rstd.io/pkg-dev

"Just in time" prep



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

* Via devtools, this will also install usethis.

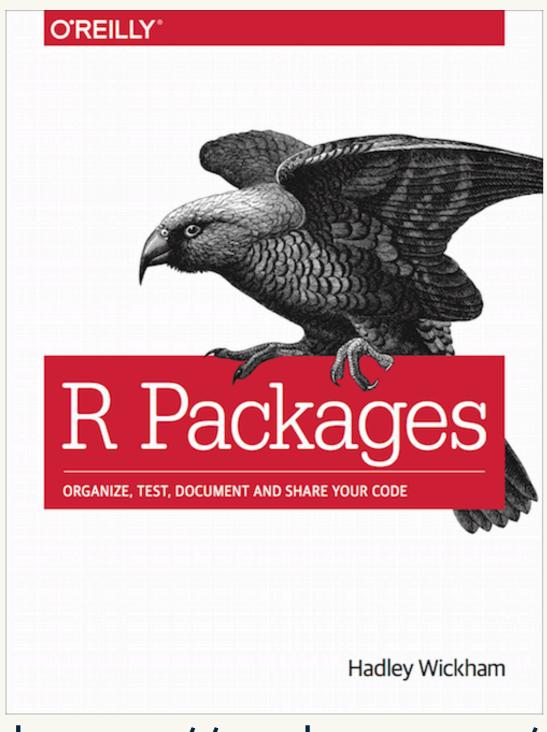
Intro & basic package workflow

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rstd.io/pkg-dev



https://r-pkgs.org/

???

Hadley Wickham

https://principles.tidyverse.org

Your turn

This course is hands-on and, while we're here to help, the best resource may be the person sitting next to you.

Introduce yourself to your neighbours. Who are you and what are you using R for?

This means that you have to work!

Warmup

Get to know your R installation!

Your turn

```
.Library
.libPaths()
installed.packages()
```

```
# Where's your default package library?
# Which libraries are searched for packages?
# How many packages do you have installed?
# Make a frequency table of package Priority.
```

Package = natural unit for distributing R code

base R \approx 14 base + 15 recommended packages ship with all binary distributions of R

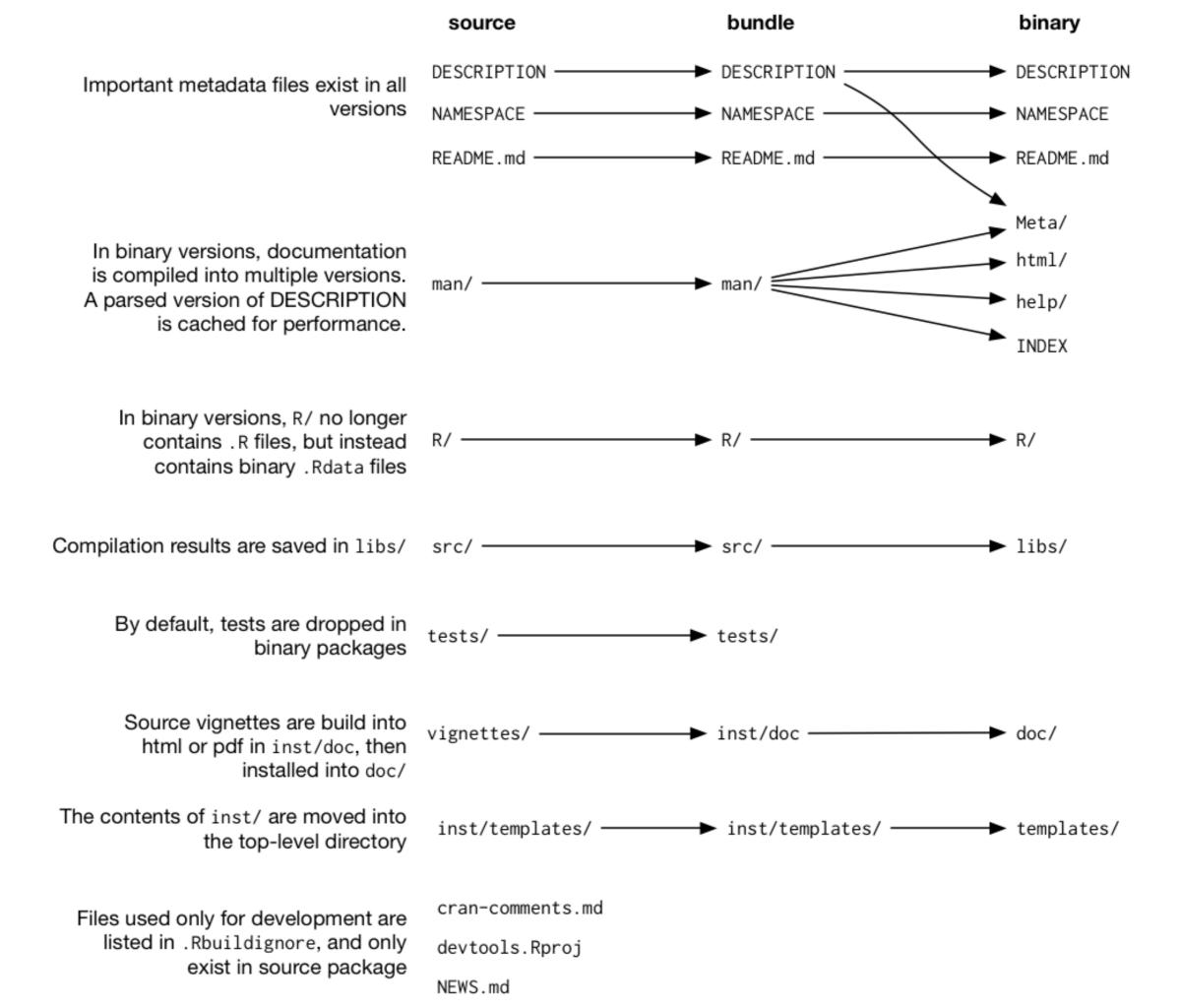
can use right out of the box:

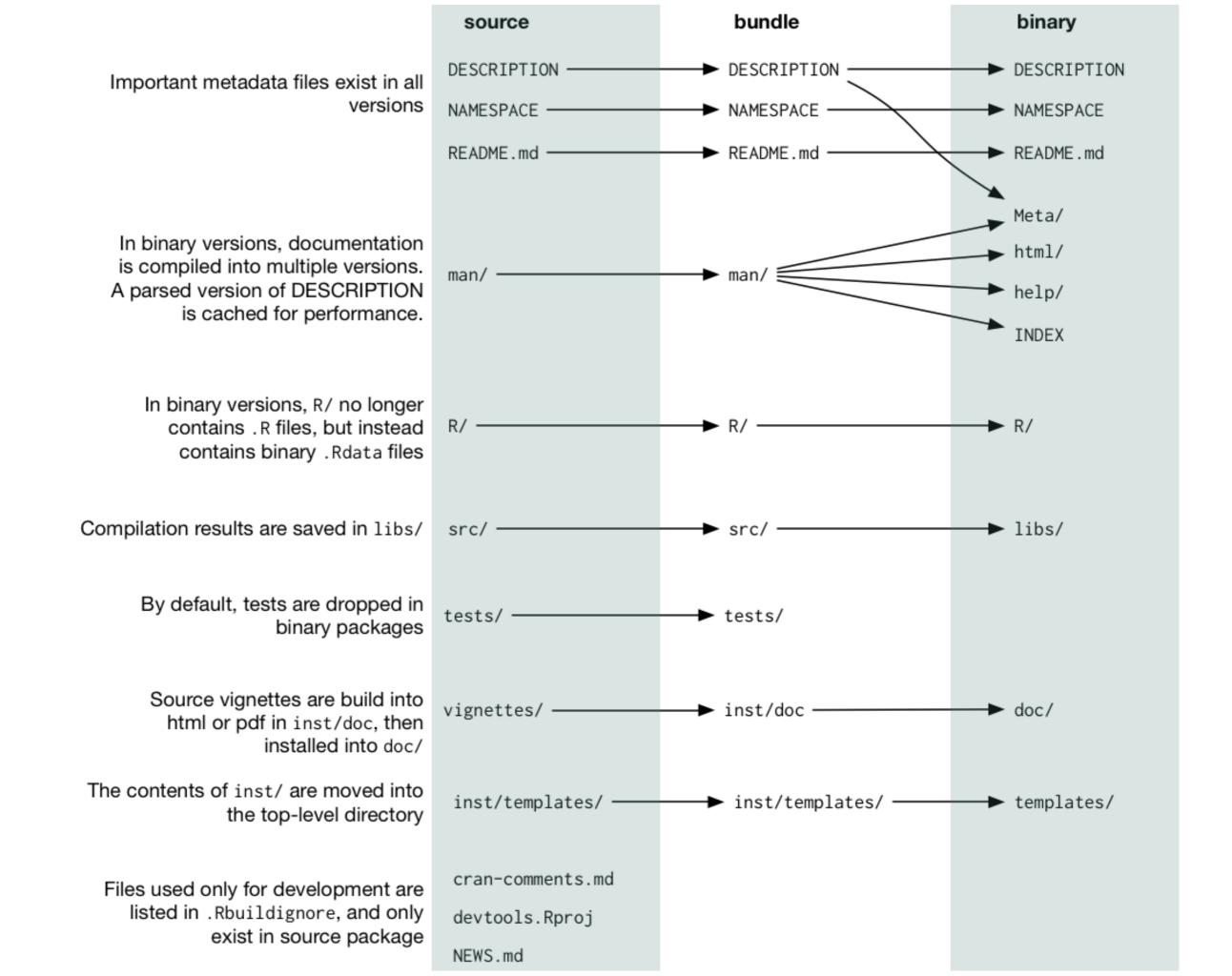
library(lattice)

