messages can be silenced by users:

```
chatty <- function() {  
  message("Hello there!")  
  warning("You have been warned!")  
}
```

```
chatty()  
#> Hello there!  
#> Warning in chatty(): You have been warned!
```

```
suppressMessages(chatty())  
#> Warning in chatty(): You have been warned!
```

```
suppressWarnings(chatty())  
#> Hello there!
```

beautiful command line interfaces

With the `{cli}` package

- Build a CLI using semantic elements: headings, bullet lists, etc.
- All cli text can contain interpreted string literals, via the glue package.
- Support for pluralization and list concatenation.
- Inline text formatting.
- Text wrapping.
- Theming via a CSS-like language.
- Progress bars from R and C code.
- Emit conditions -- messages, warnings, and errors -- with rich text formatting.

usethis's UI is built with cli

Character vector of lines.

Names control the type of bullet.

```
> ui_bullets(c(
+   "v" = "A great success!",
+   "_" = "Something you need to do.",
+   "x" = "Bad news.",
+   "i" = "The more you know.",
+   " " = "I'm just here for the indentation.",
+   "No indentation at all. Not used much in usethis."
+ ))
✓ A great success!
□ Something you need to do.
✗ Bad news.
i The more you know.
  I'm just here for the indentation.
No indentation at all. Not used much in usethis.
```

cli's greatest hits

- Single line alerts:
 - `cli_alert_success()`, `cli_alert_danger()`, `cli_alert_info()`, `cli_alert_warning()`
- Multi-line bullet lists:
 - `cli_bullets()`
- Conditions with all the cli (+ rlang) goodness
 - `cli_inform()`, alternative for `message()` or `rlang::inform()`
 - `cli_warn()`, alternative for `warning()` or `rlang::warn()`
 - `cli_abort()`, alternative for `stop()` or `rlang::abort()`

