Practice Solutions to Intro to R and Rstudio for EDA - Part 2

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OCTRI Biostatistics, Epidemiology, Research & Design (BERD) Workshop

2020/09/17

Practice 3

- 1. Continue adding code chunks to your Rmd (or, start a new one! But remember to load the libraries and data at the top.)
- 2. How many different years are in the data? (Hint: use tabyl() or n_distinct())
- 3. Count the number of penguins measured each year.
- 4. Calculate the median body mass by each species and sex subgroup. Use **summarize()** and **group_by()** to do this.
- 5. Create a 2x2 table of number of penguins measured in each year by each island.

Practice 3 Answers

2. How many different years are in the data? (Hint: use tabyl() or n_distinct())

Option 1:

```
penguins %>%
  summarize(n_distinct(year))
```

Option 2:

```
penguins %>% tabyl(year)
```

```
year n percent
2007 109 0.3187135
2008 114 0.3333333
2009 119 0.3479532
```

penguins %>% tabyl(year) %>% nrow

[1] 3

3. Count the number of penguins measured each year.

Option 1:

```
penguins %>% count(year)

# A tibble: 3 x 2
   year   n
   <dbl> <int>
1 2007 109
2 2008 114
3 2009 119
```

Option 2:

```
penguins %>% tabyl(year)
```

```
year n percent
2007 109 0.3187135
2008 114 0.3333333
2009 119 0.3479532
```

4. Calculate the median body mass by each species and sex subgroup. Use summarize() and group_by() to do this.

```
penguins %>%
 group_by(species, sex) %>%
  summarize(median(body_mass_g))
# A tibble: 8 x 3
# Groups: species [3]
 species sex
               `median(body mass g)`
 <chr> <chr>
                                  <dbl>
1 Adelie female
                                  3400
2 Adelie
        male
                                  4000
3 Adelie
           <NA>
                                  3475
4 Chinstrap female
                                  3550
5 Chinstrap male
                                  3950
          female
6 Gentoo
                                  4700
          male
7 Gentoo
                                  5500
8 Gentoo
           <NA>
                                  4688.
```

5. Create a 2x2 table of number of penguins measured in each year by each island.

```
penguins %>% tabyl(island, year)

island 2007 2008 2009

Biscoe 44 64 59

Dream 46 34 44

Torgersen 19 16 16
```

Practice 4 (old practice 2)

- 1. Create a new script and save it as Practice2.R
- 2. Create data frames for males and females separately.
- 3. Do males and females have similar BMIs? Weights? Compares means, standard deviations, range, and boxplots.
- 4. Plot BMI vs. weight for each gender separately. Do they have similar relationships?
- 5. Are males or females more likely to be bullied in the past 12 months? Calculate the percentage bullied for each gender.

Practice 4 Answers

#2 Create data frames for males and females separately.

```
boys <- mydata[mydata$sex == "Male", ]
dim(boys)

[1] 8 10

girls <- mydata[mydata$sex == "Female", ]
dim(girls)

[1] 12 10</pre>
```

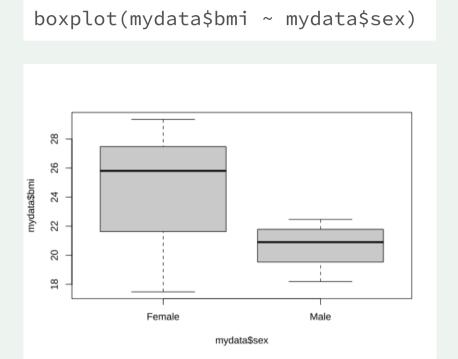
Check number of boys & girls:

```
Summary(mydata$sex)

Length Class Mode
20 character character
```

#3 Do males and females have similar BMIs? Weights? Compares means, standard deviations, range, and boxplots.

