



EXECUTIVE SUMMARY REPORT 2

Module 2 assignment

Abstract

In this assignment, I will create a summary of some data and learn how to use R

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Objective:

The objective of this assignment is to get used to following

- Calculate basic descriptive statistics to describe a set of data
- Create various types of graphs based on data provided
- Use R to visualize data

Introduction

In this summary, I will use the data from the BullTroutRML2 dataset and use the APA system for the bibliography. Also, the R code that I used to process the data is available on my GitHub account, which its address is mentioned in the bibliography part. I also put my code in the appendix part.

1. Printing my name at the top of the script
 - `print("Plotting Basics:Movahedi")`
2. Importing and loading required libraries
 - `install.packages("FSA")`
 - `install.packages("FSAdata")`
 - `install.packages("magrittr")`
 - `install.packages("dplyr")`
 - `install.packages("plotrix")`
 - `install.packages("ggplot2")`
 - `install.packages("moments")`
 - `#loading installed libraries`
 - `library(FSA)`
 - `library(FSAdata)`
 - `library(magrittr)`
 - `library(dplyr)`
 - `library(plotrix)`
 - `library(ggplot2)`
 - `library(moments)`
3. Loading the BullTroutRML2 dataset
 - `data(BullTroutRML2)`
4. Printing the first and last 3 records from the BullTroutRMS2 dataset
 - `#showing first three lines`

- `print(head(BullTroutRML2, n = 3L))`
- `#showing last three lines`
- `print(tail(BullTroutRML2, n = 3L))`

```

      age  fl    lake    era
1   14 459 Harrison 1977-80
2   12 449 Harrison 1977-80
3   10 471 Harrison 1977-80

```

```

      age  fl    lake    era
94    4 298 Osprey 1997-01
95    3 279 Osprey 1997-01
96    3 273 Osprey 1997-01

```

5. Removing all records except those from Harrison Lake

- `#make a copy of original dataset so my mistakes don't affect original data`
- `Harrison_Lake <- BullTroutRML2`
- `#cheak the structure of dataset to be able to set filter`
- `str(Harrison_Lake)`
- `#set the filter`
- `Harrison_Lake <- dplyr::filter(Harrison_Lake, lake == "Harrison")`

6. Displaying the first and last 5 records from the filtered BullTroutRML2 dataset

- `#display first and last 5 lines`
- `head(Harrison_Lake, n = 5L)`
- `tail(Harrison_Lake, n = 5L)`

```

      age  fl    lake    era
1   14 459 Harrison 1977-80
2   12 449 Harrison 1977-80
3   10 471 Harrison 1977-80
4   10 446 Harrison 1977-80
5    9 400 Harrison 1977-80

```

```

      age  fl    lake    era
57    0  41 Harrison 1997-01
58    0  20 Harrison 1997-01
59    7 245 Harrison 1997-01
60    7 279 Harrison 1997-01
61    5 245 Harrison 1997-01

```

7. Displaying the structure of the filtered BullTroutRML2dataset

```
- str(Harrison_Lake)
```

```
'data.frame': 61 obs. of 4 variables:
 $ age : int 14 12 10 10 9 9 9 8 8 7 ...
 $ fl : int 459 449 471 446 400 440 462 480 449 437 ...
 $ lake: Factor w/ 2 levels "Harrison","Osprey": 1 1 1 1 1 1 1 1 1 1 ...
 $ era : Factor w/ 2 levels "1977-80","1997-01": 1 1 1 1 1 1 1 1 1 1 ...
```