* All cli outputs are `message` conditions and can be handled as such

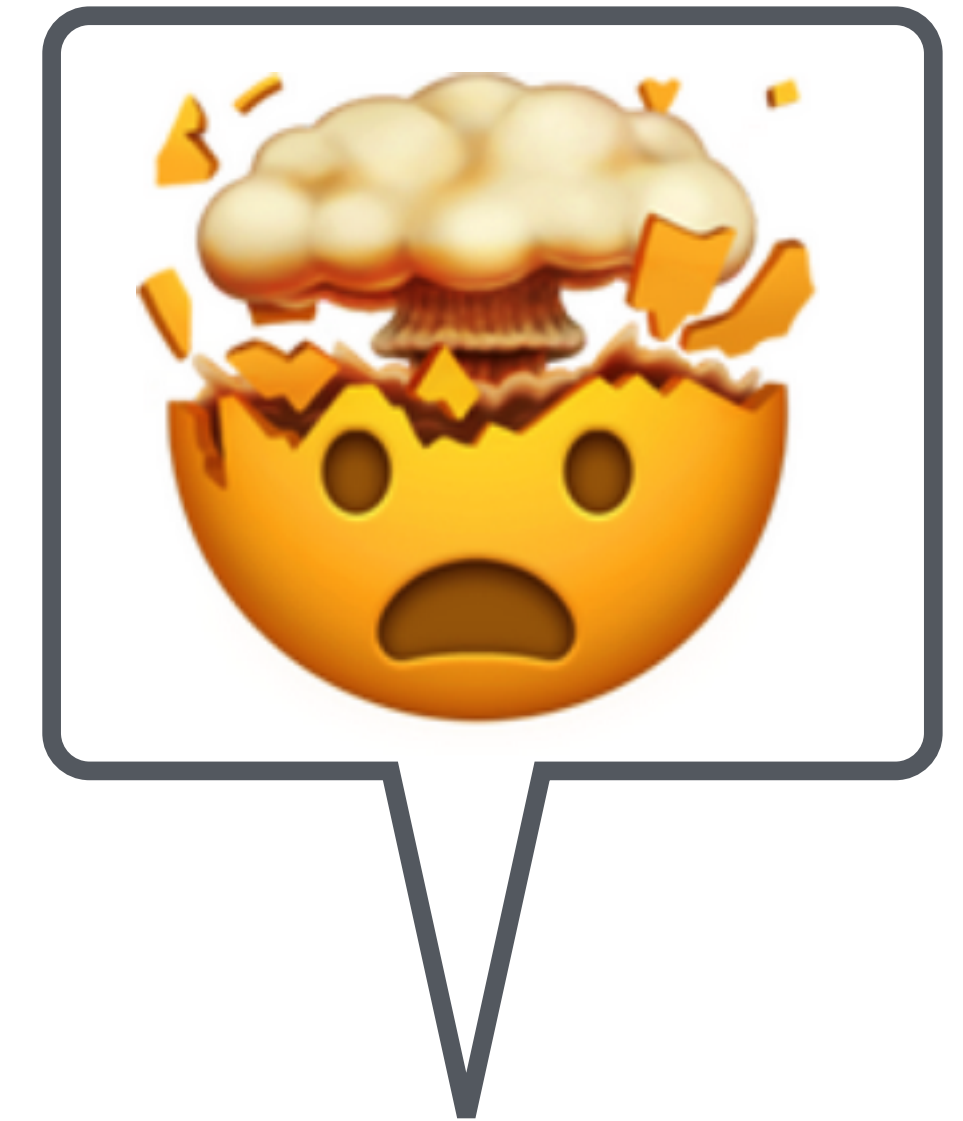
Interpolation

Code inside curly braces is evaluated, a la the glue package.

```
> pkgs <- "pkg1"  
> cli_text("Will install the {pkgs} package")  
Will install the pkg1 package
```

Pluralization

```
> nfiles <- 0; cli_text("Found {nfiles} file{?s}")  
Found 0 files  
> nfiles <- 1; cli_text("Found {nfiles} file{?s}")  
Found 1 file  
> nfiles <- 2; cli_text("Found {nfiles} file{?s}")  
Found 2 files
```



Pluralization and concatenation

```
> pkgs <- "pkg1"  
> cli_text("Will install the {pkgs} package{?s}")  
Will install the pkg1 package
```

```
> pkgs <- c("pkg1", "pkg2", "pkg3")  
