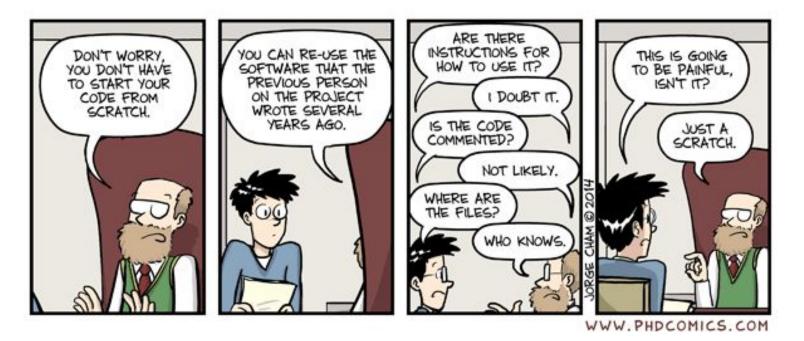
Introduction to R, RStudio, and RStudio Server

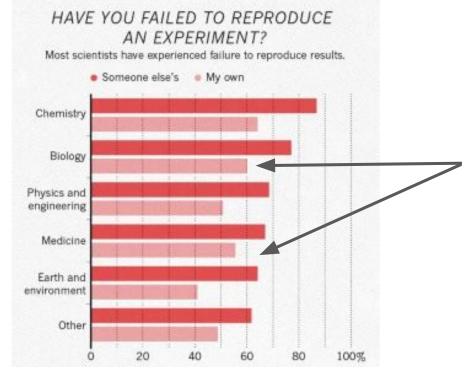
The CCDL

Who's been here before?



"Piled Higher and Deeper" by Jorge Cham www.phdcomics.com Used here with permission.

Reproducibility in 2016



55% and 60% of biologists and clinicians, respectively, could not reproduce <u>their own</u> results.

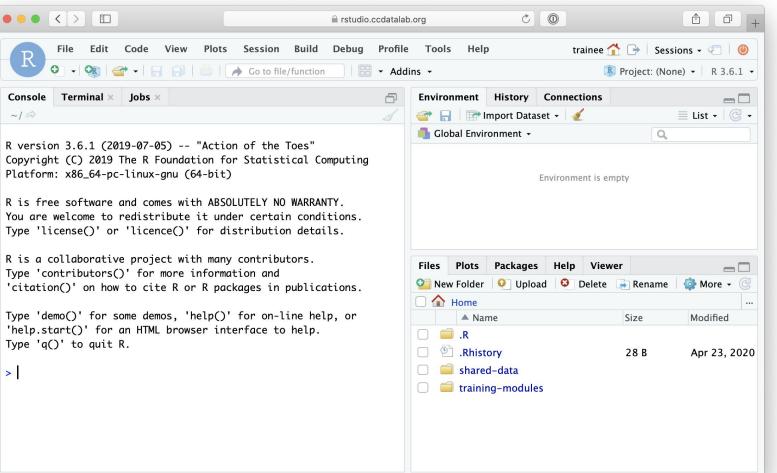
Baker, M. 1,500 scientists lift the lid on reproducibility. *Nature* 533, 452–454 (2016). https://doi.org/10.1038/533452a

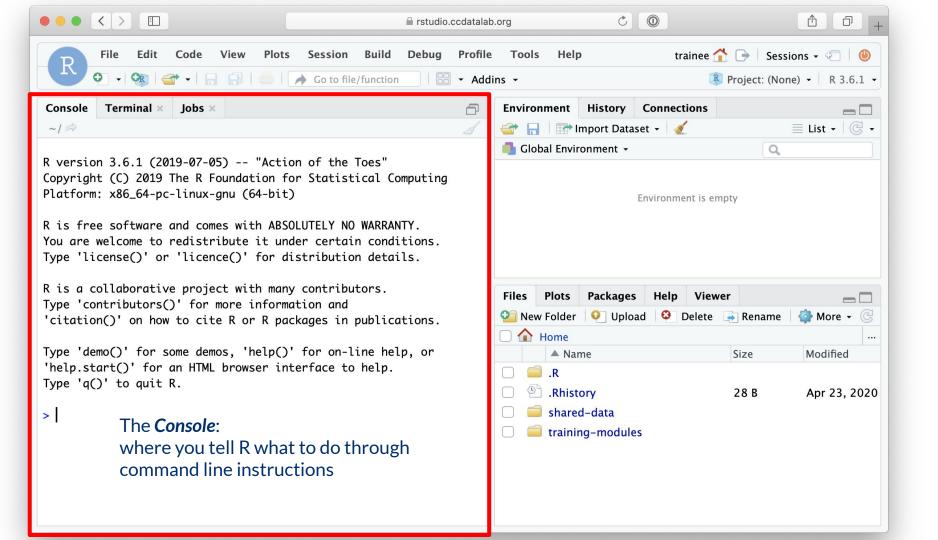
Command line vs GUI (graphics user interface)