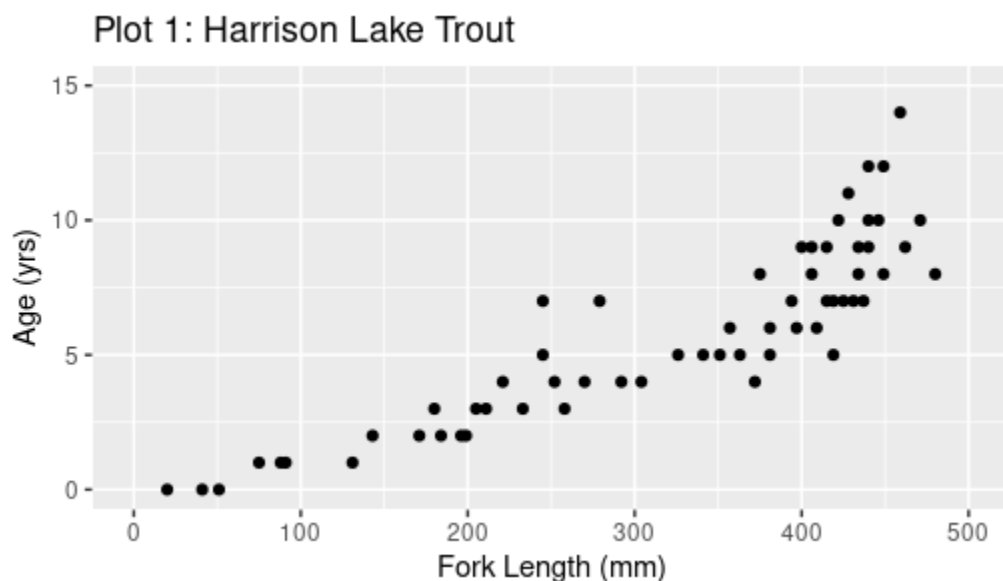
8. Display the summary of the filtered BullTroutRML2dataset

```
- summary(Harrison_Lake)
```

age	fl	lake	era
Min. : 0.000	Min. : 20	Harrison:61	1977-80:23
1st Qu.: 3.000	1st Qu.:221	Osprey : 0	1997-01:38
Median : 6.000	Median :372		
Mean : 5.754	Mean :319		
3rd Qu.: 8.000	3rd Qu.:425		
Max. :14.000	Max. :480		

9. creating Plot 1: Harrison Lake Trout

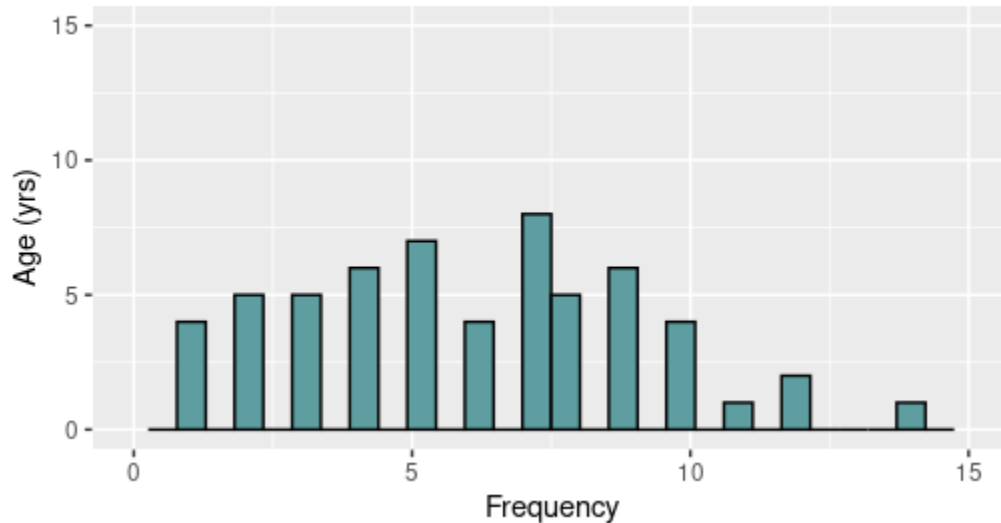
```
- ggplot(data = Harrison_Lake) +
- geom_point(mapping = aes(x = fl, y = age))+
- xlim(0, 500) + ylim(0,15)+
- xlab("Fork Length (mm)") + ylab("Age (yrs)") +
- ggtitle("Plot 1: Harrison Lake Trout")
```



