

functions_and_iteration

Functions and Iterations by Molly and Eric

```
#Reading in packages we need  
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --  
v dplyr      1.1.4      v readr      2.1.6  
v forcats    1.0.1      v stringr    1.6.0  
v ggplot2     4.0.1      v tibble     3.3.1  
v lubridate  1.9.4      v tidyr      1.3.2  
v purrr       1.2.0  
-- Conflicts ----- tidyverse_conflicts() --  
x dplyr::filter() masks stats::filter()  
x dplyr::lag()     masks stats::lag()  
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
library(here)
```

here() starts at /Users/ericmedina/functions_and_iteration

Building a function

```
# Create a new function called add_one  
# x will be the only input to the function  
add_one <- function(x){  
  
  # Add x and 1 together, store as the object "output"  
  answer <- x + 1
```

```
# Print out whatever is stored in "output"
return(answer)

}

add_one(x=4)
```

[1] 5

```
#working with multiple arguments

# Create a new function called add_together
# x and y will be the two arguments to the function
add_together <- function(x, y){

  # Add x and y together, store as the object "output"
  output <- x + y

  # Print out whatever is stored in "output"
  return(output)

}
```

Q1.1 Feed our new function two numbers

```
add_together(3,5)
```

[1] 8

Q1.2 Feed our new function a number and a character string

```
add_together(3,"five")
```

non numeric argument in operator

Accessing source code

```
add_together
```

```
function (x, y)
{
  output <- x + y
  return(output)
}
```

Q1.3 Create your own function!

```
math_time <- function(x,y,z){
  output <- (x-y)^2/z
  return(output)
}

math_time(5,2,9)
```

```
[1] 1
```

Vectors as inputs

```
# Create a function called lbs_to_kg that takes a data object 'weights' as input
lbs_to_kg <- function(weights){

  # Multiply weights by 0.454, store as the object "output"
  output <- weights*0.454

  # Print out whatever is stored in "output"
  return(output)
}

#Vector of bison weights
bison <- c(1000, 800, 1200, 1400)

#Feeding it to our function

lbs_to_kg(weights = bison)
```

```
[1] 454.0 363.2 544.8 635.6
```

Q1.4 Calculate deviation from a mean

```
deviation <- function(cow){  
  output <- cow-mean(cow)  
  return(output)  
}  
deviation(cow = bison)
```

```
[1] -100 -300 100 300
```

Working with iterations

```
# Look at the first 6 rows of iris  
head(iris)
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa

Q2.1: What are the units that the sepal and petal columns are measured in?

The measurements are in centimeters.

```
#Getting averages of all measurements by species  
iris %>%  
  group_by(Species) %>%  
  summarize(Sepal.Length = mean(Sepal.Length),  
            Sepal.Width = mean(Sepal.Width),  
            Petal.Length = mean(Petal.Length),  
            Petal.Width = mean(Petal.Width))
```

```
# A tibble: 3 x 5
  Species    Sepal.Length Sepal.Width Petal.Length Petal.Width
  <fct>         <dbl>         <dbl>         <dbl>         <dbl>
1 setosa         5.01           3.43           1.46           0.246
2 versicolor     5.94           2.77           4.26           1.33
3 virginica      6.59           2.97           5.55           2.03
```

```
#Using the across function
```

```
iris %>%
  group_by(Species) %>%
  summarize(across(.cols = c(Sepal.Length, Sepal.Width, Petal.Length, Petal.Width),
    .fns = mean))
```

```
# A tibble: 3 x 5
  Species    Sepal.Length Sepal.Width Petal.Length Petal.Width
  <fct>         <dbl>         <dbl>         <dbl>         <dbl>
1 setosa         5.01           3.43           1.46           0.246
2 versicolor     5.94           2.77           4.26           1.33
3 virginica      6.59           2.97           5.55           2.03
```

```
#Some tricks for selecting columns
```

```
iris %>%
  group_by(Species) %>%
  summarize(across(.cols = Sepal.Length:Petal.Width,
    .fns = mean))
```