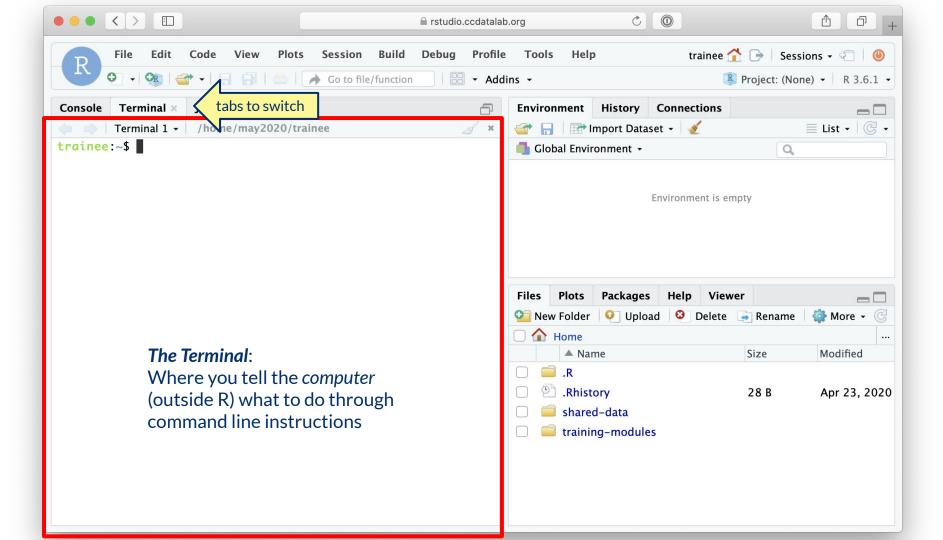
- An interface is how you interact with a program

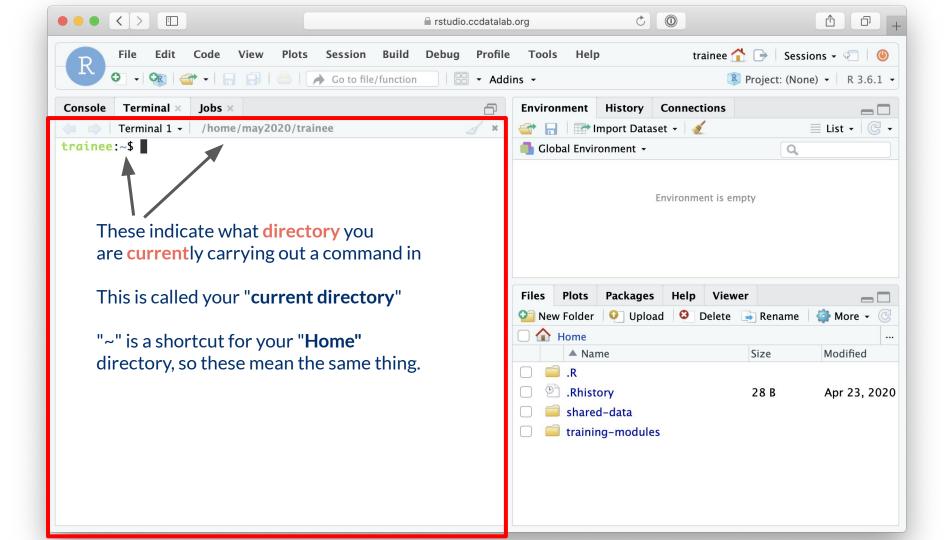
- GUI's have buttons you can *click* to do things, but...
- Command-line interfaces (CLI) have you *type* out things to do them

