

Sasha's Mini Data Analysis Project

Sasha Tuttle October 14, 2021

```
{r setup, include=FALSE} knitr::opts_chunk$set(echo = FALSE)
```

Loading packages:

```
#install.packages("devtools")
#devtools::install_github("UBC-MDS/datateachr")
suppressPackageStartupMessages(library(rmarkdown))
suppressPackageStartupMessages(library(devtools))
suppressPackageStartupMessages(library(datateachr))
suppressPackageStartupMessages(library(tidyverse))
suppressPackageStartupMessages(library(ggplot2))
```

Loading the pilot study data

```
Pilot<-read.csv("C:/Users/stutt/OneDrive/Desktop/Greenhouse_study/Pilot_study.csv",
  header=T)
head(Pilot)
```

```
##   Plant_num Cluster_ID Variety Pollen_donor num_buds end_num_buds num_berries
## 1         1         1 Bluecrop   Bluecrop     68         68         42
## 2         1         2 Bluecrop     Duke     89         81         74
## 3         1         3 Bluecrop     Reka     62         52         52
## 4         1         4 Bluecrop   Bluecrop     76         76         59
## 5         1         5 Bluecrop     Duke     84         83         55
## 6         1         6 Bluecrop     Reka    102        102         81
##   fruit_set berry_weight stem_diam Average_TSS Average_pH Volume      TA
## 1 0.6176471      16.09      5.96      20.00      2.8557    6.00 25600.00
## 2 0.9135802      33.35      4.15      20.10      2.8197    5.99 25557.33
## 3 1.0000000      23.45      6.53      21.40      2.8850    6.20 26453.33
## 4 0.7763158      15.49      6.67      19.47      2.8260    5.89 25130.67
## 5 0.6626506      22.94      4.73      19.13      2.7690    6.44 27477.33
## 6 0.7941176      32.40      7.20      20.30      2.9130    5.81 24789.33
##   TA.true.
## 1 25.60000
## 2 25.55733
## 3 26.45333
## 4 25.13067
## 5 27.47733
## 6 24.78933
```

```
Stem<-rename(Pilot, stem_diam_cm = stem_diam)
```

Task 1: Investigating the datasets in the datateachr package

```
dim(Pilot)
```

```
## [1] 36 15
```

```
typeof(Pilot)
```

```
## [1] "list"
```

```
class(Pilot)
```

```
## [1] "data.frame"
```

The pilot study dataset has 36 rows and 15 columns. It is classified as a data frame.

```
##?cancer_sample  
print(cancer_sample)
```

```
## # A tibble: 569 x 32  
##       ID diagnosis radius_mean texture_mean perimeter_mean area_mean  
##   <dbl> <chr>      <dbl>      <dbl>      <dbl>      <dbl>  
## 1  842302 M         18.0       10.4       123.       1001  
## 2  842517 M         20.6       17.8       133.       1326  
## 3 84300903 M         19.7       21.2       130        1203  
## 4 84348301 M         11.4       20.4       77.6        386.  
## 5 84358402 M         20.3       14.3       135.       1297  
## 6  843786 M         12.4       15.7       82.6        477.  
## 7  844359 M         18.2       20.0       120.       1040  
## 8 84458202 M         13.7       20.8       90.2        578.  
## 9  844981 M          13        21.8       87.5        520.  
## 10 84501001 M         12.5       24.0       84.0        476.  
## # ... with 559 more rows, and 26 more variables: smoothness_mean <dbl>,  
## # compactness_mean <dbl>, concavity_mean <dbl>, concave_points_mean <dbl>,  
## # symmetry_mean <dbl>, fractal_dimension_mean <dbl>, radius_se <dbl>,  
## # texture_se <dbl>, perimeter_se <dbl>, area_se <dbl>, smoothness_se <dbl>,  
## # compactness_se <dbl>, concavity_se <dbl>, concave_points_se <dbl>,  
## # symmetry_se <dbl>, fractal_dimension_se <dbl>, radius_worst <dbl>,  
## # texture_worst <dbl>, perimeter_worst <dbl>, area_worst <dbl>, ...
```

```
typeof(cancer_sample)
```

```
## [1] "list"
```

```
dim(cancer_sample)
```

```
## [1] 569 32
```

```
class(cancer_sample)
```

```
## [1] "spec_tbl_df" "tbl_df"      "tbl"         "data.frame"
```

