# Programming Practices Winter Institute in Data Science

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2025-01-04



Style

https://style.tidyverse.org

fit\_models.R
clean\_data.R

fit\_models.R
clean\_data.R

(not stuff.R, trying.R)

00-read-data.R 01-clean-data.R

00-read-data.R 01-clean-data.R

(not work.R, laterwork.R)

## Good Variable Names

janitor::clean\_names()

### Good Internal Organization

- 1. Start .R with a comment preamble (context, data, author?, etc.)
- 2. Start with

```
library(dplyr)
library(readr)
```

3. Set off sections in .R with Cmd-Shft-R

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- meaningful
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- numbers
- separate words with \_
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not

```
T <- FALSE
c <- 10
mean <- function(x){
   sum(x)
}</pre>
```

1. Comma space as in English

```
df[, 5] c(1, 2, 3)
```

1. Comma space as in English

2. Parentheses without space

```
mean(x)
(not mean (x) nor mean(x))
```

1. Comma space as in English

2. Parentheses without space

3. Space around operators

4. No space around high-priority operators

1:10 x[6]

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- 6. Next pipes on new lines

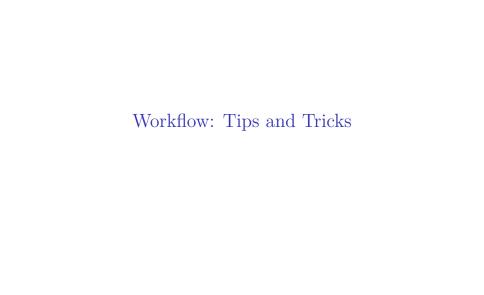
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- 5. Align delimiters { with }
- 6. Next pipes on new lines
- 7. Next args on new lines, when long, or helpful

## Usage

- 1. Use <- for assignment.
- 2. Use # for comments.
- 3. Comments with sentence case, but no period, unless two sentences.



# Keyboard Tips

Tab completion	speed, accuracy
Up arrow	previous command
Cmd-Return	$\operatorname{code} \rightsquigarrow \operatorname{Console}$
Cmd-Shift-0	restart R
Cmd-Shift-A	reformat highlit code
Alt-Shift-K	shortcut for shortcuts
Cmd-Shift-S	${ t source()} \ { t a} \ . { t R} \ { t file}$
Cmd-Shft-R	make section in .R

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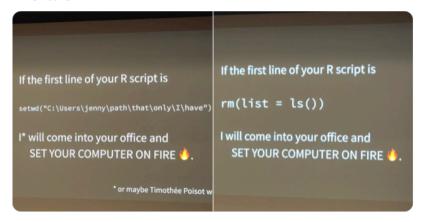
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#### Why?







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Cmd-Shft-0 regularly.

#### RStudio Diagnostics (turn them on)

```
165 - calc_ava <- function(x){
166
        s \leftarrow sum(x)
167
       n <- length(x)
168
        avg <- sum(x) / length(x)
169
170
        return(avg)
171
```

# R Projects

# What is real?

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The code is real.

# What is not real?

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Objects (currently) in the workspace

# How could this possibly matter?

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matter?

It does.

```
full_df <- read_csv("...")
t_out <- t.test(outcome ~ treatment, data = full_df)
# Process into tmp df ...
ggplot(df, aes(treatment, prop)) + geom bar()</pre>
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subset_df <- filter(full_data, <condition>)
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Then, redo first plot adding aesthetics, error bars, ...
What is real t\_out? df? gg <- ggplot() object?

# The source code is real.

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The objects are realizations of the source code. Source for EVERY user modified object is placed in a particular directory or directories, for later editing and retrieval.

- the ESS manual

There are notebooks.

▶ Joel Grus, author of *Data Science from Scratch* 

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- ► Talk at JupyterCon 2018: http://j.mp/2QHxjHB
- "hidden state" problem
  - "state": contents of memory locs at pt in time

# Ensure that Code Captures Reality

- ► Start fresh (Cmd-Shft-0)
- ➤ Turn off "Restore history/workspace" prefs
- Run the file (Cmd-Shift-S)
- Use relative paths from the working dir

# How to Behave (featuring Jenny Bryan)

- Adopt a "project-oriented workflow":

  https://www.tidyverse.org/articles/2017/12/workfl
  vs-script/

  (a.k.a., avoid arson)
  - About pkg here:
    https://github.com/jennybc/here\_here

#### The fine print

here::here() figures out the top-level of your current project using some sane heuristics. It looks at working directory, checks a criterion and, if not satisfied, moves up to parent directory and checks again. Lather, rinse, repeat.

Here are the criteria. The order doesn't really matter because all of them are checked for each directory before moving up to the parent directory:

- Is a file named .here present?
- Is this an RStudio Project? Literally, can I find a file named something like foo. Rproj?
- Is this an R package? Does it have a DESCRIPTION file?
- Is this a remake project? Does it have a file named remake.vml?
- Is this a projectile project? Does it have a file named .projectile?
- Is this a checkout from a version control system? Does it have a directory named .git or .svn? Currently, only Git
  and Subversion are supported.

library(here)

library(here)

library(here)

path <- here("data", "anes\_pilot\_2016.csv")
path</pre>