> cli_text("Will install the {pkgs} package{?s}")  
Will install the pkg1, pkg2, and pkg3 packages
```

Inline text formatting

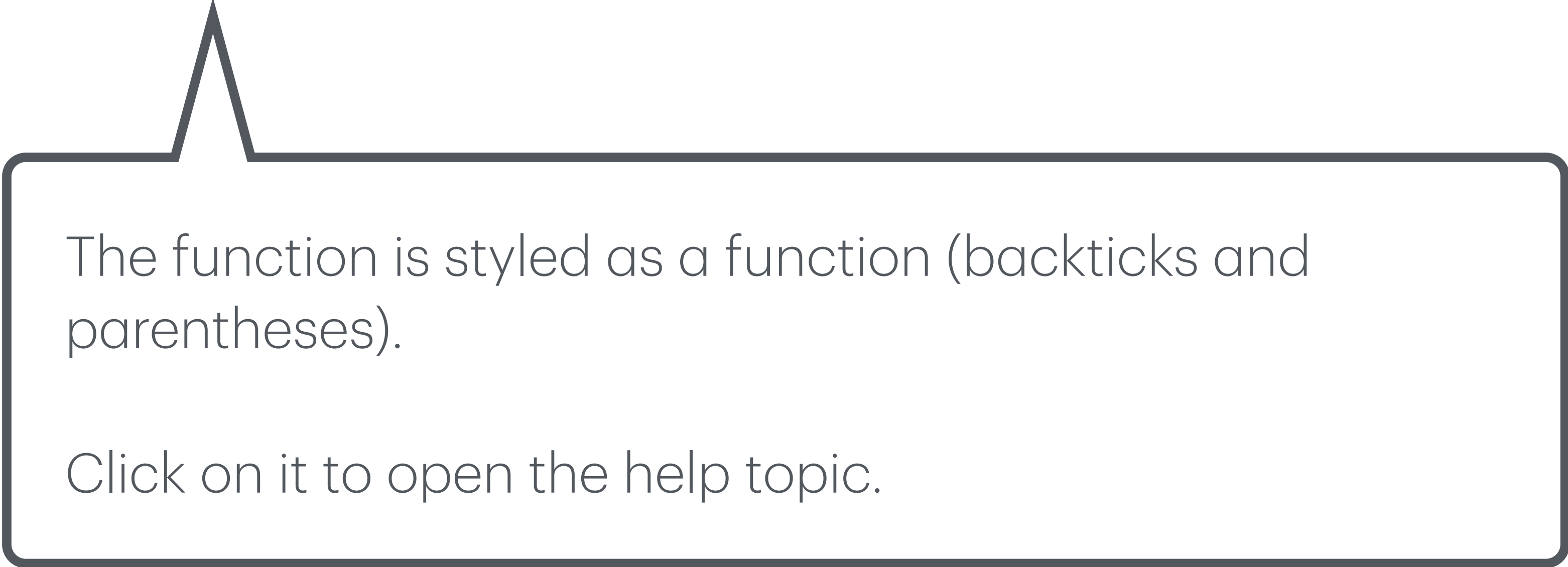
"{.thm text}"

```
cli_bullets(c(  
  "*" = "{.emph Emphasized} text",  
  "*" = "{.strong Strong} importance",  
  "*" = "A piece of code: {.code sum(a) / length(a)}",  
  "*" = "A package name: {.pkg cli}",  
  "*" = "A function name: {.fn cli_text}",  
  "*" = "A keyboard key: press {.kbd ENTER}",  
  "*" = "A file name: {.file /usr/bin/env}",  
  "*" = "An email address: {.email bugs.bunny@acme.com}",  
  "*" = "A URL: {.url https://acme.com}",  
  "*" = "An environment variable: {.envvar R_LIBS}",  
  "*" = "Some {.field field}"  
))
```