10. Plotting an "Age" histogram

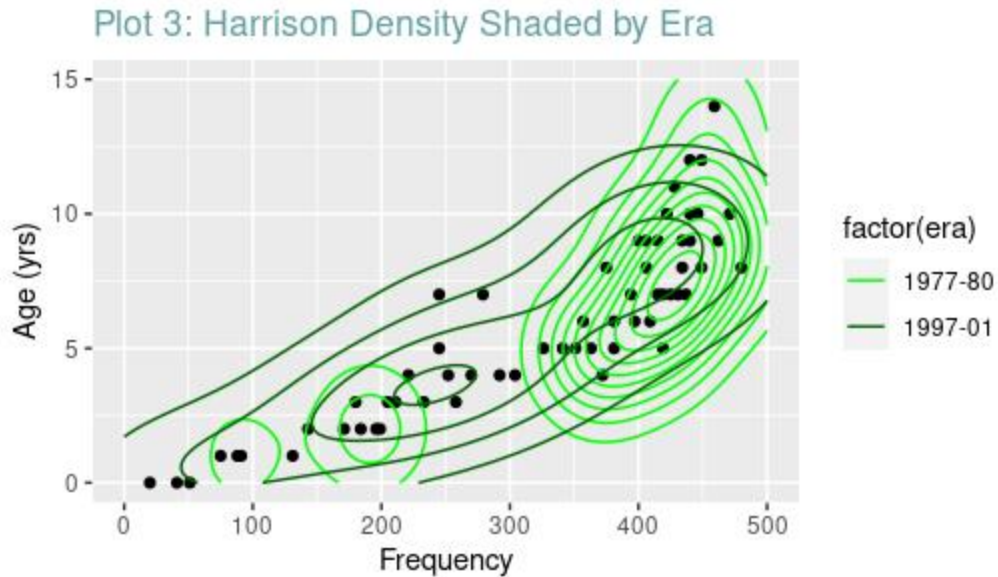
- `ggplot(data = Harrison_Lake, aes(x=age)) + geom_histogram(color="black", fill="cadetblue")+`
- `xlab("Frequency")+ylab("Age (yrs)")+`
- `labs(title="Plot 2: Harrison Fish Age Distribution")+`
- `xlim(0,15) + ylim(0,15)+`
- `theme(plot.title = element_text(color = "cadetblue"))`

Plot 2: Harrison Fish Age Distribution



11. Creating an over dense plot using the same specifications as the previous scatterplot

- `#Create 2d plot`
-
- `p <- ggplot(data = Harrison_Lake, aes(y=age,x=fl))+`
- `xlab("Frequency")+ylab("Age (yrs)")+`
- `labs(title="Plot 3: Harrison Density Shaded by Era")+`
- `theme(plot.title = element_text(color = "cadetblue"))+`
- `xlim(0,500) + ylim(0,15)+`
- `geom_point(colour = "black")`
-
- `# color 2d plot`
- `p +`
- `stat_density2d(aes(color = factor(era))) +`
- `scale_colour_manual(values = c("green", "darkgreen"))`



12. Creating a new object called “tmp” that includes the first 3 and last 3 records of the BullTroutRML2 data set

- tmp <- head(Harrison_Lake, n = 3L)
- tmp <- rbind(tmp,tail(Harrison_Lake, n = 3L))

13. Display the “era” column (variable) in the new “tmp” object

- era <- subset(tmp,select=c("era"))
- print(era)

```

      era
1 1977-80
2 1977-80
3 1977-80
59 1997-01
60 1997-01
61 1997-01

```

14. Creating a pchs vector with the argument values for + and x.

- pchs <- c("+","x")

15. Creating a cols vector with the two elements “red” and “gray60”

- cols <- c("red","gray60")

16. Convert the tmp era values to numeric values.

- tmp\$era <- as.numeric(as.factor(tmp\$era))

17. Initialize the cols vector with the tmp era values

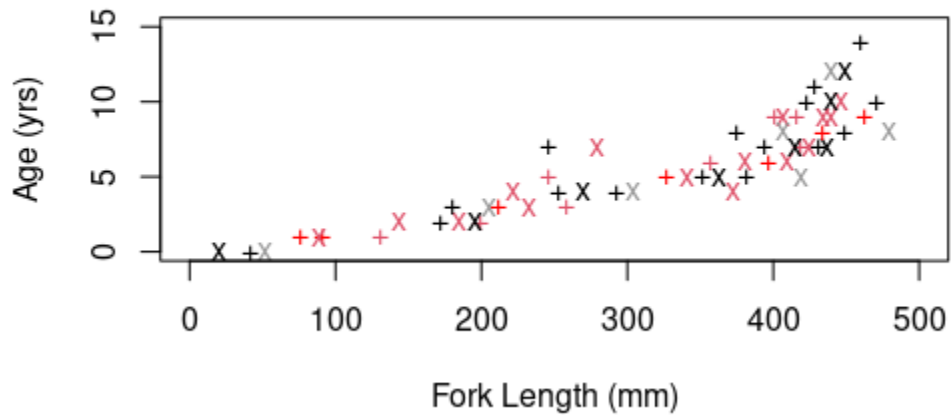
- cols <- append(tmp\$era,cols)

18. creating Plot 4: Symbol & Color by Era

- plot(y=Harrison_Lake\$age,x=Harrison_Lake\$f1,main="Plot 4: Symbol & Color by Era",

