Programming Practices Winter Institute in Data Science

Ryan T. Moore

2024-01-04



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

Good Object Names

- meaningful
- lower case
- numbers
- separate words with _
- short enough
- not common R usages

Good Object Names

- meaningful
- lower case
- numbers
- separate words with _
- short enough
- not common R usages

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]

4. No space around high-priority operators

1:10 x[6]

5. Align delimiters { with }

4. No space around high-priority operators

```
1:10
x[6]
```

- 5. Align delimiters { with }
- 6. Next pipes on new lines

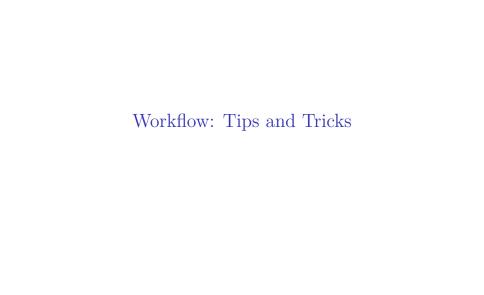
4. No space around high-priority operators

```
1:10
x[6]
```

- 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

Some metadata (what, who, ...)

- ➤ Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

- ➤ Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

- ➤ Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

Avoid

▶ install.packages("...")

- ➤ Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

Avoid

- install.packages("...")
- setwd("C:Users/me/my/unique/dir/structure/")

- Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

Avoid

- install.packages("...")
- setwd("C:Users/me/my/unique/dir/structure/")
- ▶ rm(list = ls())

- Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

Avoid

- install.packages("...")
- setwd("C:Users/me/my/unique/dir/structure/")
- ▶ rm(list = ls())

- ➤ Some metadata (what, who, ...)
- library(fivethirtyeight)
 library(tidyverse)

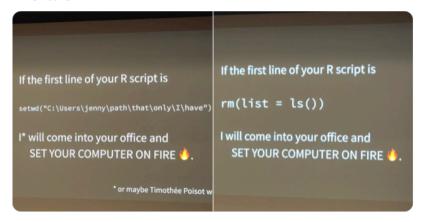
Avoid

- install.packages("...")
- setwd("C:Users/me/my/unique/dir/structure/")
- rm(list = ls())

Why?







Reproducibility: Starting Workflow

You should start fresh.

Reproducibility: Starting Workflow

You should start fresh.

Turn off "Restore my history/workspace" prefs.

Reproducibility: Starting Workflow

You should start fresh.

Turn off "Restore my history/workspace" prefs.

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?

What is real?

The code is real.

What is not real?

What is not real?

Objects (currently) in the workspace

How could this possibly matter?

How could this possibly

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>
```

```
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>
```

Then, time pressure ...a meeting ...

```
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>
```

Then, time pressure ...a meeting ...

```
subset_df <- filter(full_data, <condition>)
t_out <- t.test(outcome ~ treatment, data = subset_df)
# Process into tmp df ...
ggplot(df, aes(treatment, prop)) + geom_bar()</pre>
```

```
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>
```

Then, time pressure ...a meeting ...

```
subset_df <- filter(full_data, <condition>)
t_out <- t.test(outcome ~ treatment, data = subset_df)
# Process into tmp df ...
ggplot(df, aes(treatment, prop)) + geom_bar()</pre>
```

Then, redo first plot adding aesthetics, error bars, ...

```
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>
```

Then, time pressure ...a meeting ...

```
subset_df <- filter(full_data, <condition>)
t_out <- t.test(outcome ~ treatment, data = subset_df)
# Process into tmp df ...
ggplot(df, aes(treatment, prop)) + geom_bar()</pre>
```

Then, redo first plot adding aesthetics, error bars, ...
What is real t_out? df? gg <- ggplot() object?

The source code is real.

The source code is real.

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*

- ▶ Joel Grus, author of *Data Science from Scratch*
- ► Talk at JupyterCon 2018: http://j.mp/2QHxjHB

- ▶ Joel Grus, author of Data Science from Scratch
- ► Talk at JupyterCon 2018: http://j.mp/2QHxjHB
- hidden state" problem

- ▶ Joel Grus, author of Data Science from Scratch
- ► 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/rtm/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/rtm/Documents/github/winter-inst/data/anes_pilot_2016.csv"
   anes <- read csv(path)
   head(anes, 2)
   # A tibble: 2 \times 594
     version caseid weight weight spss follow turnout12 to
     <chr> <dbl> <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

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

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

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

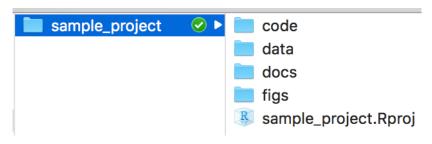
Then, always start work by opening the .Rproj file.

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

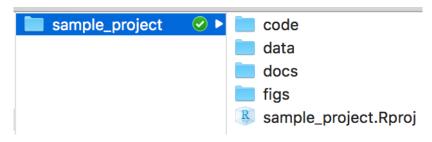
library(here) will look for your .Rproj file.

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

Main project dir, subdirs:

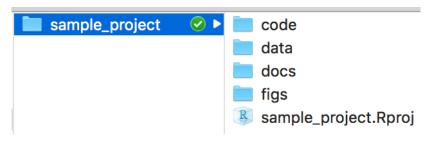


Main project dir, subdirs:



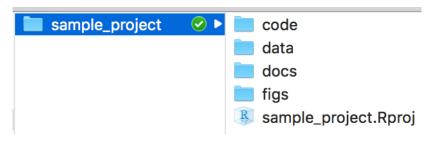
Click/open the .Rproj file in main project dir

▶ Main project dir, subdirs:



- Click/open the .Rproj file in main project dir
- 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

- ▶ In code files, paths are relative
- ▶ Best strategy: open .Rproj, use here

- ▶ In code files, paths are relative
- ▶ Best strategy: open .Rproj, use here

```
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

- ▶ In code files, paths are relative
- ▶ Why here? If open .Rproj or file at *same* level as /data/:

- ▶ In code files, paths are relative
- ▶ Why here? If open .Rproj or file at *same* level as /data/:

```
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):

- ▶ In code files, paths are relative
- ▶ 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 this afternoon:

- 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 5. Follow §9 or §10 to set up either HTTPS PAT or SSH kevs

(You can stop there.)