The cancer_sample data set has 596 rows, 32 columns, and is classified as a “list”/“data.frame”.

```
##?parking_meters
print(parking_meters)
```

```
## # A tibble: 10,032 x 22
##   meter_head r_mf_9a_6p r_mf_6p_10 r_sa_9a_6p r_sa_6p_10 r_su_9a_6p r_su_6p_10
##   <chr>      <chr>      <chr>      <chr>      <chr>      <chr>      <chr>
## 1 Twin      $2.00      $4.00      $2.00      $4.00      $2.00      $4.00
## 2 Pay Station $1.00      $1.00      $1.00      $1.00      $1.00      $1.00
## 3 Twin      $1.00      $1.00      $1.00      $1.00      $1.00      $1.00
## 4 Single     $1.00      $1.00      $1.00      $1.00      $1.00      $1.00
## 5 Twin      $2.00      $1.00      $2.00      $1.00      $2.00      $1.00
## 6 Twin      $2.00      $1.00      $2.00      $1.00      $2.00      $1.00
## 7 Twin      $2.00      $3.00      $2.00      $3.00      $2.00      $3.00
## 8 Single     $2.00      $3.00      $2.00      $3.00      $2.00      $3.00
## 9 Twin      $4.00      $1.00      $4.00      $1.00      $4.00      $1.00
## 10 Twin      $2.00      $1.00      $2.00      $1.00      $2.00      $1.00
## # ... with 10,022 more rows, and 15 more variables: rate_misc <chr>,
## #   time_in_effect <chr>, t_mf_9a_6p <chr>, t_mf_6p_10 <chr>, t_sa_9a_6p <chr>,
## #   t_sa_6p_10 <chr>, t_su_9a_6p <chr>, t_su_6p_10 <chr>, time_misc <chr>,
## #   credit_card <chr>, pay_phone <chr>, longitude <dbl>, latitude <dbl>,
## #   geo_local_area <chr>, meter_id <chr>
```

```
typeof(parking_meters)
```

```
## [1] "list"
```

```
dim(parking_meters)
```

```
## [1] 10032    22
```

```
class(parking_meters)
```

```
## [1] "tbl_df"      "tbl"        "data.frame"
```

The parking_meter data set has 10,032 rows, 22 columns, and is classified as a “list”/“data.frame”.

```
##?steam_games
print(steam_games)
```

```
## # A tibble: 40,833 x 21
##   id url   types name desc_snippet recent_reviews all_reviews release_date
##   <dbl> <chr> <chr> <chr> <chr>      <chr>      <chr>      <chr>
## 1 1 https~ app   DOOM Now include~ Very Positive~ Very Posit~ May 12, 2016
## 2 2 https~ app   PLAY~ PLAYERUNKNO~ Mixed,(6,214)~ Mixed,(836~ Dec 21, 2017
## 3 3 https~ app   BATT~ Take comman~ Mixed,(166),-- Mostly Pos~ Apr 24, 2018
## 4 4 https~ app   DayZ The post-so~ Mixed,(932),-- Mixed,(167~ Dec 13, 2018
## 5 5 https~ app   EVE ~ EVE Online ~ Mixed,(287),-- Mostly Pos~ May 6, 2003
## 6 6 https~ bund~ Gran~ Grand Theft~ NaN          NaN          NaN
## 7 7 https~ app   Devi~ The ultimat~ Very Positive~ Very Posit~ Mar 7, 2019
## 8 8 https~ app   Huma~ Human: Fall~ Very Positive~ Very Posit~ Jul 22, 2016
```

```
## 9      9 https~ app    They~ They Are Bi~ Very Positive~ Very Posit~ Dec 12, 2017
## 10     10 https~ app   Warh~ In a world ~ <NA>           Mixed,(904~ May 31, 2019
## # ... with 40,823 more rows, and 13 more variables: developer <chr>,
## #   publisher <chr>, popular_tags <chr>, game_details <chr>, languages <chr>,
## #   achievements <dbl>, genre <chr>, game_description <chr>,
## #   mature_content <chr>, minimum_requirements <chr>,
## #   recommended_requirements <chr>, original_price <dbl>, discount_price <dbl>
```

```
typeof(steam_games)
```

```
## [1] "list"
```

```
dim(steam_games)
```

```
## [1] 40833    21
```

```
class(steam_games)
```

```
## [1] "spec_tbl_df" "tbl_df"      "tbl"        "data.frame"
```