```
# A tibble: 3 x 5
  Species    Sepal.Length Sepal.Width Petal.Length Petal.Width
  <fct>         <dbl>         <dbl>         <dbl>         <dbl>
1 setosa         5.01           3.43           1.46           0.246
2 versicolor     5.94           2.77           4.26           1.33
3 virginica      6.59           2.97           5.55           2.03
```

```
#Using 'everything'
```

```
iris %>%
  group_by(Species) %>%
  summarize(across(.cols = everything(),
    .fns = mean))
```

```
# A tibble: 3 x 5
  Species    Sepal.Length Sepal.Width Petal.Length Petal.Width
  <fct>          <dbl>      <dbl>      <dbl>      <dbl>
1 setosa         5.01        3.43        1.46        0.246
2 versicolor     5.94        2.77        4.26        1.33
3 virginica      6.59        2.97        5.55        2.03
```

Q2.2 Summarize to calculate the median across all columns

```
iris %>%
  group_by(Species) %>%
  summarize(across(.cols = where(is.numeric),
                           .fns = median))
```

```
# A tibble: 3 x 5
  Species    Sepal.Length Sepal.Width Petal.Length Petal.Width
  <fct>          <dbl>      <dbl>      <dbl>      <dbl>
1 setosa         5        3.4        1.5        0.2
2 versicolor     5.9        2.8        4.35       1.3
3 virginica      6.5         3        5.55       2
```

Q2.3 Summarize to calculate the mean across all numeric columns in the cereal data

```
cereal <- read.csv("data/cereal.csv")

cereal %>%
  group_by(mfr) %>%
  summarize(across(.cols = where(is.numeric),
                           .fns = mean))
```

```
# A tibble: 7 x 14
  mfr      calories protein  fat sodium fiber carbo sugars potass vitamins shelf
  <chr>      <dbl>   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>   <dbl> <dbl>
1 Americ~    100     4     1     0     0    16     3     95     25     2
2 Genera~   111.    2.32  1.36  200.    1.27  14.7    7.95   85.2    35.2    2.14
3 Kellog~   109.    2.65  0.609 175.    2.74  15.1    7.57  103.    34.8    2.35
4 Nabisco    86.7    2.83  0.167  37.5     4    16     1.83  121.    8.33    1.67
5 Post     109.    2.44  0.889 146.    2.78  13.2    8.78  114.    25     2.44
```

```
6 Quaker~      95      2.62 1.75      92.5 1.34 10      5.25 74.4      12.5 2.38
7 Ralsto~     115      2.5 1.25     198.  1.88 17.6  6.12 89.2      25    2
# i 3 more variables: weight <dbl>, cups <dbl>, rating <dbl>
```

Practicing Loops

```
#The most basic example of a loop
```

```
for (i in 1:5) {
  # Print out whatever the value of i is
  print(i)
}
```

```
[1] 1
[1] 2
[1] 3
[1] 4
[1] 5
```

```
#Manually running first two iterations of the loop
```

```
# First iteration where i = 1:
i <- 1
print(i)
```

```
[1] 1
```

```
# Second iteration where i = 2
i <- 2
print(i)
```

```
[1] 2
```

```
#Using mutiplication in our loop
for (i in 1:5) {
  print(i*2)
}
```

```
[1] 2
[1] 4
[1] 6
[1] 8
[1] 10
```

Q2.4 Create a for loop that iterates from 1 to 10 and squares the value of the index i

```
for (i in 1:10) {
  # Print out whatever the value of i is
  print(i^2)
}
```

```
[1] 1
[1] 4
[1] 9
[1] 16
[1] 25
[1] 36
[1] 49
[1] 64
[1] 81
[1] 100
```

Using a for loop to simulate data

```
N0 = 100 #initial population size

years = 20 #number of years into the future

N = vector(length = years) # create an empty vector to store pop. sizes

N[1] = N0 #initial population size should be the first N

lambda = 1.2 #growth rate

print(N)
```

```
[1] 100 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
[20] 0
```



```
N[5]
```

```
[1] 0
```

```
N[1]
```

```
[1] 100
```

```
# For every year t in 2 through 20 (remember, "years" also equals 20), apply the following equation
for (t in 2:20) {
  N[t] = N[t - 1] * lambda # Apply the equation
}