- *Emphasized* text
- **Strong** importance
- A piece of code: ``sum(a) / length(a)``
- A package name: `cli`
- A function name: ``cli_text()``
- A keyboard key: press `[ENTER]`
- A file name: `/usr/bin/env`
- An email address: `bugs.bunny@acme.com`
- A URL: `<https://acme.com>`
- An environment variable: ``R_LIBS``
- Some `field`

Help topic links

```
> cli_text("Try out the {.fun cli::cli_abort} function")  
Try out the `cli::cli_abort()` function
```

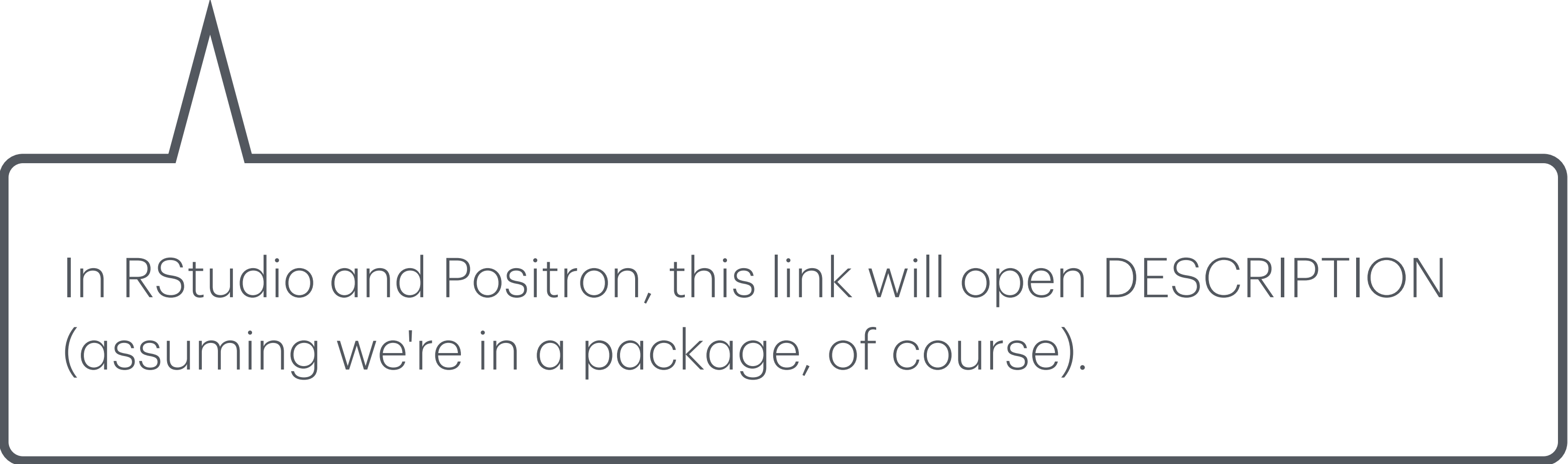


The function is styled as a function (backticks and parentheses).