- ylab="Age (yrs)",xlab="Fork Length (mm)",
- ylim=c(0,15),xlim=c(0,500),
- pch=pchs,col=cols)

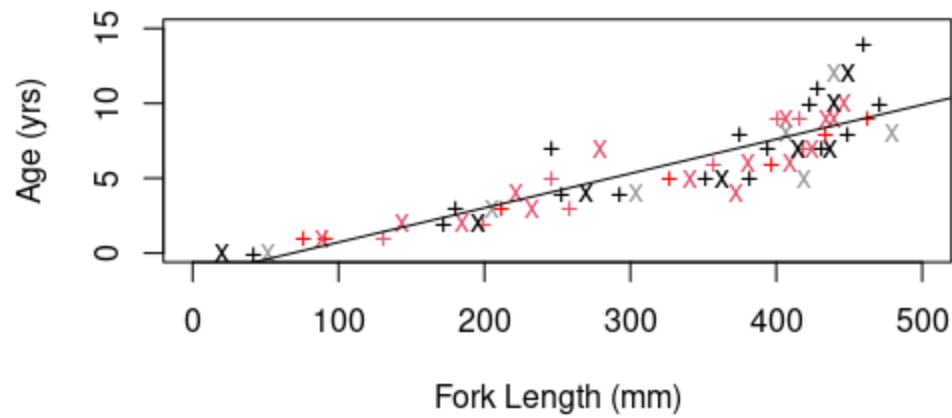
Plot 4: Symbol & Color by Era



19. Plot 5: Regression Overlay

```
- plot(y=Harrison_Lake$age,x=Harrison_Lake$fl,main=""Plot 5: Regression
,"Overlay
,"ylab="Age (yrs)",xlab="Fork Length (mm)
,(. 'd . . )ylim=c(0,15),xlim=c
pch=pchs,col=cols)+abline(lm(Harrison_Lake$age ~ Harrison_Lake$fl))
```

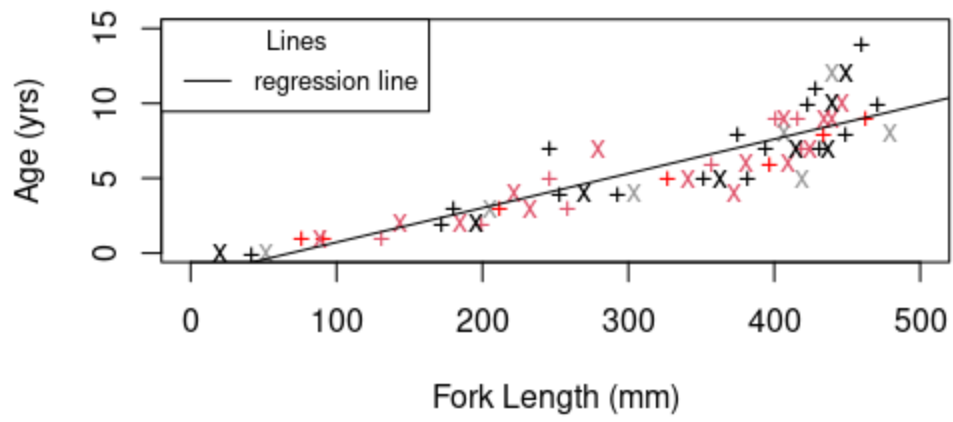
**“Plot 5: Regression
Overlay**



20. Place a legend of on Plot 5 and call the new graph “Plot 6: :Legend Overlay”

```
, "- plot(y=Harrison_Lake$age,x=Harrison_Lake$fl,main=""Plot 6: :Legend Overlay
,"ylab="Age (yrs)",xlab="Fork Length (mm)
,(. 'o . .)ylim=c(0,15),xlim=c
+((pch=pchs,col=cols)+abline(lm(Harrison_Lake$age ~ Harrison_Lake$fl
,("legend("topleft", legend=c("regression line
,col=c("black"), lty=1:2, cex=0.8
title="Lines", text.font=1, bg="white")
```


Plot 6: :Legend Overlay



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Appendix

Print my name at the top of the script#

```
print("Plotting Basics:Movahedi")
```

Importing required libraries#

```
install.packages("FSA")
```

```
install.packages("FSAdata")
```

```
install.packages("magrittr")
```

```
install.packages("dplyr")
```

```
install.packages("plotrix")
```