N
```

```
[1] 100.0000 120.0000 144.0000 172.8000 207.3600 248.8320 298.5984
[8] 358.3181 429.9817 515.9780 619.1736 743.0084 891.6100 1069.9321
[15] 1283.9185 1540.7022 1848.8426 2218.6111 2662.3333 3194.8000
```

Q2.5a Rerun the for loop with the following parameters:

```
N0 = 300
lambda = 0.95
years = 50
N = vector(length=years)
N[1] = N0
N
```

```
[1] 300 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
[20] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
[39] 0 0 0 0 0 0 0 0 0 0 0 0
```

```
for (t in 2:50) {
  N[t] = N[t - 1] * lambda}

N
```

```

[1] 300.00000 285.00000 270.75000 257.21250 244.35187 232.13428 220.52757
[8] 209.50119 199.02613 189.07482 179.62108 170.64003 162.10803 154.00262
[15] 146.30249 138.98737 132.03800 125.43610 119.16430 113.20608 107.54578
[22] 102.16849 97.06006 92.20706 87.59671 83.21687 79.05603 75.10323
[29] 71.34807 67.78066 64.39163 61.17205 58.11345 55.20777 52.44738
[36] 49.82502 47.33376 44.96708 42.71872 40.58279 38.55365 36.62596
[43] 34.79467 33.05493 31.40219 29.83208 28.34047 26.92345 25.57728
[50] 24.29841

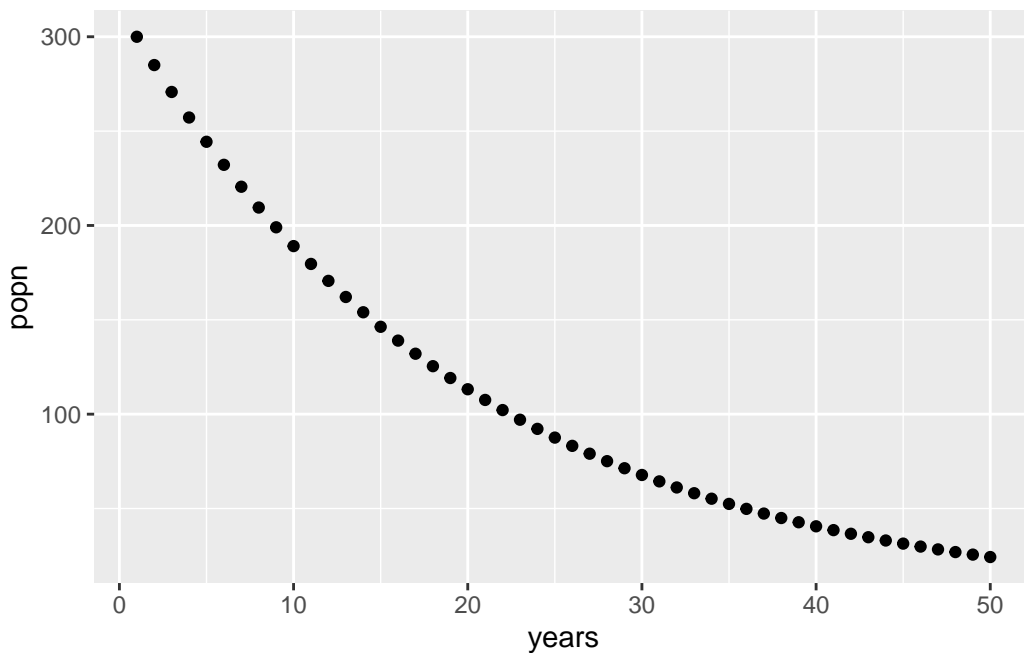
```

Q2.5b Plot the data and interpret

```

popn_data <- tibble(years = 1:years, # Make the years column = 1, 2, 3, ..., 20
                    popn = N)
popn_data %>%
  ggplot(aes(x=years,
             y=popn))+
  geom_point()

```



The population is decreasing almost exponentially.

Looping through data frames

