DataVisualization

Kalen Clifton

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R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

To run each chunk, click on the green arrow button in each code block.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document.

Set up for Data Visualization

Uncomment and run chunk to install the package ggplot2. Recomment out this section before knitting the document

```
#install.packages("ggplot2")

Open ggplot2 library
```

```
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 3.6.2
```

Plotting Example

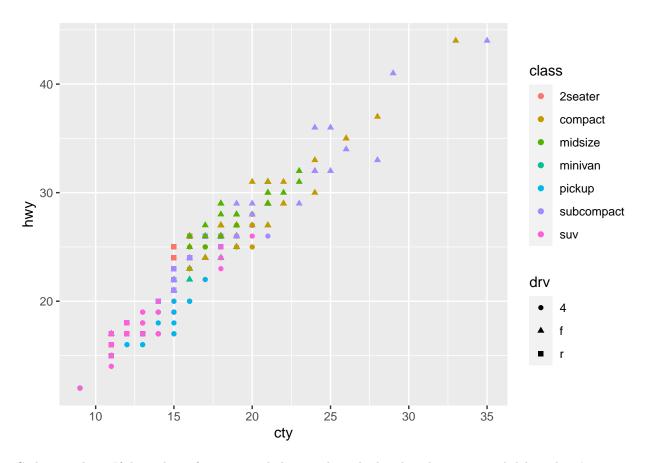
The dataset mpg has data on several car models collected the US EPA. The plot below displays some of that data. The variable on the x-axis is cty, a car's fuel efficiency in the city, in miles per gallon (mpg). The variable on the y-axis is hwy, a car's fuel efficiency on the highway in mpg.

Two other variables are included in the plot by aesthetic mapping. An aesthetic is a visual property of the objects in your plot. Aesthetics include things like the size, the shape, or the color of your points.

The class variable is added to the plot by mapping the levels of color to the values of class. (The class variable of the mpg dataset classifies cars into groups such as compact, midsize, and SUV.)

The drv variable (identifies type of drive including four wheel drive, front-wheel drive, and rear wheel drive) is added to the plot by mapping the levels of shape to the values of drv.

```
ggplot(data = mpg) +
  geom_point(mapping = aes(x = cty, y = hwy, color = class, shape = drv))
```



Code Template. If desired use for prompts below, replace the bracketed sections and delete the #'s

```
# ggplot(data = <DATA>) +
# <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Visualizing Data from Experiments on Cyanobacteria Growth

1. Load in growth data from characterization team (artificial data for example only)

```
growth <- read.csv('https://raw.githubusercontent.com/kpclifton/Baltimore_BioCrew/master/app/examples/D</pre>
```

2. View how the data is structured by printing the first 10 rows.

head(growth, 10)

```
##
      Days Iron
                    OD Trial
## 1
          1
              0x 0.05
                         1st
## 2
          2
              0x 0.05
                         1st
## 3
          3
              0x 0.05
                         1st
          4
## 4
              0x 0.05
                         1st
## 5
          5
              0x 0.05
                         1st
## 6
          6
              0x 0.05
                         1st
##
   7
          7
              0x 0.05
                         1st
## 8
          8
              0x 0.05
                         1st
## 9
          9
              0x 0.05
                         1st
## 10
         10
              0x 0.05
                         1st
```

3. View a summary of the data to understand basic statistics about the values in the columns (such as counts for categorical variables and means for continous variables).

summary(growth)