Click on it to open the help topic.

File hyperlinks

```
> cli_text("A file name: {.file DESCRIPTION}")  
A file name: DESCRIPTION.
```



In RStudio and Positron, this link will open DESCRIPTION (assuming we're in a package, of course).

URLs

```
> cli_text(  
  "See the cli homepage at {.url https://cli.r-lib.org} for details"  
)
```

See the cli homepage at <https://cli.r-lib.org> for details.



A clickable hyperlink with no custom link text.

Hyperlinks

Markdown-style hyperlinks

```
> cli_text(  
  "See the {.href [cli homepage](https://cli.r-lib.org)} for details"  
)
```

```
See the cli homepage for details.
```

A clickable hyperlink with custom link text.

Formatting plus interpolation

Use double curly braces

```
> cli_url <- "https://cli.r-lib.org"
> cli_text(
  "See the cli homepage at {.url {cli_url}} for details"
)
```

See the cli homepage at <https://cli.r-lib.org> for details.

Your turn! Some prompts

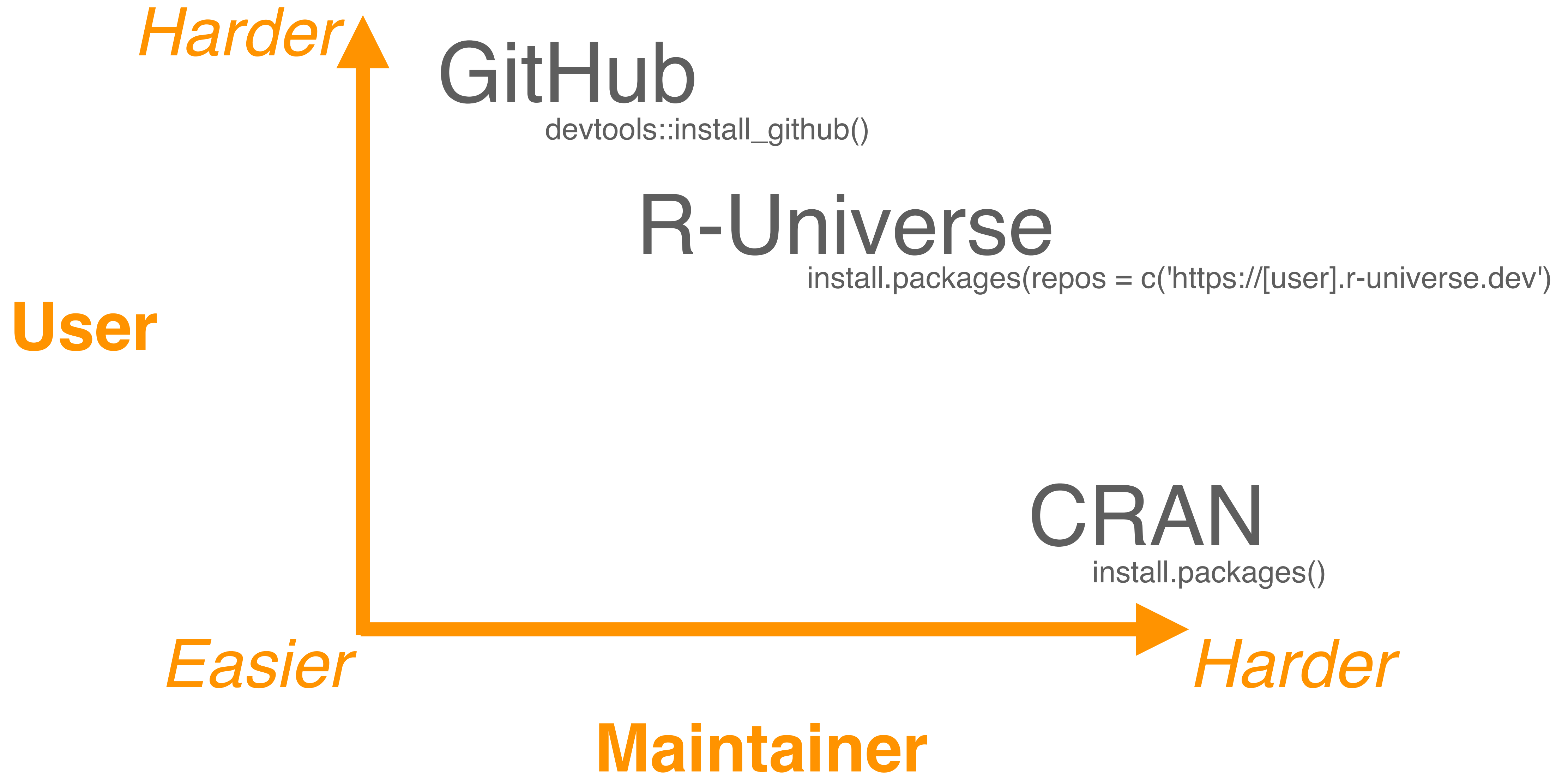
- Choose a package: libminer or one of your own
- Study the existing user interface (which functions are used?).
- Incrementally convert the UI-providing functions (e.g. `message()` or `stop()`) to its cli equivalent.
- Add some more bling
- Look at the cli documentation, especially:
 - cli's articles: <https://cli.r-lib.org/articles/index.html>
 - Help topics in the "Introduction" section here: <https://cli.r-lib.org/reference/index.html>