RStudio Server: A basic guide



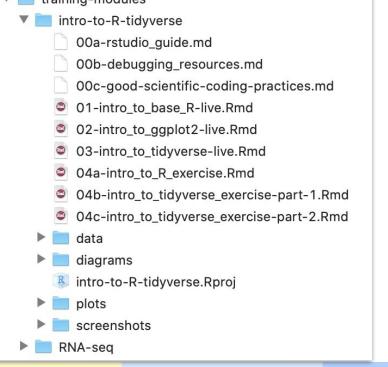


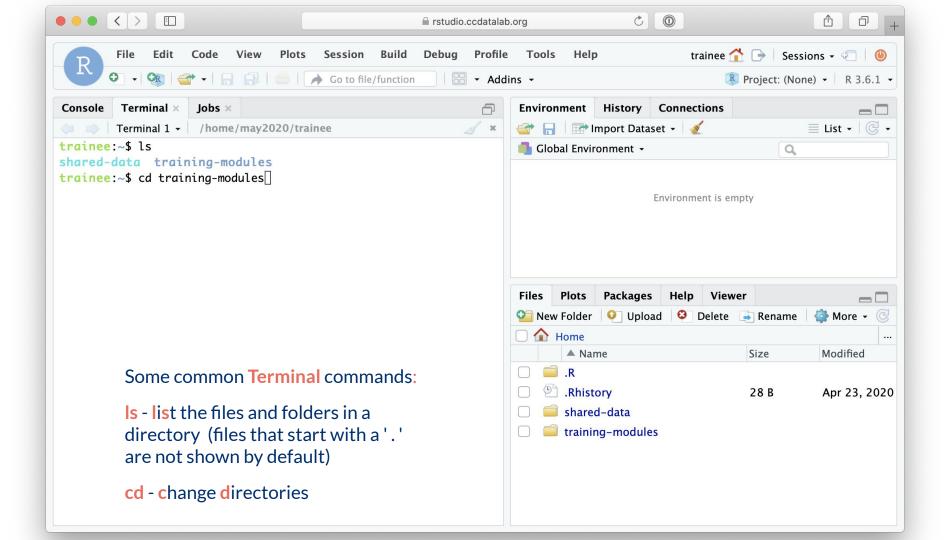


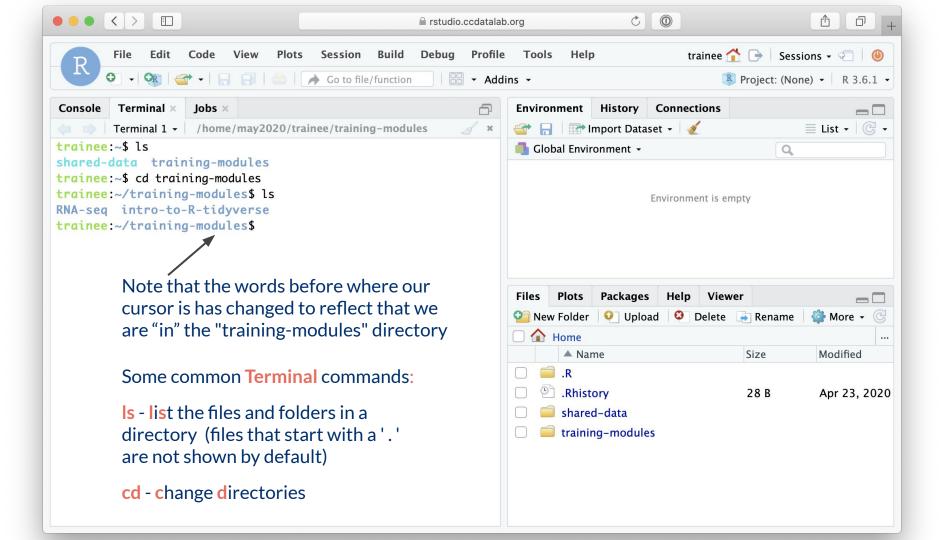


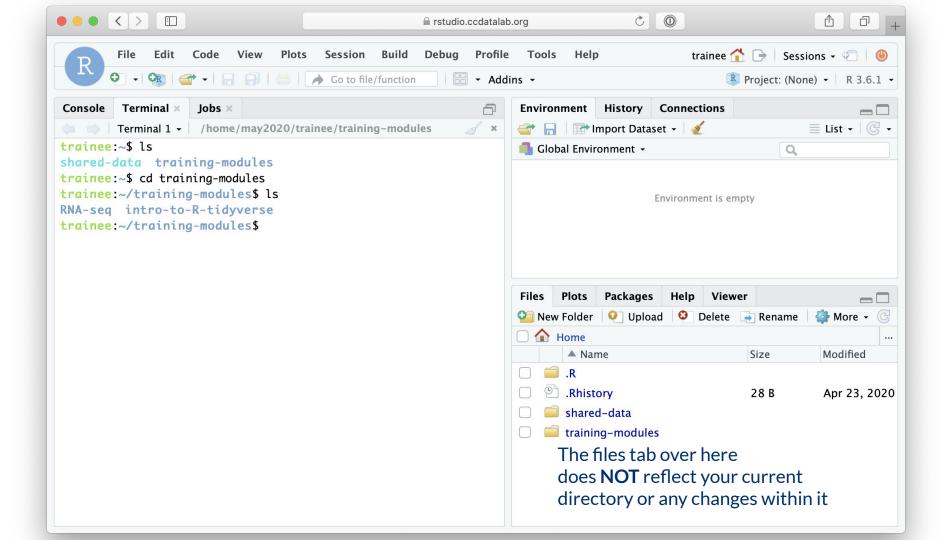
Directories = Folders

When we are working on the command line, we have to keep track of where the files we are using are being kept. Training-modules









File paths: Directions to a file or folder

Let's say we want access to "01-intro_to_base_R-live.Rmd"

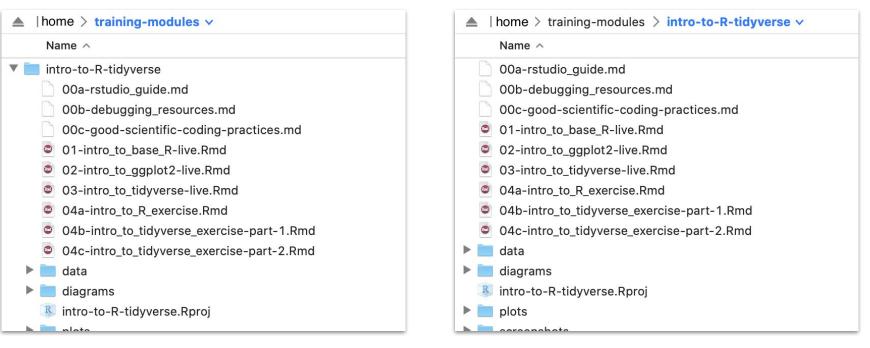
▲ home > training-modules ∨			
Name ^			
intro-to-R-tidyverse			
00a-rstudio_guide.md			
00b-debugging_resources.md			
00c-good-scientific-coding-practices.md			
01-intro_to_base_R-live.Rmd			
02-intro_to_ggplot2-live.Rmd			
03-intro_to_tidyverse-live.Rmd			
04a-intro_to_R_exercise.Rmd			
04b-intro_to_tidyverse_exercise-part-1.Rmd			
04c-intro_to_tidyverse_exercise-part-2.Rmd			
🕨 🚞 data			
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intro-to-R-tidyverse.Rproj			
plots			
screenshots			
▶ 📄 RNA-seq			

Current directory = "training-modules"

File path = "intro-to-R-tidyverse/01-intro_to_base_R-live.Rmd"

Relative file paths

Let's say we want access to "01-intro_to_base_R-live.Rmd"



Current directory = "training-modules"

Relative file path = "intro-to-R-tidyverse/01-intro_to_base_R-live.Rmd" Current directory = "training-modules/intro-to-R-tidyverse"

Relative file path = "01-intro_to_base_R-live.Rmd"

Introduction to R

The CCDL

R programming

Programming: making executable scripts for accomplishing a task (in this case, data analysis is our task)

Scripts allow others to see, step-by-step, what you did.

Why we use R:

- It's free and open-source
- People make cool packages that do stuff for us
- Many researchers in genomics use it (as well as Python)

One in five genetics papers contains errors thanks to Microsoft Excel

By Jessica Boddy | Aug. 29, 2016, 1:45 PM

What you type	What you see How Excel store		
MARCH1	1-MAR	42430	
SEPT2	2-SEP	42615	

https://www.sciencemag.org/news/2016/08/one-five-genetics-papers-contains-errors-thanks-microsoft-excel Ziemann et al. Genome Biology (2016) 17:177 DOI 10.1186/s13059-016-1044-7

R, RStudio, and RStudio Server

R is a statistical programming language.