Package = natural unit for distributing R code

```
CRAN has ~14K additional packages install, then attach: install.packages("devtools") library(devtools)
```

And then there's GitHub ...
install via devtools, then attach:
devtools::install_github("jimhester/lookup")
library(lookup)





A package is a set of conventions that (with the right tools) makes your life easier

foofactors

usethis::create_package()

Your turn

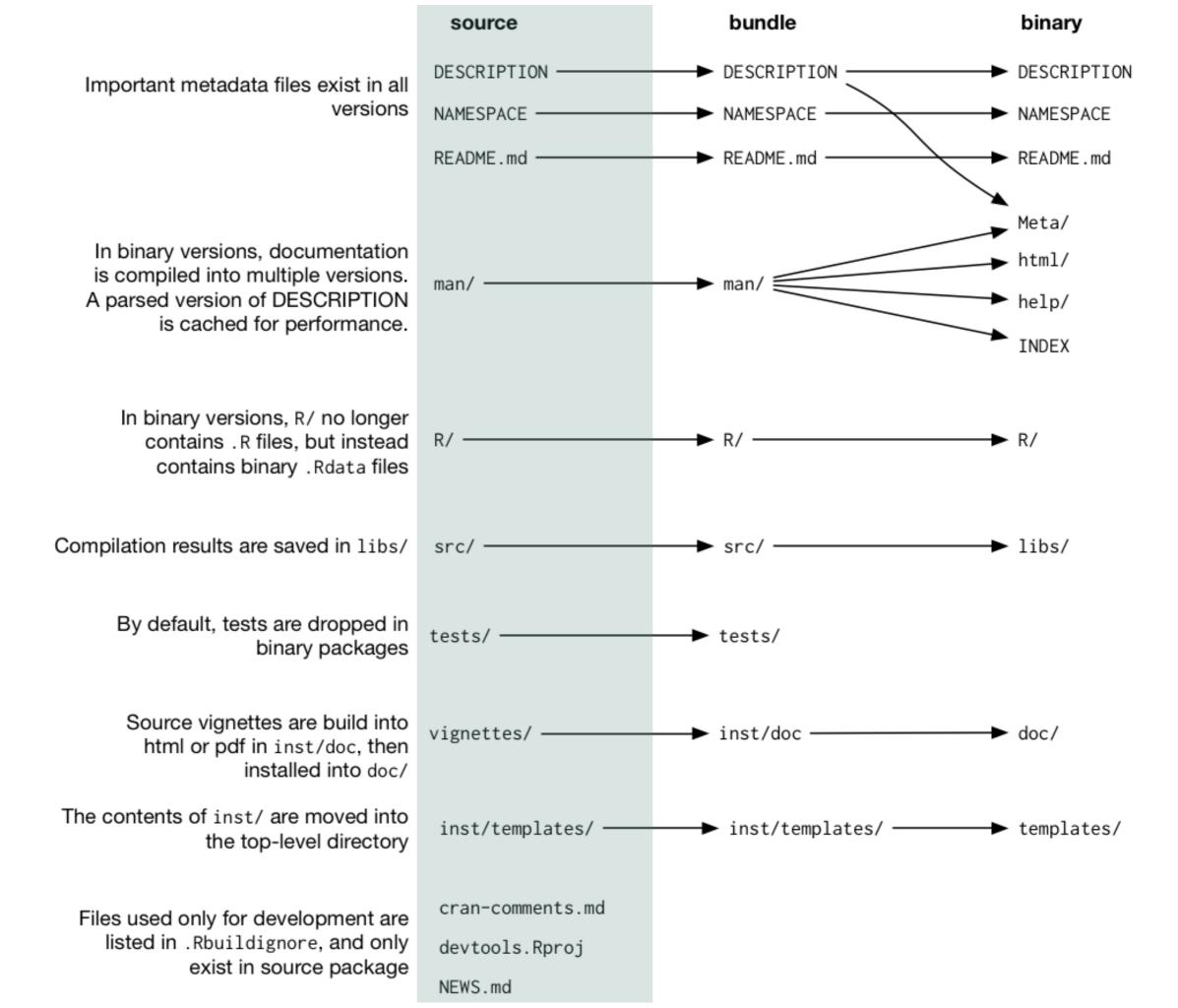
```
# Create a package with:
usethis::create_package("~/Desktop/foofactors")
```