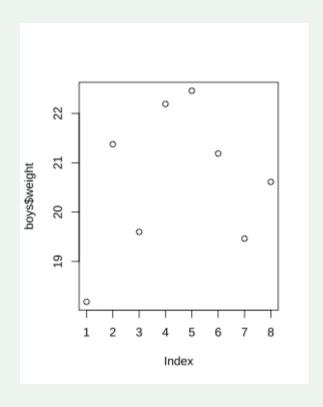
```
summary(boys$bmi); sd(boys$bmi)
  Min. 1st Qu. Median
                      Mean 3rd Qu.
                                      Max.
 18.18
         19.57
               20.90
                       20.63
                              21.58
                                       22.46
[1] 1.466896
summary(girls$bmi); sd(girls$bmi)
  Min. 1st Qu. Median
                      Mean 3rd Qu.
                                      Max.
                              27.47
 17.48
         21.95
               25.80
                        24.59
                                       29.35
[1] 3.70739
```

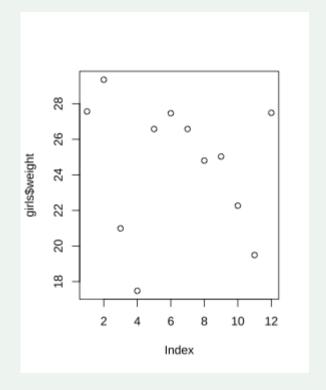


#4 Plot BMI vs. weight for each gender separately. Do they have similar relationships?

plot(boys\$bmi, boys\$weight)

plot(girls\$bmi, girls\$weight)





#5 Are males or females more likely to be bullied in the past 12 months? Calculate the percentage bullied for each gender.

```
bullied_boys <-</pre>
                                                         # Apply the same method for girls:
                                                         bullied_girls <-</pre>
  boys[boys$bullied_past_12mo == TRUE,]
nrow(bullied_boys)
                                                           girls[girls$bullied_past_12mo == TRUE,]
                                                         nrow(bullied_girls)
\lceil 1 \rceil 3
                                                         \lceil 1 \rceil 6
bullied_boys_prct <-</pre>
  nrow(bullied_boys) / nrow(boys) * 100
                                                         bullied_girls_prct <-</pre>
bullied_boys_prct
                                                           nrow(bullied_girls) / nrow(girls) * 100
                                                         bullied_girls_prct
[1] 37.5
                                                         \lceil 1 \rceil 50
# alternative
mean(boys$bullied_past_12mo, na.rm=TRUE)
                                                         # alternative. Answers don't match. Why???
                                                         mean(girls$bullied_past_12mo, na.rm=TRUE)
[1] 0.375
                                                         \lceil 1 \rceil 0.4
```

#5 cont'd

On the previous slide we saw that our two methods for calculating the percentage of girls that were bullied in the past 12 months did not match. What went wrong?

```
nrow(bullied_girls)

[1] 6

girls$bullied_past_12mo

[1] NA NA TRUE FALSE FALSE TRUE TRUE FALSE TRUE FALSE
[11] FALSE FALSE
```

To get the number of girls that were bullied we need to make sure the missing values (NA) are not included.

```
# values of bullied_past_12mo
girls$bullied_past_12mo
              NA TRUE FALSE FALSE TRUE TRUE FALSE TRUE FALSE
 \lceil 1 \rceil
[11] FALSE FALSE
# which are missing (logical)
is.na(girls$bullied_past_12mo)
     TRUE TRUE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
[11] FALSE FALSE
# which are NOT missing (logical)
!is.na(girls$bullied_past_12mo)
 [1] FALSE FALSE TRUE
                       TRUE
                              TRUE
                                    TRUE TRUE
                                                TRUE
     TRUE TRUE
\lceil 11 \rceil
```

#5 cont'd - working with NA's

#5 cont'd - fix girls' code

Exclude the missing values from the bullied_girls:

```
girls2 <- girls[!is.na(girls$bullied_past_12mo),]</pre>
nrow(girls2)
\lceil 1 \rceil 10
bullied_girls2 <- girls2[girls2$bullied_past_12mo == TRUE,]</pre>
nrow(bullied_girls2)
\lceil 1 \rceil 4
# from girls dataset, total number bullied
sum(girls$bullied_past_12mo, na.rm = TRUE)
\lceil 1 \rceil 4
```

