

STAT 215A Fall 2020

Week 2

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Announcements

- Lab 0 example code available at github.com/jpdunc23/stat-215a-fall-2020
- Make sure your **stat-215-a** repo is private
- Lab 1 will be released at the end of today's discussion section. **Due Thursday 17 at 11:59pm**

GitHub repos I have access to:

aagarwal1996
adelaidehsu
aidantmccloughlin
ajholmqu
akh9066
alexanderoa
alicecima
amishra791
anna-nguyen
aubluce
bouzaghrane
dayuapoji
denainjs
DohyeongKi
drewtnghuyen
ed-w
emilyrfl
Facta-Non-Verba
hanshengjiang
HaodongL

HeyuanLiu
Jordylek
kaylolittlejohn
ldliao
maocheng23
meiqingli
mgorenstein
mwerner28
nammaneerat
neurogomez
nikhil-ghosh-berkeley
OmerRonen
ravi-ivatari
rroggenkemper
SaptarshiC98
sfuller2
SuFangzhou
theo-s
utkarshapets
V-Chitra

wanghao98
WenxinZhang25
whryan
Yangch301
yiberkeley
yixiangLuo
zhenni-lin
zherenwang
ZitongYang

If you don't see your name,
please email / Slack me.

Today's outline

- `here()` and `across()`
- Some practice with Tidyverse
- Workflows
- Lab 1 Introduction

A quick poll on last week's discussion

Did you learn something new in last week's discussion?

sli.do

Event code #: 87666



continued...

dplyr: across ()

- `across()` supplants some of the “scoped verbs” that end with `_if()`, `_at()`, and `_all()`.
- Two primary arguments:
 - `.cols`: selects columns to operate on.
 - `.fns`: functions (can be more than one) to apply to the selected columns.

With `across()`:

```
> iris %>%  
+   mutate(across(.cols = contains("Sepal"), ~ 2 * .x)) %>%  
+   head()
```

With `mutate_at()`:

```
> iris %>%  
+   mutate_at(vars(contains("Sepal")), list(~ 2 * .)) %>%  
+   head()
```

dplyr: across () and summarize ()

- across () works particularly nicely with summarize ()

```
> iris %>%  
+   summarize(  
+     across(where(is.numeric), mean),  
+     across(where(is.factor), nlevels),  
+   )  
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species  
1      5.843333      3.057333         3.758      1.199333      3
```

- There are some examples where the across () syntax is not as nice:

```
# mutate + across  
mtcars %>% mutate_all(mean)  
# ->  
mtcars %>% mutate(across(everything(), mean))
```

- Learn more: <https://dplyr.tidyverse.org/articles/colwise.html>



here ()

here ()

- here is a very simple package that **increases reproducibility**
- When you run `library(here)` it checks the current working directory (i.e. whatever `getwd()` returns) for:
 - A file named `.here`
 - An RStudio project: `foo.Rproj`
 - An R package: `DESCRIPTION`
 - A git repo: `git`
 - Some others
- If it doesn't find any of those, it moves up to the parent directory and starts over.

here() example

PWD is week2

here() starts at the
git repo top-level
directory

```
> getwd()
[1] "/home/james/school/215a/stat-215a-fall-2020/week2"
> library(here)
here() starts at /home/james/school/215a/stat-215a-fall-2020
> here()
[1] "/home/james/school/215a/stat-215a-fall-2020"
> here("week2", "data", "mtcars.rds")
[1] "/home/james/school/215a/stat-215a-fall-2020/week2/data/mtcars.rds"
> mtcars2 <- readRDS(here("week2", "data", "mtcars.rds"))
> head(mtcars2)
```