```
head(iris)
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa

```
#extracting elements  
#getting the first row  
iris[1,]
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa

```
#getting the data in the third column as a vector  
iris[,3]
```

```
[1] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4  
[19] 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2  
[37] 1.3 1.4 1.3 1.5 1.3 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0  
[55] 4.6 4.5 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5 4.1 4.5 3.9 4.8 4.0  
[73] 4.9 4.7 4.3 4.4 4.8 5.0 4.5 3.5 3.8 3.7 3.9 5.1 4.5 4.5 4.7 4.4 4.1 4.0  
[91] 4.4 4.6 4.0 3.3 4.2 4.2 4.2 4.3 3.0 4.1 6.0 5.1 5.9 5.6 5.8 6.6 4.5 6.3  
[109] 5.8 6.1 5.1 5.3 5.5 5.0 5.1 5.3 5.5 6.7 6.9 5.0 5.7 4.9 6.7 4.9 5.7 6.0  
[127] 4.8 4.9 5.6 5.8 6.1 6.4 5.6 5.1 5.6 6.1 5.6 5.5 4.8 5.4 5.6 5.1 5.1 5.9  
[145] 5.7 5.2 5.0 5.2 5.4 5.1
```

```
#getting exact value in the first row and third column  
iris[1,3]
```

```
[1] 1.4
```

```
#iteratively performing each action
for (i in 1:5) {

  # This prints out a statement saying "Here's column i",
  #but the i gets replaced with the number that it's currently at
  print(paste("Here's column",i))

  # This prints out column i
  print(iris[,i])
}
```

```
[1] "Here's column 1"
```

```
[1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9 5.4 4.8 4.8 4.3 5.8 5.7 5.4 5.1
[19] 5.7 5.1 5.4 5.1 4.6 5.1 4.8 5.0 5.0 5.2 5.2 4.7 4.8 5.4 5.2 5.5 4.9 5.0
[37] 5.5 4.9 4.4 5.1 5.0 4.5 4.4 5.0 5.1 4.8 5.1 4.6 5.3 5.0 7.0 6.4 6.9 5.5
[55] 6.5 5.7 6.3 4.9 6.6 5.2 5.0 5.9 6.0 6.1 5.6 6.7 5.6 5.8 6.2 5.6 5.9 6.1
[73] 6.3 6.1 6.4 6.6 6.8 6.7 6.0 5.7 5.5 5.5 5.8 6.0 5.4 6.0 6.7 6.3 5.6 5.5
[91] 5.5 6.1 5.8 5.0 5.6 5.7 5.7 6.2 5.1 5.7 6.3 5.8 7.1 6.3 6.5 7.6 4.9 7.3
[109] 6.7 7.2 6.5 6.4 6.8 5.7 5.8 6.4 6.5 7.7 7.7 6.0 6.9 5.6 7.7 6.3 6.7 7.2
[127] 6.2 6.1 6.4 7.2 7.4 7.9 6.4 6.3 6.1 7.7 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8
[145] 6.7 6.7 6.3 6.5 6.2 5.9
```