#5 cont'd - Calculate percentage girls bullied

```
bullied_girls_prct2 <- nrow(bullied_girls2) / nrow(girls2) * 100
bullied_girls_prct2

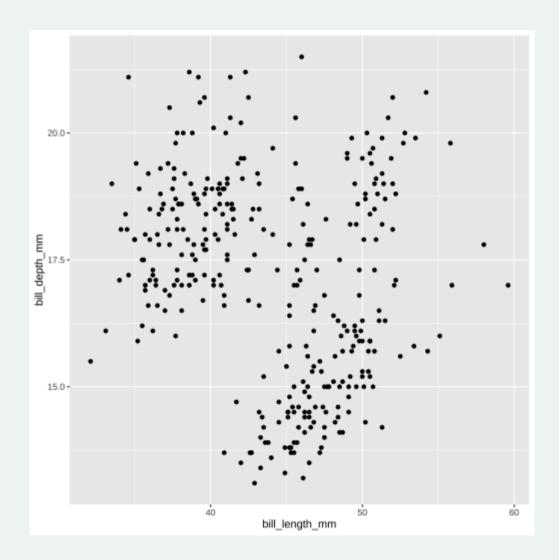
[1] 40

# Compare to alternative
mean(girls$bullied_past_12mo, na.rm=TRUE)</pre>
[1] 0.4
```

Practice 5

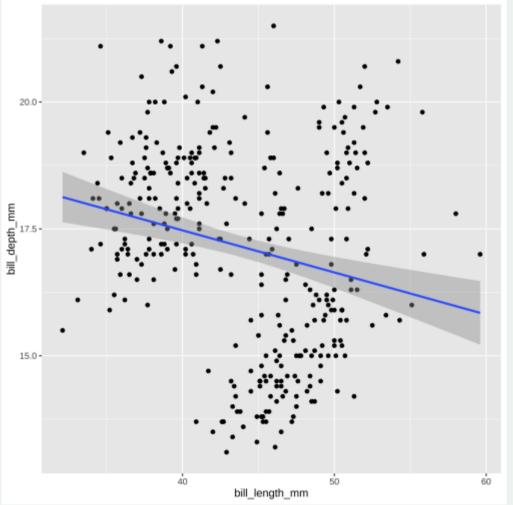
- 1. Continue adding code chunks to your Rmd (or, start a new one! But remember to load the libraries and data at the top.)
- 2. Make a scatter plot of bill depth vs bill length.
- 3. Add + geom_smooth(method="lm") to the plot. What is this saying about the association between bill depth and length?
- 4. Now add **color** = **species** to the aesthetic **aes()**. Keep **geom_smooth**. How do the associations look now?

2. Make a scatter plot of bill depth vs bill length.



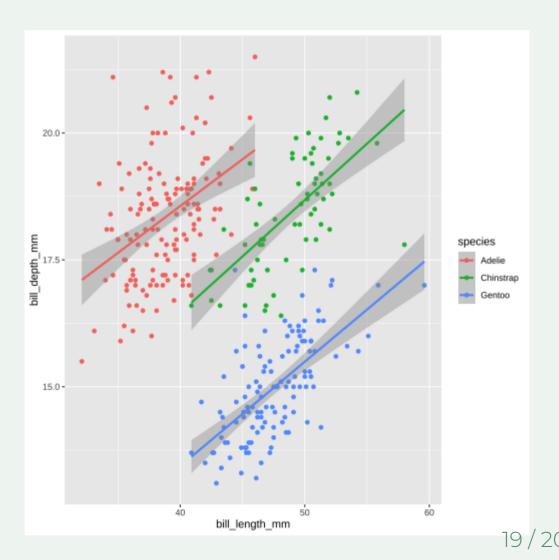
3. Add + geom_smooth(method="lm") to the plot. What is this saying about the association between bill depth and length?

It looks like as bill length increases, bill depth decreases. This is a negative association. But there is a wide variation and a lot of noise.



4. Now add color = species to the aesthetic aes(). Keep geom_smooth. How do the associations look now?

The association reverses, when we look inside species. As bill length increases, bill depth increases. This is an example of Simpson's paradox!



Here's a prettier version:

```
ggplot(data = penguins,
       aes(x = bill_length_mm,
           y = bill_depth_mm,
           color = species)) +
  geom_point()+
  geom_smooth(method="lm")+
  labs(
    title = "Flipper vs bill length",
    subtitle = "Palmer Station LTER",
    x = "Flipper length(mm)",
    y = "Bill length(mm)") +
  scale_color_viridis_d(
    name = "Species") +
  theme(legend.position = "bottom")+
  theme_bw()
```

