

# **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 becom
```

```
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 four 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 0 0
[20] 0 0 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 0 0 0 0 0 0 0 0 0
```

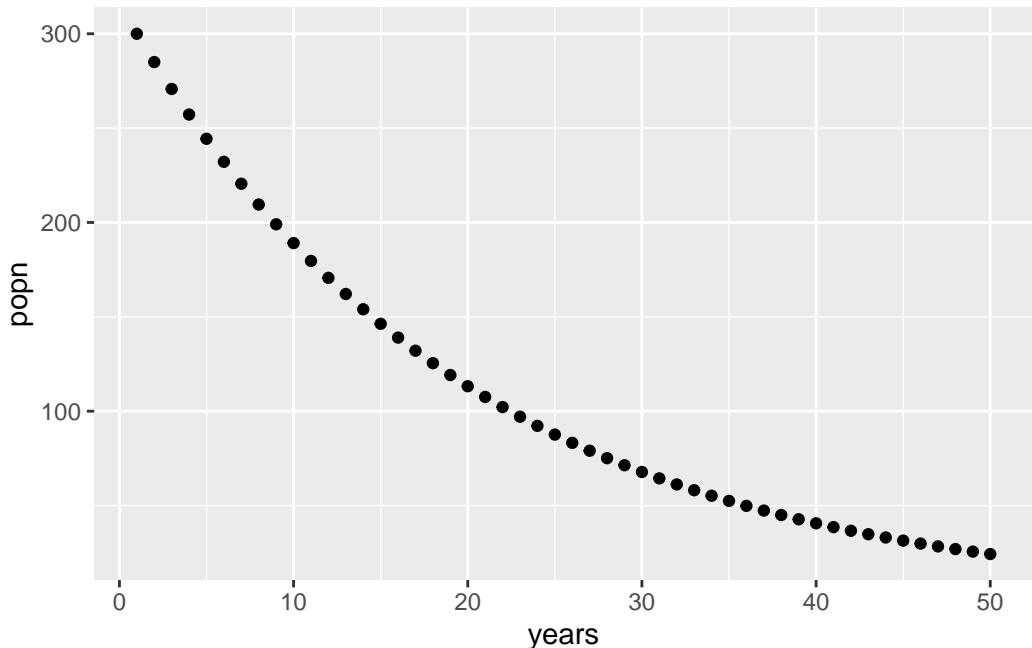
```
for (t in 2:50) {
  N[t] = N[t - 1] * lambda}
```

```
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 2.6 3.0 3.4 3.1 3.0 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 0.4 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 0.4 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 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 numnber of species
for (i in 1:length(spp_names)) {

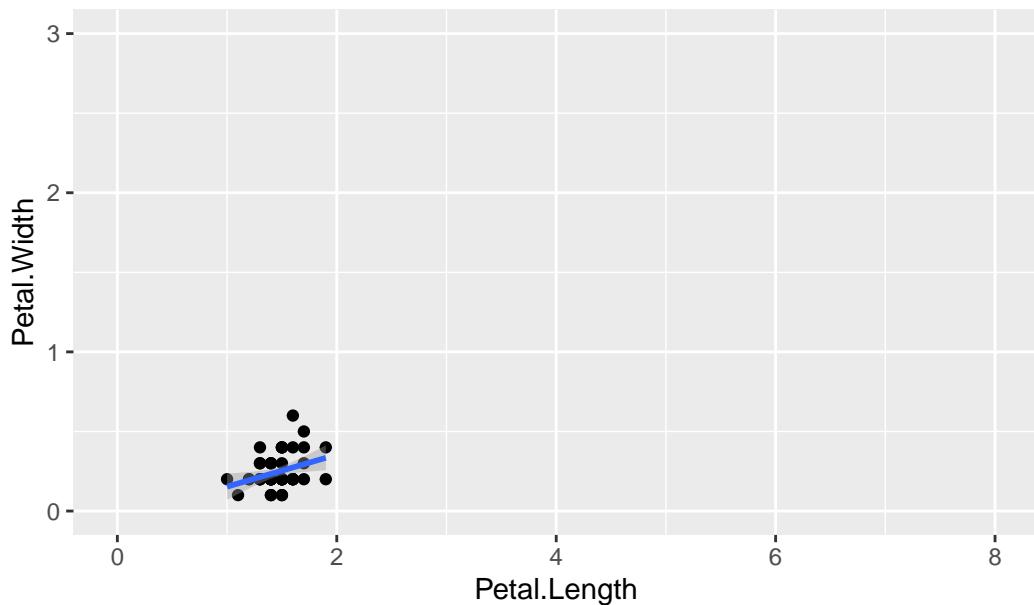
  filt_data <- iris %>%
    #Filters to species [i] indexed species name
    filter(Species == spp_names[i])

  # Plots filtered data of species [1]
  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)
}

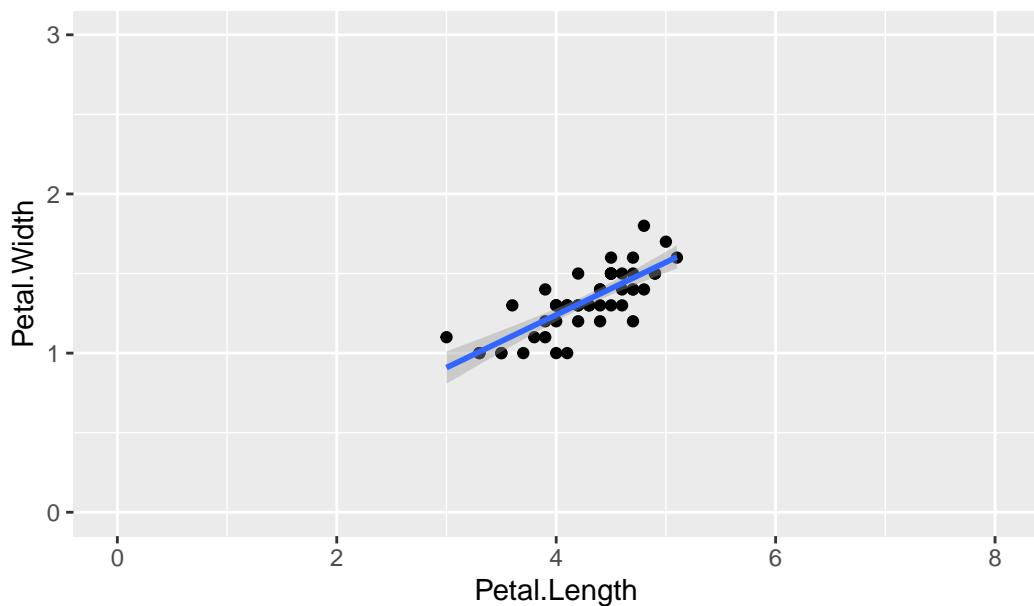
`geom_smooth()` using formula = 'y ~ x'
```

Species: setosa



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

Species: versicolor



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

Species: virginica