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4

here() concatenates the
path

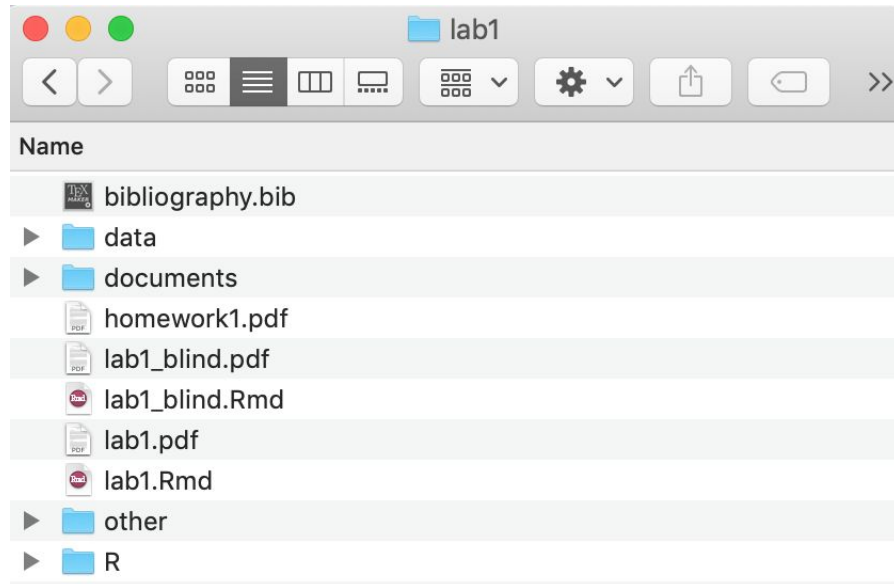
Let's get some hands-on practice

- About Gapminder: <http://www.gapminder.org/about-gapminder/>
- Resources for this tutorial:
 - ggplot: <http://swcarpentry.github.io/r-novice-gapminder/08-plot-ggplot2/>
 - dplyr: <http://swcarpentry.github.io/r-novice-gapminder/13-dplyr/>
- See **lab_gapminder.Rmd** in the week2 folder on my GitHub



Project file structure

- **data/**: store raw and processed data
- **documents/**: store relevant papers, instructions, meeting notes, etc.
- **R/**: store R code, utility functions, scripts
- **other/**: miscellaneous



Project File Structure

R/

- **load.R** – file containing function(s) for reading in the data

```
> loadData(path_to_data)
```

- **clean.R** – file containing function(x) for cleaning the loaded data

```
> cleanData(loaded_data)
```

data/

- ▶ Contains datasets
- ▶ Not uploaded to GitHub (can automate this using .gitignore)



Project File Structure

lab1.Rmd – your final report combining code (not printed in the output) and text/narrative

- Should be written like a paper; focus on communicating well

lab1.pdf – pdf output from lab1.Rmd

lab1_blind.Rmd – same as lab1.Rmd but without name

lab1_blind.pdf – pdf output from lab1_blind.Rmd

explore.Rmd (optional) – a separate .Rmd file that contains your exploratory code and figures

- A useful place for exploring the data and saving avenues of exploration that you don't necessarily want to include in your final report

bibliography.bib (optional) – a .bib file for easy citations within the lab reports

homework1.pdf – can be submitted electronically or in person at Friday lab section

Workflow: General Tips

Make code readable

- Be kind to both your peer reviewer and your future self

Keep your code modular – write functions

- Separate your functions from your analysis file (lab1.Rmd) and store them in R/
- In doing so, you create a bank of useful functions that you can load into any analysis script for your project (or future projects)

- To load in a single file:

```
> source("../R/filename.R")
```

- To load in all files in the R/ directory:

```
> library(R.utils)
```

```
> sourceDirectory("../R/", modifiedOnly = F, recursive = F)
```

- Group together related functions in the same .R script
(e.g. put all data cleaning functions in clean.R)

Workflow: General Tips

Documentation

- Write lots of comments in your code and ask yourself: why are you writing this particular piece of code?
- Document functions (think about the R help pages)
 - Always add comments section immediately below the function definition line
 - What does this function do?
 - Describe the inputs and outputs

```
CalculateSampleCovariance <- function(x, y, verbose = TRUE) {  
  # Computes the sample covariance between two vectors.  
  # Args:  
  #   x: One of two vectors whose sample covariance is to be calculated.  
  #   y: The other vector. x and y must have the same length, greater than one,  
  #       with no missing values.  
  #   verbose: If TRUE, prints sample covariance; if not, not. Default is TRUE.  
  # Returns:  
  #   The sample covariance between x and y.  
  ...  
}
```

Workflow: General Tips

Test your code

- Write tests to make sure your functions are doing the right thing
- Write these tests as you go

Don't Repeat Yourself (DRY)

- If you find yourself copying and pasting similar lines of code, write a reusable function instead

Establish consistencies – follow Google R Style Guide

Workflow: Code Style

Follow Google's R Style Guide when writing code

(See <https://google.github.io/styleguide/Rguide.xml> and part I Analyses of <https://style.tidyverse.org/syntax.html#object-names>)

Variable names

- All lowercase
- Separate words by “.” or “_” (be consistent with the one you choose)

Good: `avg.tmp`, `avg_tmp`

Bad: `AvgTmp`

Workflow: Code Style

Function names

- Camel-case
- Make function names verbs

Good: `CalculateAvgClicks`, `calculateAvgClicks`

Bad: `calculate_avg_clicks`, `calculate.avg.clicks`

Line Length: maximum length of 80 characters

- Go to: Preferences ☐ code ☐ display ☐ check show margin and set margin column = 80

Indentation: When indenting your code, use two spaces (rather than tabs)

Workflow: Code Style

Spacing

- ▶ Place spaces around all binary operators (=, +, -, <-, etc.)
- ▶ Always put a space after a comma, never before, just like in regular English