```
[1] "Here's column 2"
```

```
[1] 3.5 3.0 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 3.7 3.4 3.0 3.0 4.0 4.4 3.9 3.5
[19] 3.8 3.8 3.4 3.7 3.6 3.3 3.4 3.0 3.4 3.5 3.4 3.2 3.1 3.4 4.1 4.2 3.1 3.2
[37] 3.5 3.6 3.0 3.4 3.5 2.3 3.2 3.5 3.8 3.0 3.8 3.2 3.7 3.3 3.2 3.2 3.1 2.3
[55] 2.8 2.8 3.3 2.4 2.9 2.7 2.0 3.0 2.2 2.9 2.9 3.1 3.0 2.7 2.2 2.5 3.2 2.8
[73] 2.5 2.8 2.9 3.0 2.8 3.0 2.9 2.6 2.4 2.4 2.7 2.7 3.0 3.4 3.1 2.3 3.0 2.5
[91] 2.6 3.0 2.6 2.3 2.7 3.0 2.9 2.9 2.5 2.8 3.3 2.7 3.0 2.9 3.0 3.0 2.5 2.9
[109] 2.5 3.6 3.2 2.7 3.0 2.5 2.8 3.2 3.0 3.8 2.6 2.2 3.2 2.8 2.8 2.7 3.3 3.2
[127] 2.8 3.0 2.8 3.0 2.8 3.8 2.8 2.8 2.6 3.0 3.4 3.1 3.0 3.1 3.1 3.1 2.7 3.2
[145] 3.3 3.0 2.5 3.0 3.4 3.0
```

```
[1] "Here's column 3"
```

```
[1] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4
[19] 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2
[37] 1.3 1.4 1.3 1.5 1.3 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0
[55] 4.6 4.5 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5 4.1 4.5 3.9 4.8 4.0
[73] 4.9 4.7 4.3 4.4 4.8 5.0 4.5 3.5 3.8 3.7 3.9 5.1 4.5 4.5 4.7 4.4 4.1 4.0
[91] 4.4 4.6 4.0 3.3 4.2 4.2 4.2 4.3 3.0 4.1 6.0 5.1 5.9 5.6 5.8 6.6 4.5 6.3
[109] 5.8 6.1 5.1 5.3 5.5 5.0 5.1 5.3 5.5 6.7 6.9 5.0 5.7 4.9 6.7 4.9 5.7 6.0
[127] 4.8 4.9 5.6 5.8 6.1 6.4 5.6 5.1 5.6 6.1 5.6 5.5 4.8 5.4 5.6 5.1 5.1 5.9
[145] 5.7 5.2 5.0 5.2 5.4 5.1
```

```
[1] "Here's column 4"
```

```

[1] 0.2 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 0.2 0.2 0.1 0.1 0.2 0.4 0.4 0.3
[19] 0.3 0.3 0.2 0.4 0.2 0.5 0.2 0.2 0.4 0.2 0.2 0.2 0.2 0.2 0.4 0.1 0.2 0.2 0.2
[37] 0.2 0.1 0.2 0.2 0.3 0.3 0.2 0.6 0.4 0.3 0.2 0.2 0.2 0.2 0.2 1.4 1.5 1.5 1.3
[55] 1.5 1.3 1.6 1.0 1.3 1.4 1.0 1.5 1.0 1.4 1.3 1.4 1.5 1.0 1.5 1.1 1.8 1.3
[73] 1.5 1.2 1.3 1.4 1.4 1.7 1.5 1.0 1.1 1.0 1.2 1.6 1.5 1.6 1.5 1.3 1.3 1.3
[91] 1.2 1.4 1.2 1.0 1.3 1.2 1.3 1.3 1.1 1.3 2.5 1.9 2.1 1.8 2.2 2.1 1.7 1.8
[109] 1.8 2.5 2.0 1.9 2.1 2.0 2.4 2.3 1.8 2.2 2.3 1.5 2.3 2.0 2.0 1.8 2.1 1.8
[127] 1.8 1.8 2.1 1.6 1.9 2.0 2.2 1.5 1.4 2.3 2.4 1.8 1.8 2.1 2.4 2.3 1.9 2.3
[145] 2.5 2.3 1.9 2.0 2.3 1.8

```

