Programming Practices Winter Institute in Data Science

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Style

Workflow: Tips and Tricks

R Projects

Style

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https://style.tidyverse.org

fit_models.R
clean_data.R

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clean_data.R

(not stuff.R, trying.R)

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

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(not work.R, laterwork.R)

Good Internal Organization

- 1. Start 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

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- ▶ lower case
- numbers
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not

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

Legibility

- Comma space as in English (df[, 5], c(1, 2, 3))
- 2. Parentheses without space
 (mean(x), not mean (x) or mean(x))
- 3. Space around operators (x <- y, 4 / 11)
- 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.

Workflow: Tips and Tricks

Keyboard Tips

Tab completion	speed, accuracy
Up arrow	previous command
Cmd-Return	$\operatorname{code} \leadsto \operatorname{Console}$
Cmd-Shift-F10	restart R
Cmd-Shift-A	reformat highlit code
Alt-Shift-K	shortcut for shortcuts
Cmd-Shift-S	source() a .R file
Cmd-Shft-R	make section in .Rmd

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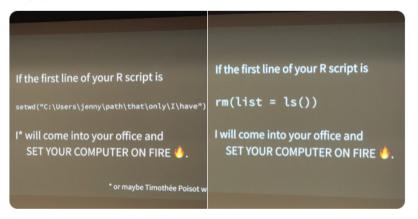
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Why not?







4:50 PM - 10 Dec 2017

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Cmd-Shft-F10 regularly.

RStudio Diagnostics (turn them on)

```
165 - calc_avg <- function(x){
166
        s \leftarrow sum(x)
167
        n <- length(x)
        avg <- sum(x) / length(x)
168
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?

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It does.

Recently, at The Lab ...

```
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>)
t out <- t.test(outcome ~ treatment, data = subset df)</pre>
# Process into tmp df ...
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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-Fn-F10)
- ► 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/ workflow-vs-script/ (a.k.a., avoid arson)

► About pkg here: https://github.com/jennybc/here_here

Use here (not setwd())

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.yml?
- 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.

Use here — not setwd() library(here)

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library(here)

here() starts at /Users/rtm/Documents/github/winter-inst

Use here — not setwd() library(here) ## here() starts at /Users/rtm/Documents/github/winter-inst path <- here("data", "anes_pilot_2016.csv") path</pre>

Use here — not setwd() library(here)

```
## here() starts at /Users/rtm/Documents/github/winter-inst
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)
   ## here() starts at /Users/rtm/Documents/github/winter-inst
   path <- here("data", "anes pilot 2016.csv")</pre>
   path
                                                              t1:
                                                              bl:
```

## [1] "/Users/rtm/Documents/github/winter-inst/de	ata/anes_pilot_2016.csv"
<pre>anes <- read_csv(path) head(anes)</pre>	
	spss follow turnout dbl> <dbl> <dbl> <dbl< td=""></dbl<></dbl></dbl>

2 ANES 2~ 2 2.67 1.52

3 ANES 2~ 3 1.43 0.815

0.150

55 / 80

4 ANES 2~ 4 0.914 0.521

0.264

5 ANES 2~

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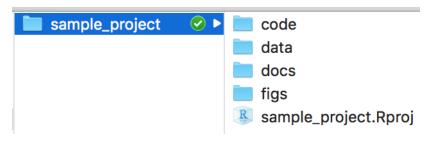
- ▶ Starts fresh R
- Sets wd to project directory
- Opens files you were working on
- ▶ Restores other settings (prefs, data, etc.)

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

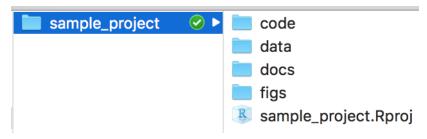
library(here) will look for your .Rproj file.

That's (part) of how it knows the top-level dir.

▶ Main project dir, subdirs:

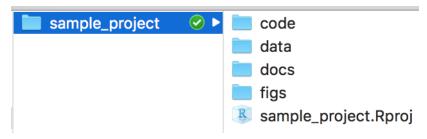


▶ Main project dir, subdirs:



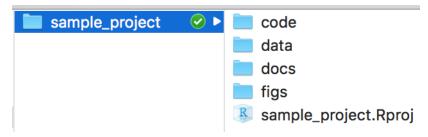
► Click/open the .Rproj file in main project dir

▶ Main project dir, subdirs:



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- ► Opens files you had open

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- ► Click/open the .Rproj file in main project dir
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- \blacktriangleright Sets . R
proj 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>
```

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- ▶ Why here? If open .Rproj file at *same* level as /data/:

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- ▶ Why here? If open .Rproj file at *same* level as /data/:

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df_1 <- read_csv("data/first_data.csv")
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ggplot(df, aes(x, y)) + geom_point()
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```

- ▶ 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")
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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

But which code version is real?

But which code version is real? My desktop?

But which code version is real? My desktop? My laptop?

But which code version is real?

My desktop? My laptop? Your desktop?

But which code version is real?

My desktop? My laptop? Your desktop? ...?

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

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Distributed version control. (Agree on origin.)