```
library(here)
```

```
path <- here("data", "anes_pilot_2016.csv")
path</pre>
```

[1] "/Users/ryanmoore/Documents/github/winter-inst/data/anes\_pilot\_2016.csv"

```
Use here — not setwd()
  library(here)
   path <- here("data", "anes pilot 2016.csv")</pre>
   path
   [1] "/Users/ryanmoore/Documents/github/winter-inst/data/anes_pilot_2016.csv"
   anes <- read csv(path)</pre>
   head(anes, 2)
   # A tibble: 2 \times 594
     version caseid weight weight spss follow turnout12 to
     <chr> <dbl> <dbl>
                            <dbl> <dbl> <dbl>
   1 ANES 2~ 1 0.951 0.542 1
   2 ANES 2~ 2 2.67 1.52
```

# i 585 more variables: meet <dbl>, givefut <dbl>, info sign <dbl>, give12mo <dbl>, compromise <dbl>, ftoba

ftwhite <dbl>, fthisp <dbl>, ftgay <dbl>, ftjeb <dl

#

In top-level directory, create a .Rproj file for each project

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Then, always start work by opening the .Rproj file.

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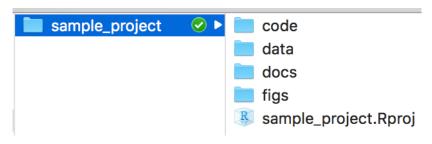
Then, always start work by opening the .Rproj file.

- ► Starts fresh R
- Sets wd to project directory
- Opens files you were working on
- Restores other settings (prefs, data, etc.)

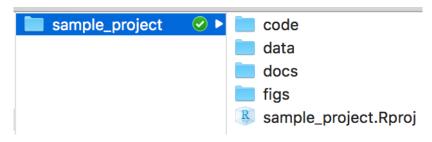
library(here) will look for your .Rproj file.

library(here) will look for your .Rproj file. That's (part) of how it knows the top-level dir.

Main project dir, subdirs:

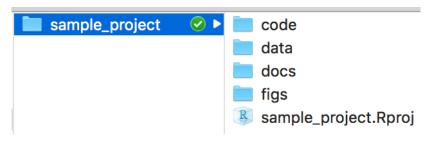


Main project dir, subdirs:



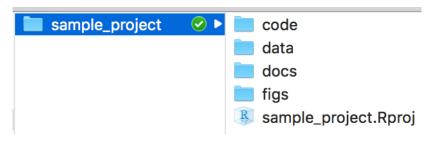
Click/open the .Rproj file in main project dir

▶ Main project dir, subdirs:



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- Dens files you had open

▶ Main project dir, subdirs:



- Click/open the .Rproj file in main project dir
- Dens files you had open
- ▶ Sets .Rproj dir as working dir

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- ▶ Best strategy: open .Rproj, use here

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```
df_1 <- read_csv(here("data", "first_data.csv"))
df_2 <- read_csv(here("data", "second_data.csv"))
ggplot(df, aes(x, y)) + geom_point()
ggsave(here("figs", "myplot.pdf"))</pre>
```

▶ In code files, paths are relative

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```
df_1 <- read_csv("data/first_data.csv")
df_2 <- read_csv("data/second_data.csv")
ggplot(df, aes(x, y)) + geom_point()
ggsave("figs/myplot.pdf")</pre>
```

- ▶ In code files, paths are relative
- Why here? If no .Rproj, open .R file from /code/ (same-level dir):

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- ▶ Why here? If no .Rproj, open .R file from /code/ (same-level dir):

```
df_1 <- read_csv("../data/first_data.csv")
df_2 <- read_csv("../data/second_data.csv")
ggplot(df, aes(x, y)) + geom_point()
ggsave("../figs/myplot.pdf")</pre>
```

#### Opening the .Rproj File

#### From http://j.mp/2QHeKDt:

When a project is opened within RStudio the following actions are taken:

- · A new R session (process) is started
- The .Rprofile file in the project's main directory (if any) is sourced by R
- The .RData file in the project's main directory is loaded (if project options indicate that it should be loaded).
- The .Rhistory file in the project's main directory is loaded into the RStudio History pane (and used for Console Up/Down arrow command history).
- The current working directory is set to the project directory.
- · Previously edited source documents are restored into editor tabs
- Other RStudio settings (e.g. active tabs, splitter positions, etc.) are restored to where they were the last time the project was closed.

#### sample\_project Directory

- Walk through sample\_project code in my research/sample\_project/
- Create .Rproj
- Make, save, load data
- Create figures

My desktop?

My desktop? My laptop?

My desktop? My laptop?

Your desktop?

My desktop? My laptop?

Your desktop? ...?

## The HEAD of origin, on

GitHub?

# The **HEAD** of **origin**, on GitHub?

Distributed version control. (Agree on origin.)

#### For next time:

- 1. Create a (free) GitHub account
- 2. Read §1 at https://happygitwithr.com/index.html
- 3. Complete §6 "Install Git"
- 4. Complete §7.0 "Introduce yourself" first paragraph
- 4. Complete §7.0 "Introduce yourself" first paragraph5. Follow §9 or §10 to set up either HTTPS PAT or SSH keys

(You can stop there.)