Substitute your preferred location.

- # What files and directories are created? # Compare to source package diagram.
- # FYI, you can also create new project using # RStudio but it has some slight differences.

What does create_package() do?

✔ Creating '/Users/jenny/tmp/foofactors2/' ✓ Setting active project to '/Users/jenny/tmp/foofactors2' ✔ Creating 'R/' ✓ Writing 'DESCRIPTION' Package: foofactors2 Title: What the Package Does (One Line, Title Case) Version: 0.0.0.9000 Authors@R (parsed): * Jennifer Bryan <jenny@rstudio.com> [aut, cre] Description: What the package does (one paragraph). License: MIT + file LICENSE Encoding: UTF-8 LazyData: true ✓ Writing 'NAMESPACE' ✔ Writing 'foofactors2.Rproj' ✓ Adding '.Rproj.user' to '.gitignore' ✓ Adding '^foofactors2\\.Rproj\$', '^\\.Rproj\\.user\$' to '.Rbuildignore' ✔ Opening '/Users/jenny/tmp/foofactors2/' in new RStudio session ✓ Setting active project to '<no active project>'



What follows is adapted from

The Whole Game

chapter in the revised version of R Packages.

https://r-pkgs.org/whole-game.html

A proper package for the care and feeding of factors:

forcats

https://forcats.tidyverse.org

foofactors::fbind()

Factors can be vexing

```
(a <- factor(c("character", "hits", "your", "eyeballs")))</pre>
#> [1] character hits your eyeballs
#> Levels: character eyeballs hits your
(b <- factor(c("but", "integer", "where it", "counts")))</pre>
#> [1] but integer where it counts
#> Levels: but counts integer where it
c(a, b)
#> [1] 1 3 4 2 1 3 4 2
```

Factors can be vexing

```
factor(c(as.character(a), as.character(b)))
#> [1] character hits your eyeballs but integer where it
#> [8] counts
#> Levels: but character counts eyeballs hits integer where it your
```

Let's turn this into our first function: fbind()

Where do we define functions?

```
Beautiful pairing:
                         use_r() & use_test()
# There's a usethis helper for that too!
usethis::use_r("file-name")
# Organise files so that related code
# lives together. If you can give a file
# a concise and informative name, it's
# probably about right
```

Your turn

Use usethis::use_r("fbind") to create a new file In it, define a function named "fbind" that combines its inputs (presumably factors) like so:

- coerce each input to character
- combine inputs
- make output a factor

factor(c(as.character(a), as.character(b)))

Now what?

source("R/fbind.R")

Use IDE tricks to send definition of fbind() to the R Console

Now what?