The `steam_games` dataset has 40833 rows, 21 columns, and is classified as a “list”/“data.frame”.

Datasets ranked from first to last choice:

1. Pilot_study
2. parking_meters
3. steam_games
4. cancer_sample

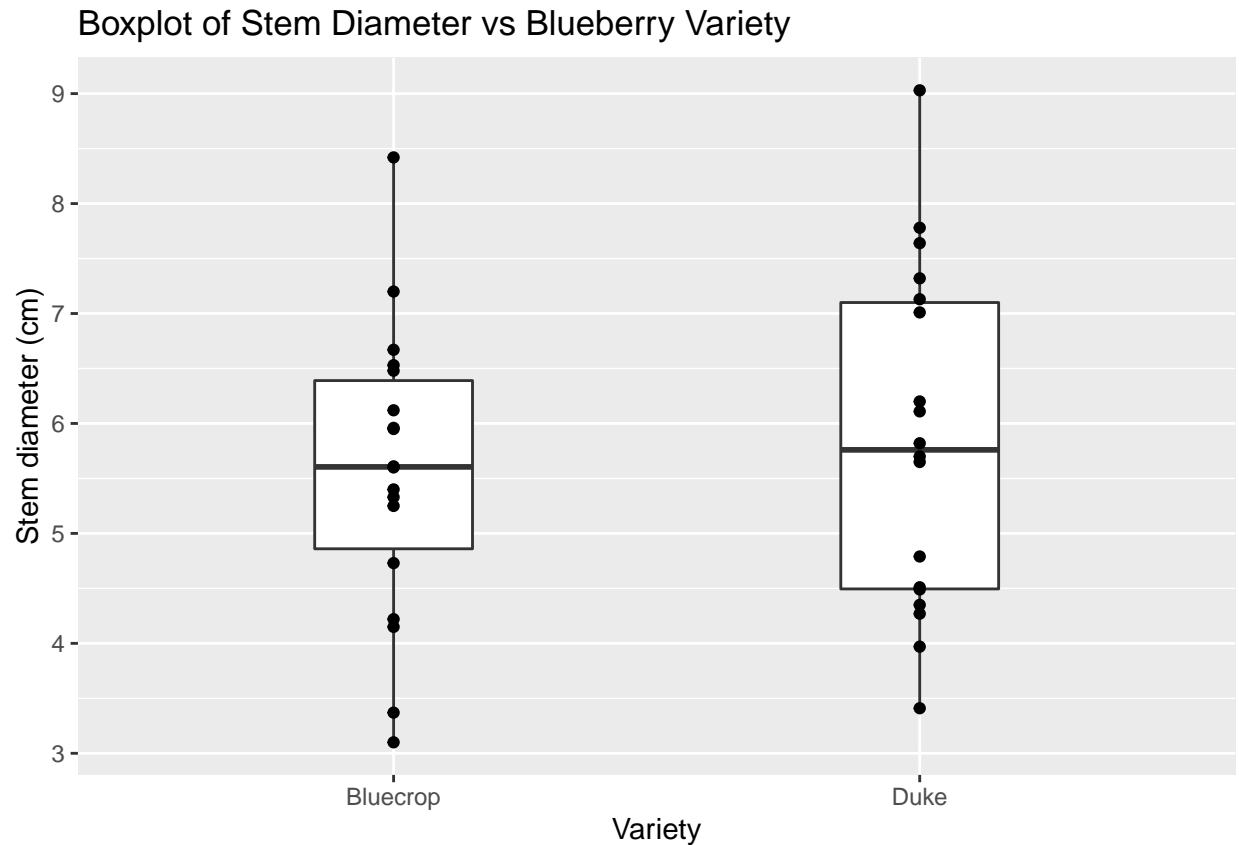
Of the four datasets investigated, my top two choices are the pilot study and parking meter data. The pilot study interests me because this dataset needs organizing and I am familiar with it. The parking meter data set is also ideal because it is smaller relative to the other datasets provided and it’s easier to understand (relative to the breast cancer dataset).