```
[1] "Here's column 5"
```

```

[1] setosa      setosa      setosa      setosa      setosa      setosa
[7] setosa      setosa      setosa      setosa      setosa      setosa
[13] setosa      setosa      setosa      setosa      setosa      setosa
[19] setosa      setosa      setosa      setosa      setosa      setosa
[25] setosa      setosa      setosa      setosa      setosa      setosa
[31] setosa      setosa      setosa      setosa      setosa      setosa
[37] setosa      setosa      setosa      setosa      setosa      setosa
[43] setosa      setosa      setosa      setosa      setosa      setosa
[49] setosa      setosa      versicolor  versicolor  versicolor  versicolor
[55] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[61] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[67] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[73] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[79] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[85] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[91] versicolor  versicolor  versicolor  versicolor  versicolor  versicolor
[97] versicolor  versicolor  versicolor  versicolor  virginica   virginica
[103] virginica   virginica   virginica   virginica   virginica   virginica
[109] virginica   virginica   virginica   virginica   virginica   virginica
[115] virginica   virginica   virginica   virginica   virginica   virginica
[121] virginica   virginica   virginica   virginica   virginica   virginica
[127] virginica   virginica   virginica   virginica   virginica   virginica
[133] virginica   virginica   virginica   virginica   virginica   virginica
[139] virginica   virginica   virginica   virginica   virginica   virginica
[145] virginica   virginica   virginica   virginica   virginica   virginica

```

Levels: setosa versicolor virginica

```
#taking the mean of each function
```

```
for (i in 1:4) {
```

```

  # This prints out a statement saying "Here's column i", but the i gets replaced with the n
  print(paste("Here's column",i))

```

```
# This prints out column i
print(mean(iris[,i]))
}
```

```
[1] "Here's column 1"
[1] 5.843333
[1] "Here's column 2"
[1] 3.057333
[1] "Here's column 3"
[1] 3.758
[1] "Here's column 4"
[1] 1.199333
```

```
#Using across to do the same
iris %>%
  summarize(across(.cols = 1:4,
                    .fns = mean))
```

```
      Sepal.Length Sepal.Width Petal.Length Petal.Width
1      5.843333      3.057333      3.758      1.199333
```

Q2.6 Which do you prefer?

In this instance I prefer across as it maintains the table format.

```
#Adding names to the four loop
for (i in 1:4) {
  # Fetch the column names of the dataframe, store in a vector "names"
  names <- colnames(iris)

  # Print out the "i"th element of the vector to print alongside the output
  print(names[i])

  # This prints out column i
  print(mean(iris[,i]))
}
```

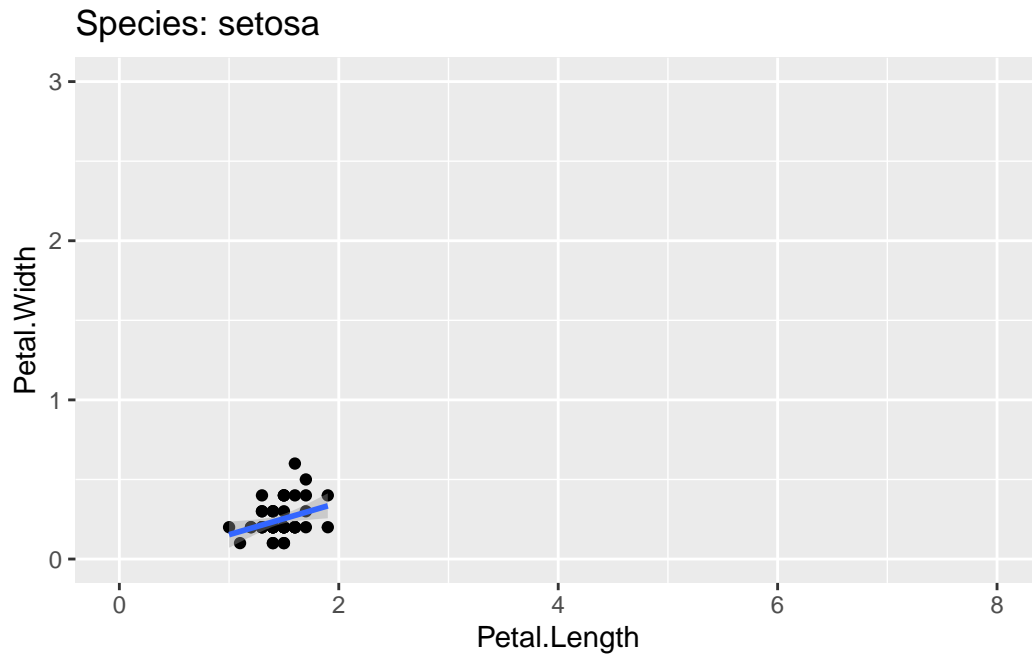
```
[1] "Sepal.Length"
[1] 5.843333
```