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

```
install.packages("moments")
```

loading installed libraries#

```
library(FSA)
```

```
library(FSAdata)
```

```
library(magrittr)
```

```
library(dplyr)
```

```
library(plotrix)
```

```
library(ggplot2)
```

```
library(moments)
```

loading data set#

```
data(BullTroutRML2)
```

showing first three lines#

```
print(head(BullTroutRML2, n = 3L))
```

```
showing last three lines#
```

```
print(tail(BullTroutRML2, n = 3L))
```

```
make a copy of original dataset so my mistakes don't affect original data#
```

```
Harrison_Lake <- BullTroutRML2
```

```
cheak the structure of dataset to be able to set filter#
```

```
str(Harrison_Lake)
```

```
set the filter#
```

```
Harrison_Lake <- dplyr::filter(Harrison_Lake, lake == "Harrison")
```

```
display first and last 5 lines#
```

```
head(Harrison_Lake, n = 5L)
```

```
tail(Harrison_Lake, n = 5L)
```

```
displaying structure of the filtered BullTroutRML2 data set#
```

```
str(Harrison_Lake)
```

```
Display the summary of the filtered BullTroutRML2 data set #
```

```
summary(Harrison_Lake)
```

```
creating scatter plot#
```

```
determining limits#
```

```
determining labels#
```

```
+ ggplot(data = Harrison_Lake)
```

```
+geom_point(mapping = aes(x = fl, y = age))
```

```
+(·'∧)xlim(0, 500) + ylim
```

```
+xlab("Fork Length (mm)")+ylab("Age (yrs)")
```

```
ggtitle("Plot 1: Harrison Lake Trout")
```

```
ploting histogram for age#
```

```
+ggplot(data = Harrison_Lake, aes(x=age)) + geom_histogram(color="black", fill="cadetblue")
```

```
+xlab("Frequency")+ylab("Age (yrs)")
```

```
+labs(title="Plot 2: Harrison Fish Age Distribution")
```

```
+(. ' \ Δ)xlim(0,15) + ylim
```

```
theme(plot.title = element_text( color = "cadetblue"))
```

```
Create 2d plot#
```

```
+p <- ggplot(data = Harrison_Lake, aes(y=age,x=fl))
```

```
+xlab("Frequency")+ylab("Age (yrs)")
```

```
+labs(title="Plot 3: Harrison Density Shaded by Era")
```

```
+theme(plot.title = element_text( color = "cadetblue"))
```

```
+(. ' \ Δ)xlim(0,500) + ylim
```

```
geom_point(colour = "black")
```

```
color 2d plot #
```

```
+ p
```

```
+ stat_density2d(aes(color = factor(era)))
```

```
scale_colour_manual(values = c("green", "darkgreen"))
```

```
creating tmp object containing first and last 3 lines of BullTroutRML2#
```

```
tmp <- head(Harrison_Lake, n = 3L)
```

```

tmp <- rbind(tmp,tail(Harrison_Lake, n = 3L))

print(tmp)

display era only#

era <- subset(tmp,select=c("era"))

print(era)

creating pchs vector #

pchs <- c("+","x")

creating cols vector #

cols <- c("red","gray60")

turn era to numeric factor#

tmp$era <- as.numeric(as.factor(tmp$era))

Initialize the cols vector with the tmp era values #

cols <- append(tmp$era,cols)

create the + and x plot#

, "plot(y=Harrison_Lake$age,x=Harrison_Lake$fl,main="Plot 4: Symbol & Color by Era

,"ylab="Age (yrs)",xlab="Fork Length (mm)

,(. 'Δ . . )ylim=c(0,15),xlim=c

(pch=pchs,col=cols

create regression line plot 5#

plot(y=Harrison_Lake$age,x=Harrison_Lake$fl,main=""Plot 5: Regression

,"Overlay

,"ylab="Age (yrs)",xlab="Fork Length (mm)

,(. 'Δ . . )ylim=c(0,15),xlim=c

((pch=pchs,col=cols)+abline(lm(Harrison_Lake$age ~ Harrison_Lake$fl

```


create plot 6#

```
, "plot(y=Harrison_Lake$age,x=Harrison_Lake$fl,main="Plot 6: :Legend Overlay
```

```
, "ylab="Age (yrs)",xlab="Fork Length (mm)
```

```
,(·'∂· ·)ylim=c(0,15),xlim=c
```

```
+((pch=pchs,col=cols)+abline(lm(Harrison_Lake$age ~ Harrison_Lake$fl
```

```
,("legend("topleft", legend=c("regression line
```

```
,col=c("black"), lty=1:2, cex=0.8
```

```
("title="Lines", text.font=1, bg="white
```