Task 2: Exploring data from the pilot study.

1. Plot the distribution of a numeric variable.

I plotted the stem diameter(cm) of the different blueberry varieties to see if one of the varieties had a greater stem diameter relative to the other. Thicker stems may have more leaves, allowing branches to produce a greater number of berries or heavier berries. So, if thicker branches were accidentally chosen for one variety compared to the other, we may see a greater yield and berry weight for said variety.

```
ggplot(Pilot, aes(Variety, stem_diam)) +
  geom_boxplot(width=0.3) + geom_point() + ylab("Stem diameter (cm)") +
  ggtitle("Boxplot of Stem Diameter vs Blueberry Variety ")
```



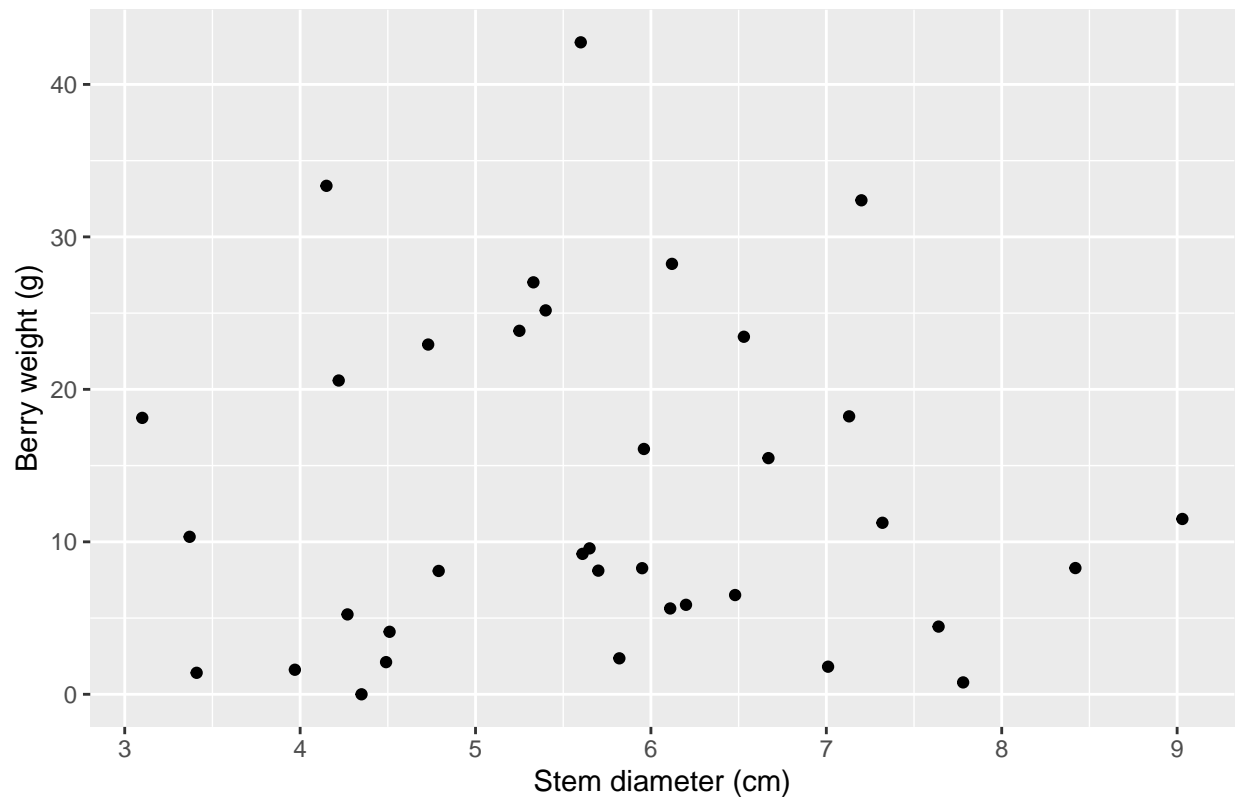
From the boxplot above, it doesn't seem like there is a significant difference between the stem diameter of the two varieties. However, I should also see if stem diameter differs significantly between different treatments within the same variety. This could act as a confounding variable and affect my results.