```
[1] "Sepal.Width"  
[1] 3.057333  
[1] "Petal.Length"  
[1] 3.758  
[1] "Petal.Width"  
[1] 1.199333
```

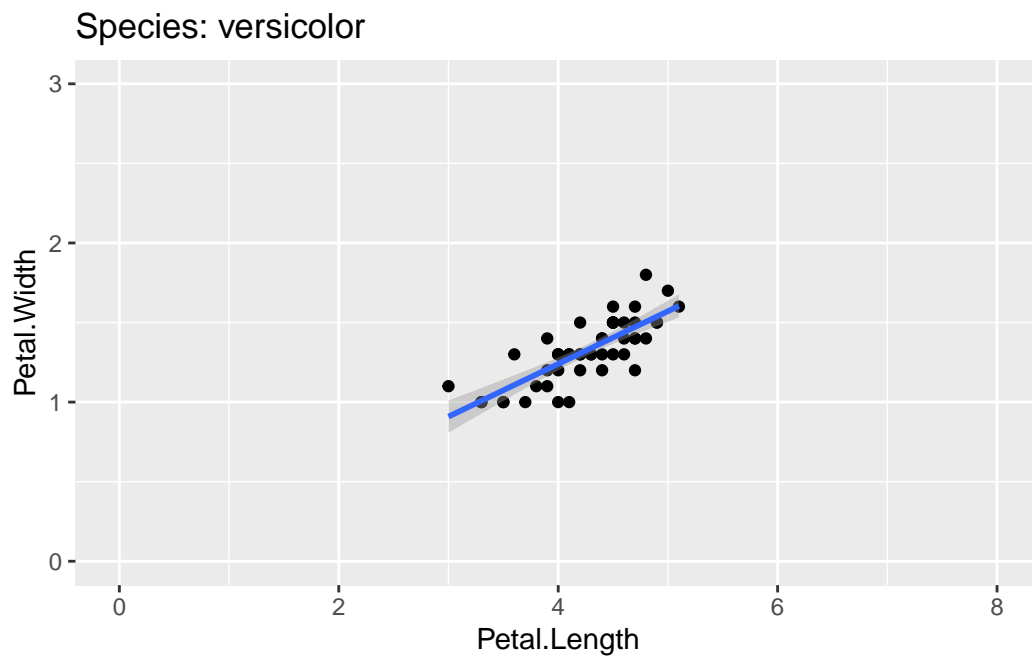
Q2.7 Annotate this code with what you think it's doing

```
# Store a vector of unique species names from the Species column of Iris  
spp_names <- unique(iris$Species)  
  
# Create a loop that iterates across the total number of species  
for (i in 1:length(spp_names)) {  
  
  filt_data <- iris %>%  
    # Filters to species [i] indexed species name  
    filter(Species == spp_names[i])  
  
  # Plots filtered data of species [i]  
  plot <- filt_data %>%  
    # Petal length on the x-axis and width on the y-axis  
    ggplot(aes(x = Petal.Length,  
               y = Petal.Width)) +  
    # Creates points  
    geom_point() +  
    # Creates a linear model showing the relationship across points  
    geom_smooth(method = "lm") +  
    # Limits x and y to specified values  
    lims(x = c(0,8),  
          y = c(0,3)) +  
    # Titles plot with species of [i]  
    ggtitle(paste("Species:", spp_names[i]))  
  
  # Shows the plot  
  print(plot)  
}
```

`geom_smooth()` using formula = 'y ~ x'



``geom_smooth()`` using formula = 'y ~ x'



``geom_smooth()`` using formula = 'y ~ x'

Species: virginica