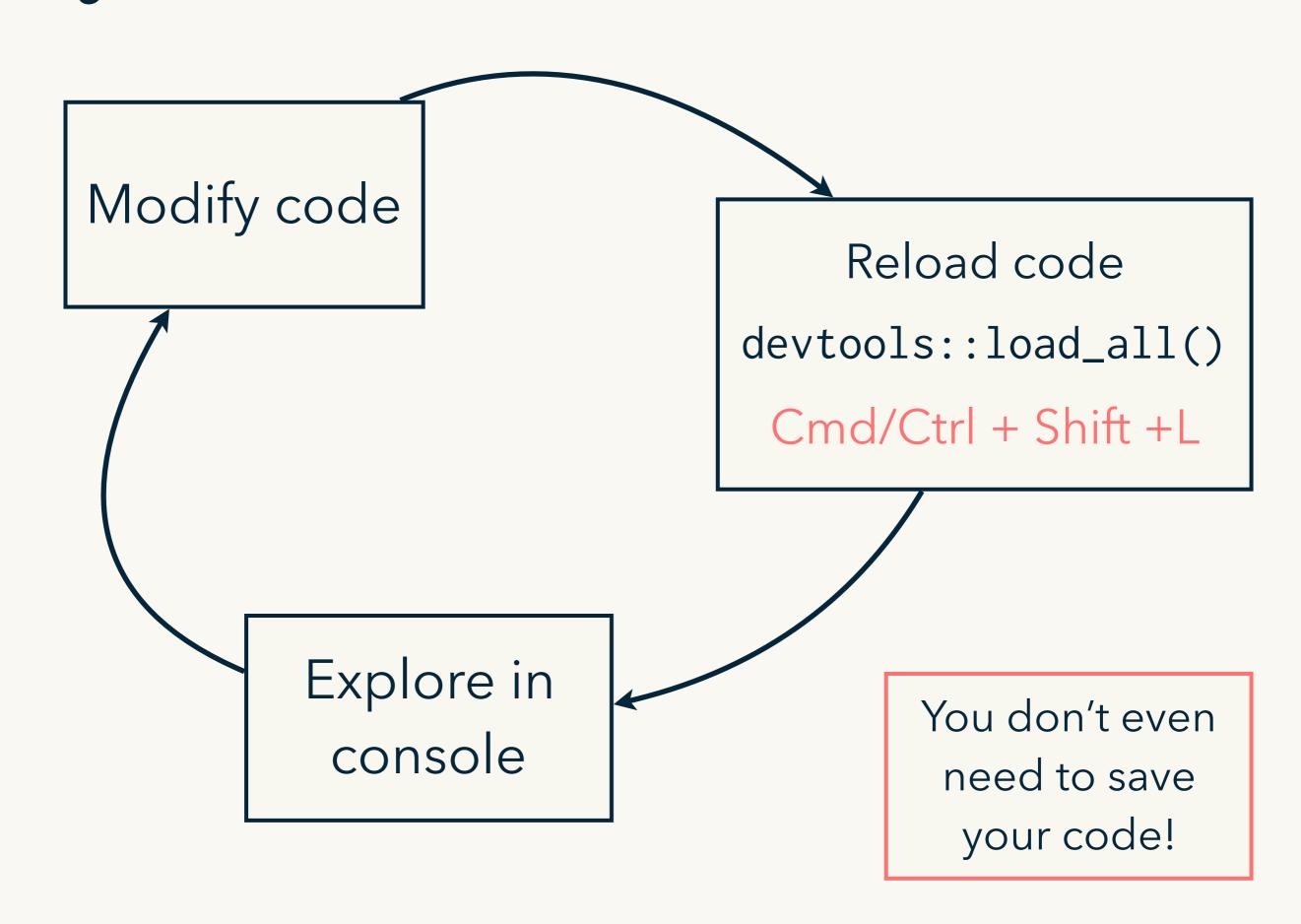
source("R/fbind.R")

Use IDE tricks to send definition of fbind() to the R Console

devtools::load_all()

devtools::load_all()

Why do we love devtools? Workflow!