2. Explore the relationship between two variables in a plot.

I plotted the stem diameter of the variety against the berry weight to see if I could find a linear relationship. I hypothesized that clusters with thicker branches would have more leaves and energy to produce heavier berries.

```
ggplot(Pilot, aes(x=stem_diam, y=berry_weight)) +
  geom_point() + ylab("Berry weight (g)") + xlab("Stem diameter (cm)") +
  ggtitle("Scatterplot of Blueberry Weight vs Stem diameter")
```

Scatterplot of Blueberry Weight vs Stem diameter



The plot produced doesn't suggest a relationship is present between the two variables. I should analyze the data further with a regression to be certain.

3. Filter observations

I filtered my data into two different blueberry varieties (the pollen recipients in the experiment). Blueberry variety can significantly affect blueberry weight (among other parameters) and I'm not interested in this variable. So, I will assess the effect on each variety separately.

```
Blue<-filter(Pilot, Variety == "Bluecrop")
head(Blue)
```

```
##   Plant_num Cluster_ID Variety Pollen_donor num_buds end_num_buds num_berries
## 1         1         1 Bluecrop   Bluecrop     68         68         42
## 2         1         2 Bluecrop     Duke     89         81         74
## 3         1         3 Bluecrop     Reka     62         52         52
## 4         1         4 Bluecrop   Bluecrop     76         76         59
## 5         1         5 Bluecrop     Duke     84         83         55
## 6         1         6 Bluecrop     Reka    102        102         81
##   fruit_set berry_weight stem_diam Average_TSS Average_pH Volume      TA
## 1 0.6176471      16.09      5.96      20.00      2.8557    6.00 25600.00
## 2 0.9135802      33.35      4.15      20.10      2.8197    5.99 25557.33
## 3 1.0000000      23.45      6.53      21.40      2.8850    6.20 26453.33
## 4 0.7763158      15.49      6.67      19.47      2.8260    5.89 25130.67
## 5 0.6626506      22.94      4.73      19.13      2.7690    6.44 27477.33
```

```
## 6 0.7941176      32.40      7.20      20.30      2.9130      5.81 24789.33
##   TA.true.
## 1 25.60000
## 2 25.55733
## 3 26.45333
## 4 25.13067
## 5 27.47733
## 6 24.78933
```

```
Duke<-filter(Pilot, Variety == "Duke")
head(Duke)
```

```
##   Plant_num Cluster_ID Variety Pollen_donor num_buds end_num_buds num_berries
## 1      1      1      Duke      Duke      71      71      0
## 2      1      2      Duke    Bluecrop      75      67      7
## 3      1      3      Duke      Reka      73      65     13
## 4      1      4      Duke      Duke      68      66      3
## 5      1      5      Duke    Bluecrop      93      87     11
## 6      1      6      Duke      Reka      94      94      2
##   fruit_set berry_weight stem_diam Average_TSS Average_pH Volume      TA
## 1 0.0000000      0.00      4.35      NA      NA      NA      NA
## 2 0.10447761      2.11      4.49      20.10      NA      NA      NA
## 3 0.20000000      8.09      4.79      21.10      3.272      2.79 11904.00
## 4 0.04545454      1.41      3.41      16.33      NA      NA      NA
## 5 0.12643678      4.44      7.64      18.00      2.920      3.95 16853.33
## 6 0.02127660      0.78      7.78      17.57      NA      NA      NA
##   TA.true.
## 1      NA
## 2      NA
## 3 11.90400
## 4      NA
## 5 16.85333
## 6      NA
```