Break Time!

Sharing your Package



Package Distribution



Releasing your package to CRAN



Releasing to CRAN

TLDR:

- `release()`
 - Runs through an additional list of checks
 - Builds package bundle and submits to CRAN

use_news_md()

Adds a NEWS.md file to your package

- Tracks version numbers
- Tracks user-facing changes between versions

use_cran_comments()

Comments that go along with your submission

```
## R CMD check results
```

```
0 errors | 0 warnings | 1 note
```

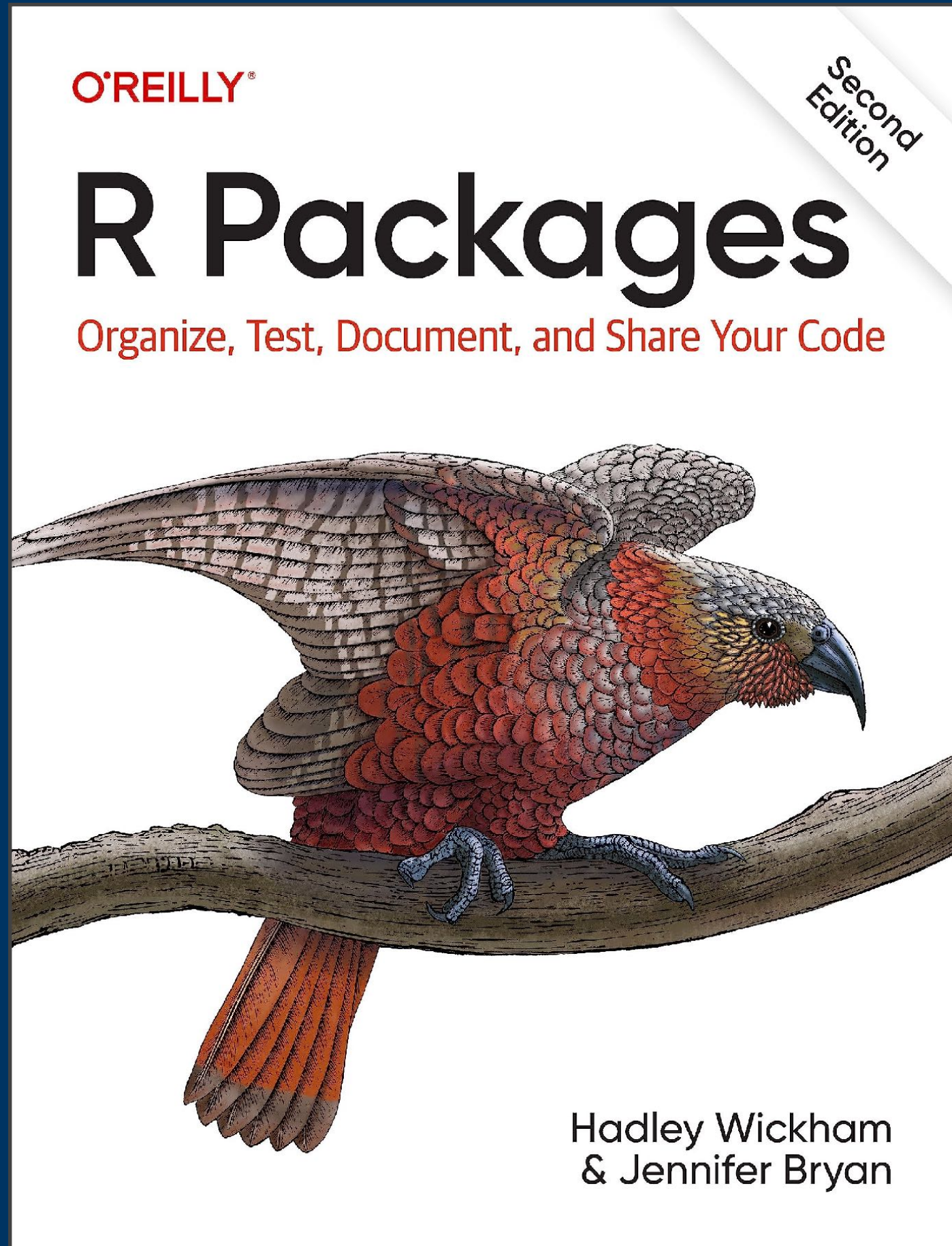
```
* This is a new release.
```

- Most times don't need more than this
 - Note if you are fixing failures seen in CRAN checks
 - Note reverse dependency checks
 - If your submission was rejected, note that it is a resubmission and say how you have addressed the issues
- *There is always one NOTE with initial submission

DFO Package Discussion

Thank You!

Resources



r-pkgs.org

Happy Git and GitHub for the userR

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Connect Git, GitHub, RStudio

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Let's Git started

Still from Heaven King video

Happy Git provides opinionated instructions on how to:

- Install Git and get it working smoothly with GitHub, in the shell and in the RStudio IDE.
- Develop a few key workflows that cover your most common tasks.
- Integrate Git and GitHub into your daily work with R and R Markdown.

happygitwithr.com

Package Development : : CHEAT SHEET

DESCRIPTION devtools

Package Structure

Workflow

Getting Started

DESCRIPTION

NAMESPACE

DESCRIPTION

NAMESPACE

posit

Tidy design principles

Welcome

The goal of this book is to help you write better R code. It has four main components:

- Identifying design **challenges** that often lead to suboptimal outcomes.
- Introducing useful **patterns** that help solve common problems.
- Defining key **principles** that help you balance conflicting patterns.
- Discussing **case studies** that help you see how all the pieces fit together with real code.

While I've called these principles "tidy" and they're used extensively by the tidyverse team to promote consistency across our packages, they're not exclusive to the tidyverse. Think tidy in the sense of tidy data (broadly useful regardless of what tool you're using) not tidyverse (a collection of functions designed with a singular point of view in order to facilitate learning and use).

This book will be under heavy development for quite some time; currently we are loosely aiming for completion in 2025. You'll find many chapters contain disjointed text that mostly serve as placeholders for the authors, and I do not recommend attempting to systematically read the book at this time. If you'd like to follow along with my journey writing this book, and learn which chapters are ready to read, please sign up for my [tid design substack mailing list](#).

1 Unifying principles →

design.tidyverse.org

tidydesign.substack.com

posit.co/resources/cheatsheets/

Attribution

- <https://github.com/posit-conf-2024/pkg-dev>
- <https://cli.r-lib.org/articles/index.html>
- <https://adv-r.hadley.nz/conditions.html>
- <https://rstudio-conf-2022.github.io/build-tidy-tools/>
- <https://dplyr.tidyverse.org/articles/programming.html>
- <https://design.tidyverse.org/>

Course Materials

<https://andyteucher.ca/pkg-dev-dfo-2024-11>

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Survey

Your feedback is crucial! Please complete the post-workshop survey! 🙏

Data from the survey informs curriculum and format decisions for future workshops, and we really appreciate you taking the time to provide it.