Important metadata files exist in all versions

In binary versions, documentation is compiled into multiple versions. A parsed version of DESCRIPTION is cached for performance.

In binary versions, R/ no longer contains .R files, but instead contains binary .Rdata files

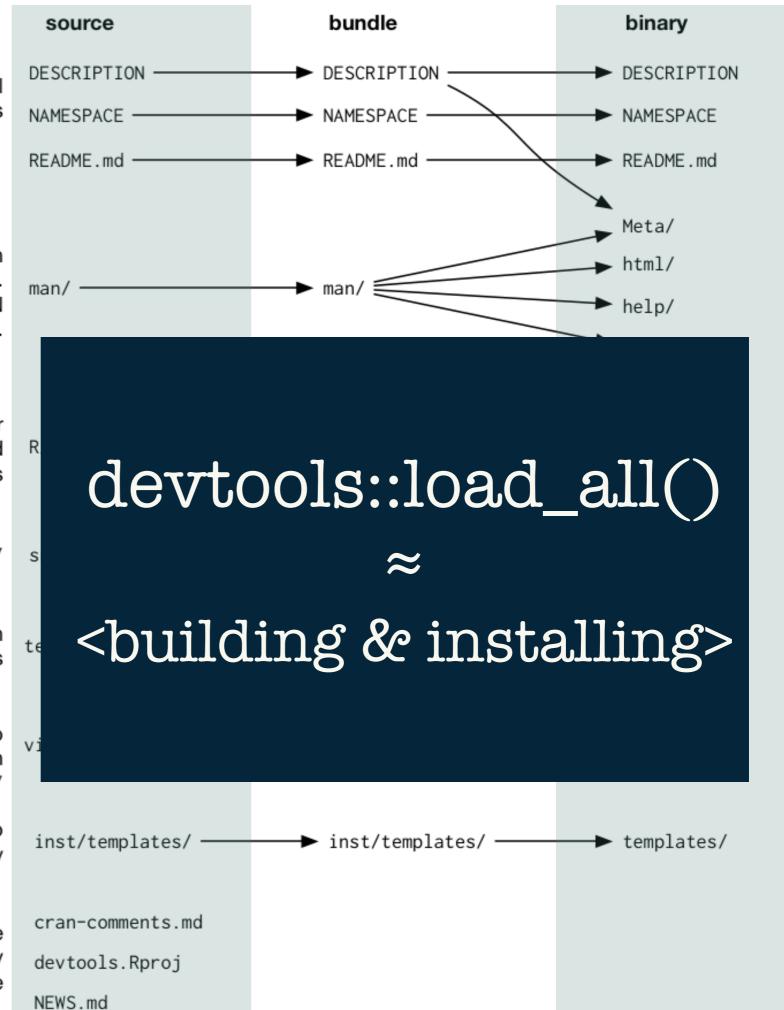
Compilation results are saved in libs/

By default, tests are dropped in binary packages

Source vignettes are build into html or pdf in inst/doc, then installed into doc/

The contents of inst/ are moved into the top-level directory

Files used only for development are listed in .Rbuildignore, and only exist in source package