4. Create a tibble

I created a tibble to separate the Bluecrop variety from the Duke variety. Blueberry variety can have a significant effect on the variables I measured so I want to assess the effects of cross-pollination on each variety separately. I arranged pollen_donor column in descending order so I can more easily compare the self-crossed treatments against the out-crossed treatments. I also only selected the fruit set and berry weight columns so I can ignore the columns used in calculations and for sample IDs.

```
as_tibble(Pilot) %>%
  select(Variety, Pollen_donor, fruit_set, berry_weight,
         Average_TSS, Average_pH, TA.true.) %>%
  filter(Variety == "Bluecrop") %>%
  arrange(Pollen_donor, desc(berry_weight)) %>%
  rename(Berry_weight_g = berry_weight) %>%
  rename(TSS = Average_TSS) %>%
  rename(pH = Average_pH) %>%
  rename(Titratable_Acid = TA.true.)
```

```
## # A tibble: 18 x 7
```

##	Variety	Pollen_donor	fruit_set	Berry_weight_g	TSS	pH	Titrateable_Acid
##	<chr>	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
## 1	Bluecrop	Bluecrop	0.618	16.1	20	2.86	25.6
## 2	Bluecrop	Bluecrop	0.776	15.5	19.5	2.83	25.1
## 3	Bluecrop	Bluecrop	0.462	10.3	19.7	3.04	11.0
## 4	Bluecrop	Bluecrop	0.458	9.21	17.0	2.74	22.2
## 5	Bluecrop	Bluecrop	0.471	8.28	18.3	2.78	23.4
## 6	Bluecrop	Bluecrop	0.415	8.27	17.3	2.83	19.8
## 7	Bluecrop	Duke	0.914	33.4	20.1	2.82	25.6
## 8	Bluecrop	Duke	0.616	28.2	17.5	3.06	13.6
## 9	Bluecrop	Duke	0.702	23.8	19.2	2.83	22.3
## 10	Bluecrop	Duke	0.663	22.9	19.1	2.77	27.5
## 11	Bluecrop	Duke	0.548	18.1	21.2	3.13	15.8
## 12	Bluecrop	Duke	0.359	6.51	18.1	2.80	15.7
## 13	Bluecrop	Reka	0.62	42.8	17.8	3.20	8.96
## 14	Bluecrop	Reka	0.794	32.4	20.3	2.91	24.8
## 15	Bluecrop	Reka	0.511	27.0	19.6	2.99	17.1
## 16	Bluecrop	Reka	0.651	25.2	19.1	2.94	17.4
## 17	Bluecrop	Reka	1	23.4	21.4	2.88	26.5
## 18	Bluecrop	Reka	0.7	20.6	19.9	3.40	7.85

Task 3: Write your research questions

- Do blueberry flowers of the bluecrop variety experience a greater fruit set when they receive pollen from another blueberry variety?
- Do heavier berries have a higher TSS? a lower pH/TA?
- Does stem diameter have an effect on blueberry weight and/or TSS?
- Does the Bluecrop variety experience greater benefits from receiving pollen from another variety relative to the Duke variety?