```
##
         Days
                        Iron
                                       OD
                                                   Trial
##
    Min.
           : 1.0
                    0x
                           :40
                                 Min.
                                         :0.0475
                                                   1st:50
##
    1st Qu.: 3.0
                    1/100x:40
                                 1st Qu.:0.0550
                                                   2nd:50
   Median: 5.5
                    1/10x :40
                                 Median :0.6150
                                                   3rd:50
##
##
    Mean
           : 5.5
                    10x
                           :40
                                 Mean
                                         :1.0926
                                                   4th:50
    3rd Qu.: 8.0
                                 3rd Qu.:2.0000
##
                           :40
##
    Max.
           :10.0
                                 Max.
                                         :3.3000
```

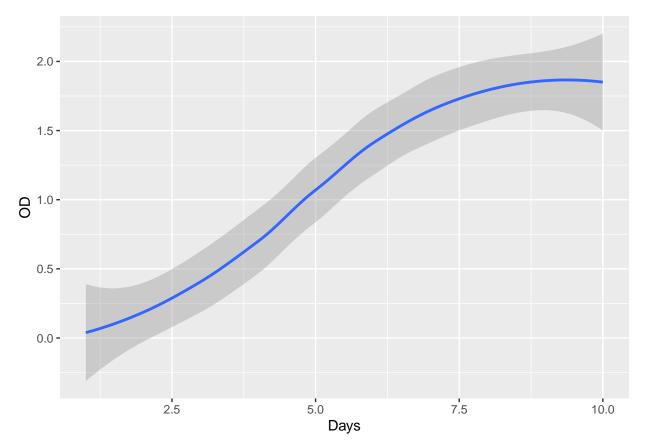
Optional: The View function opens the dataframe in another tab where you can scroll through the whole dataset

#View(growth)

4. Plot Days vs OD as smooth line using geom_smooth

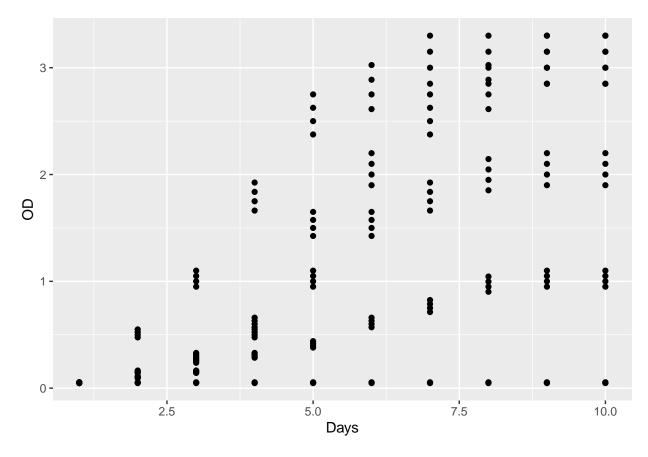
```
ggplot(data = growth) +
geom_smooth(mapping = aes(x = Days, y = OD))
```

```
## geom_smooth() using method = 'loess' and formula 'y ~ x'
```



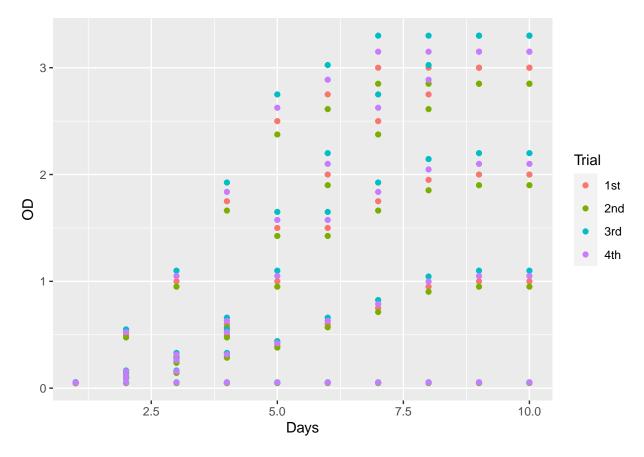
5. Plot Days vs OD as scatter plot using geom_point

```
ggplot(data = growth) +
geom_point(mapping = aes(x = Days, y = OD))
```



6. Plot Days vs OD as scatter plot and add a label by mapping the variable Trial to color

```
ggplot(data = growth) +
geom_point(mapping = aes(x = Days, y = OD, color = Trial))
```



7. Try to create new plot(s) of your own design. Consider what you think would be useful for the project or simply test out different different functions and settings. You can use the cheat sheet at https://rstudio.com/wp-content/uploads/2016/11/ggplot2-cheatsheet-2.1.pdf

```
ggplot(data = growth) +
  geom_smooth(mapping = aes(x = Days, y = OD, color = Iron))
```

`geom_smooth()` using method = 'loess' and formula 'y ~ x'