R/RStudio setup

Workflow setup: your .Rprofile

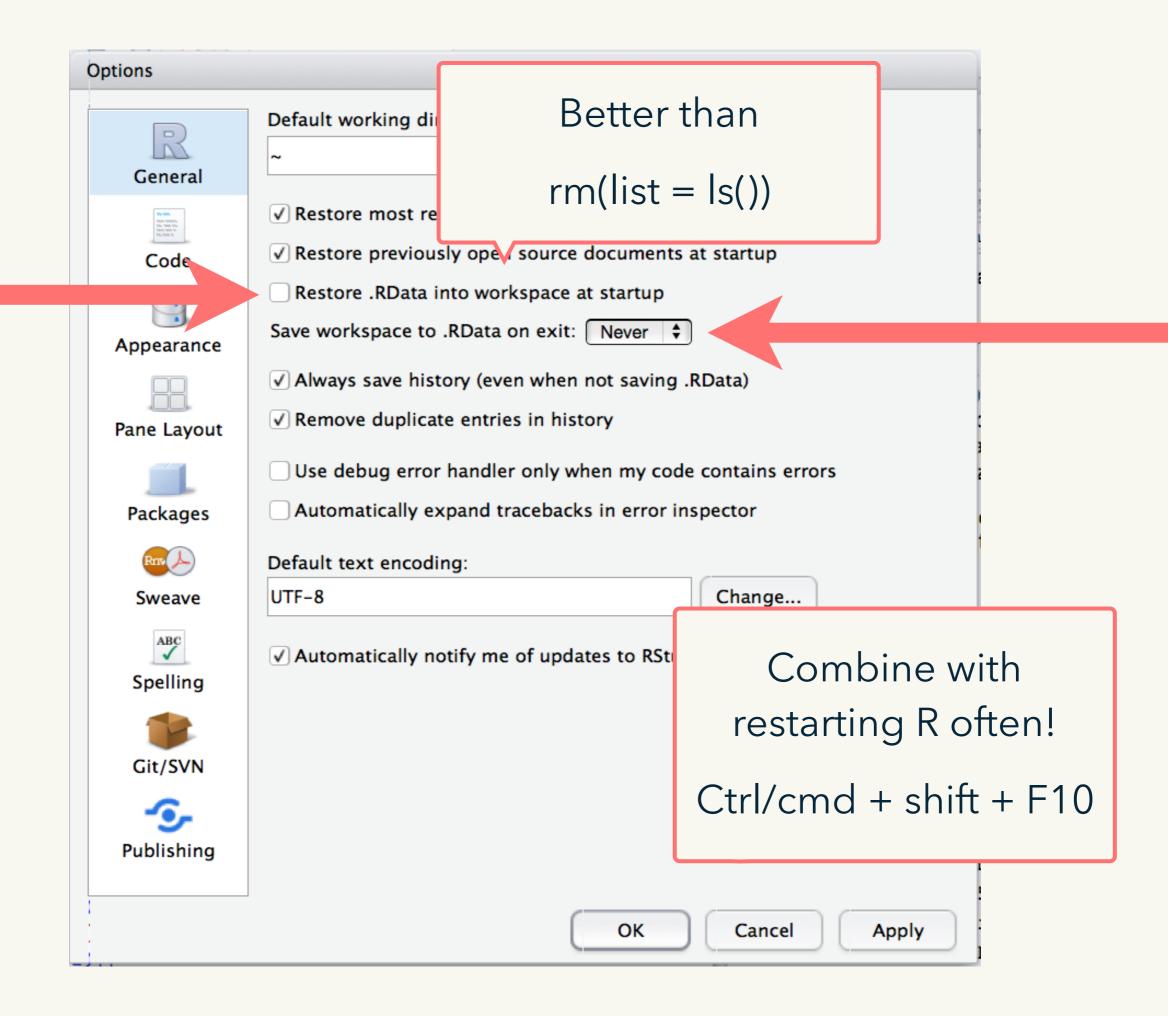
```
# Setup code that is run at R startup:
# usethis::edit_r_profile()
# Helper to add devtools specifically:
# usethis::use_devtools()
                                devtools makes
                              usethis available too!
if (interactive()) {
  suppressMessages(require(devtools))
```

Never include analysis packages here

```
if (interactive()) {
   suppressMessages(require(ggplot2))
   suppressMessages(require(dplyr))
}
```

While you're in there, also add:

```
# Helper to add devtools specifically:
# usethis::use_partial_warnings()
options(
  warnPartialMatchArgs = TRUE,
  warnPartialMatchDollar = TRUE,
  warnPartialMatchAttr = TRUE
```



devtools::check()



check() ≈ R CMD check

Checks package for technical validity

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Do from R (or RStudio Ctrl/cmd + shift + e)

check() early, check() often

Get it working, keep it working

Necessary (but not sufficient) for CRAN

Excellent way to run your tests (and more)

Your turn

Edit DESCRIPTION (optional)