Good: `df.prior <- df[df$days.from.opt < 0, "campaign.id"]
x[, 1]`

Bad: `calculate_avg_clicks, calculate.avg.clicks
x[,1], x[, 1]`

Line Length: maximum length of 80 characters

- ▶ Go to: Preferences ☐ code ☐ display ☐ check show margin and set margin column = 80

Indentation: When indenting your code, use two spaces (rather than tabs)

Workflow: Code Style

Assignment

- ▶ Use `<-` instead of `=`

Curly Braces

- ▶ An opening curly brace should never go on its own line; a closing curly brace should always go on its own line
- ▶ Always begin the body of a block on a new line

Good: `if (x > 0) {
 print(x)
}`

Bad: `if (x > 0) print(x)`

Workflow: Code Style

Most importantly, BE CONSISTENT

Lab 1

Redwood Trees

Due: Thurs,
Sep 17 @ 11:59pm



Lab 1 Goals



Data cleaning



Exploratory Data Analysis and
Visualization

Lab 1 Redwood Introduction

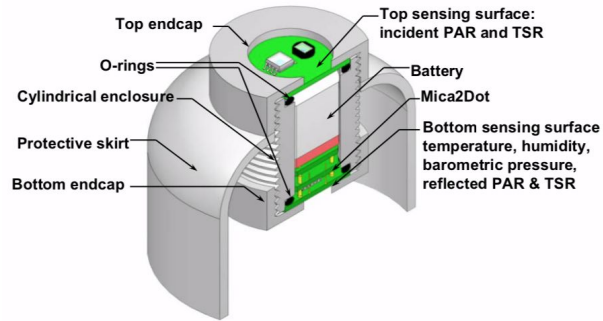


Figure 2: Sensor node and packaging

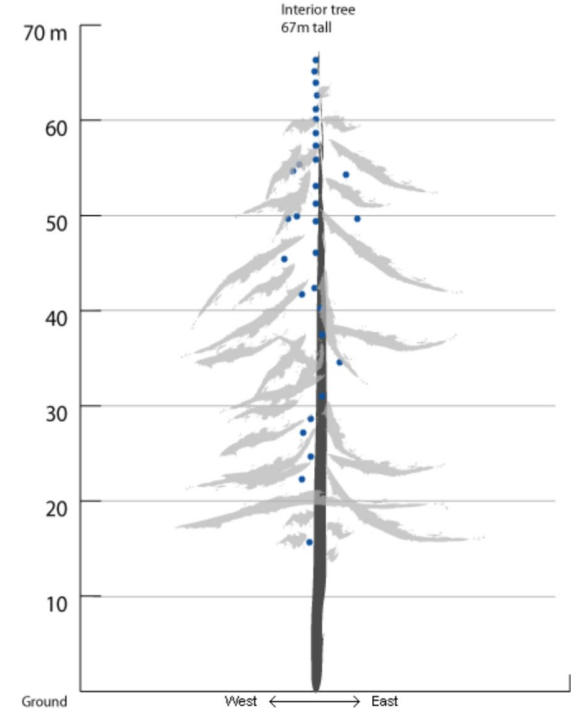


Figure 1: The placement of nodes within the tree

Lab 1 Introduction

- Read the paper carefully **sensys05-TollePolastreEtAl-redwoods.pdf** in the lab1 folder on Bcourses
- The lab1 folder on Bcourses will also contain a template to follow when putting together your lab as well as loading and cleaning functions that you may use/fill in
- The `exploration.Rmd` file was put together by Rebecca, a previous GSI, to get you started looking at the data, but you cannot use these plots as part of your lab report
- Do **not** push this `exploration.Rmd` file (or your own explore files) or the data folder to your stat-215-a repo
 - Can easily do this with `.gitignore` file

Collaboration Policy

- You are allowed to discuss **ideas** with others, but you must submit your own report
- Do not share code or copy/paste any part of the writeup
- If you do discuss ideas with others, be sure to acknowledge these students in your report

Lab 1 Rubric

Redwood Tree Lab (~60 points)

- Readability and grammar
- Readability of code (+ comments)
 - Follow Google's R Style Guide (a slight modification of the Tidyverse Style Guide)
- Reproducibility of report
 - I should be able to pull your `lab1/` folder from GitHub, manually add the `data/` folder, open `lab1.Rmd`, click knit, and get the same `.pdf` file as you
- Data cleaning (description and validity)
 - Describe *any* problems/inconsistencies you see with the data, how you cleaned the data, and why you cleaned the data in that way
- Three findings (creativity, interestingness, and quality of figure)
 - Fix titles, axis and legend titles, choose appropriate color schemes, adjust size of figure
- Graphical critique
- Figures that are not for the findings (relevance and quality)
- Overall quality and level of detail of report
 - Attempts to incorporate domain information (from the paper) and place your analysis in the domain context

Homework – Some Basic Statistics (8 points)

Start Early!!!!!!!