RStudio is an IDE for working in R

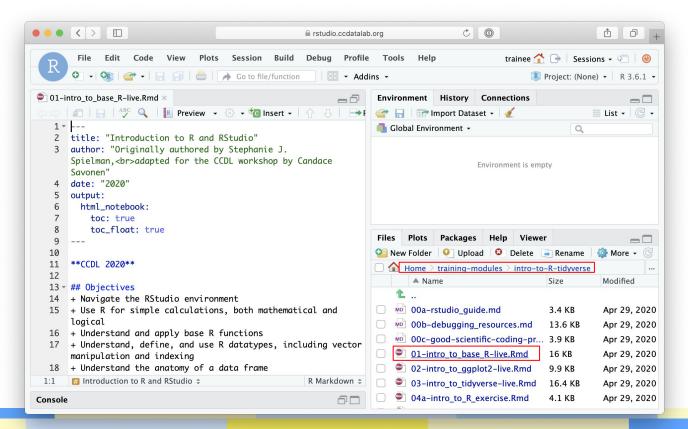
- IDE: Integrated Development Environment
- We write R code using the (free!) RStudio IDE



RStudio Server allows us to run the RStudio IDE from a browser

R Notebooks

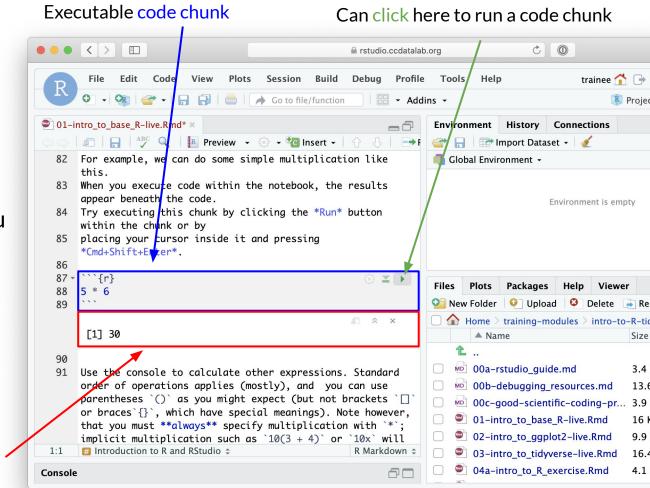
Use the "Files" tab to open: training-modules/intro-to-R-tidyverse/01-intro_to_base_R-live.Rmd



R Notebooks

 R Notebooks allow you to have files that show both your code and results

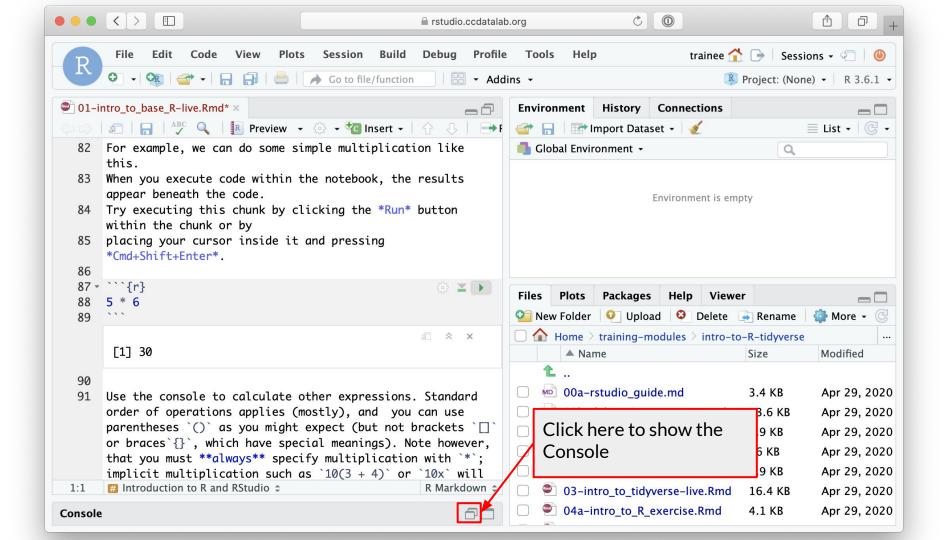
Output from above code chunk



R Notebooks

- Code that runs in R Notebooks uses <u>wherever the file is</u> <u>saved</u> as its current directory
- Warning! That may not be the directory shown in the files pane *or* the console!

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