- Make yourself the author
- Update Title and Description

use_mit_license("Your Name")

Fixes 1 of our 2 warnings.

check() again, if you wish

Nice to do, but skippable.

devtools::document()

roxygen2 turns comments into help

```
#' Bind two factors
# '
  Create a new factor from two existing factors, where the new
  factor's levels are the union of the levels of the input
  factors.
# '
                                            RStudio helper:
  @param a factor
  @param b factor
                                     Code > Insert roxygen skeleton
# '
  @return factor
  @export
  @examples
#' fbind(factor(letters[1:3]), factor(letters[26:24]))
fbind <- function(a, b) {
  factor(c(as.character(a), as.character(b)))
```

Your turn

Add docs for fbind() as a roxygen comment

- RStudio helper: Code > Insert roxygen skeleton
- Lines MUST start with #'

document() < Makes an .Rd file from the comment

check() again and ... rejoice!

devtools::check()



devtools::install()



install() ≈ R CMD install

- Makes an installed pkg from your source pkg
- Do from R (or RStudio Install and Restart)
- install() less often than you load_all() or check()
- Marks transition from developing your package to using your package

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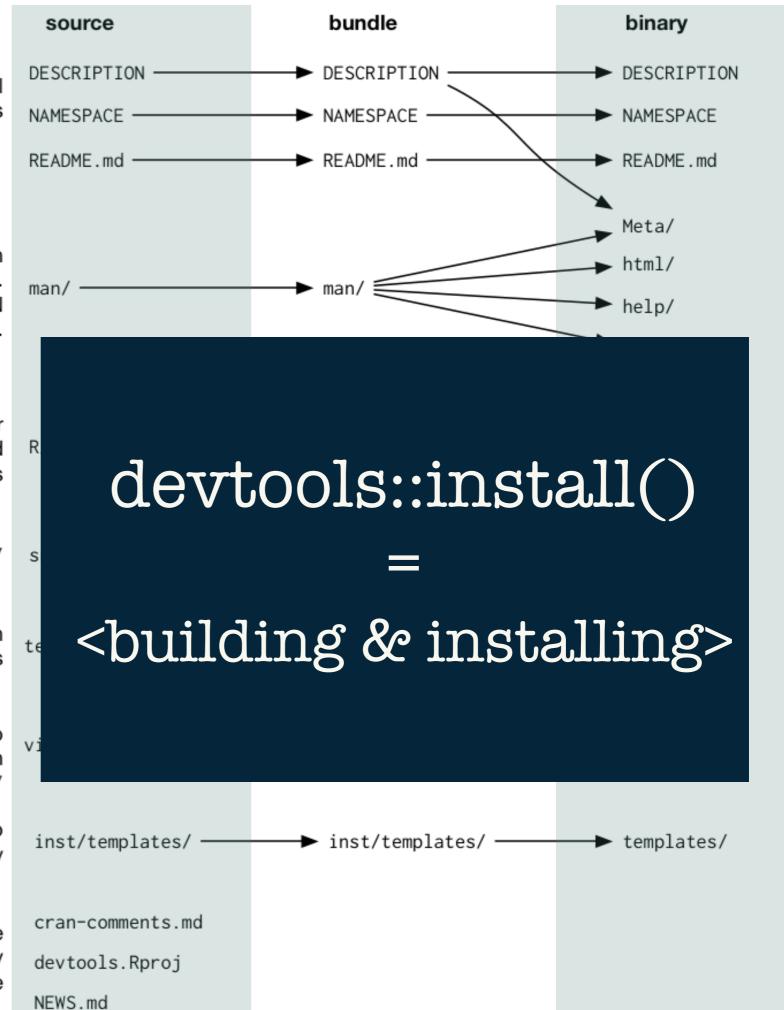
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Your turn

Install your foofactors package

- Call install() in R
- RStudio Build & Restart
- Shell: R CMD build + R CMD install

Restart R

Attach like a "regular" package with library()

Call fbind()

Revel in your success!

Other goodies if time allows

- use_package_doc()
- use_git() + use_github()

```
usethis::create_package()
devtools::load_all()
devtools::check()
devtools::document()
devtools::install()
```

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