Exercise 4 - Data Visualisation using R

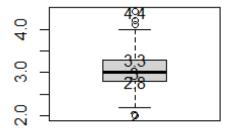
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06.02.2024

R Markdown

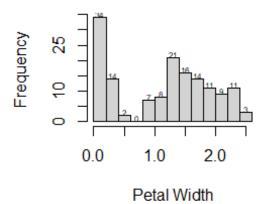
1. Extract the maximum information from a Histogram and Boxplot.

```
boxplot(iris$Sepal.Width)
text(y=fivenum(iris$Sepal.Width),x=1,labels = fivenum(iris$Sepal.Width))
```



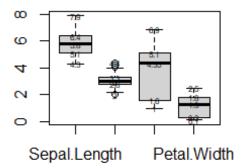
```
h <- hist(iris$Petal.Width,xlab="Petal Width",main="Petal Width")
text(h$mids,h$counts,labels=h$counts,cex=0.5, adj=c(0.5, 0))</pre>
```

Petal Width



2. Consider the iris dataset, create new variable called boxplot_data that excludes the species column.

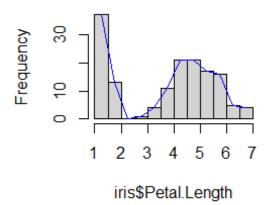
```
library(tidyverse)
boxplot_data <- iris %>% select(-Species)
boxplot(boxplot_data)
text(y=fivenum(boxplot_data$Sepal.Length),x=1,labels =
fivenum(iris$Sepal.Length),cex=0.5)
text(y=fivenum(boxplot_data$Sepal.Width),x=2,labels =
fivenum(iris$Sepal.Width),cex=0.5)
text(y=fivenum(boxplot_data$Petal.Length),x=3,labels =
fivenum(iris$Petal.Length),cex=0.5)
text(y=fivenum(boxplot_data$Petal.Width),x=4,labels =
fivenum(iris$Petal.Width),cex=0.5)
```



3. Create Histogram and illustrate the distribution look like within the petal length feature of the Iris data set.

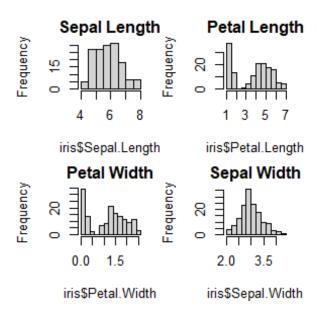
```
h<-hist(iris$Petal.Length)
mid <- (h$breaks[-1] + h$breaks[-length(h$breaks)]) / 2
lines(mid, h$counts, col = "blue")</pre>
```

Histogram of iris\$Petal.Lengt



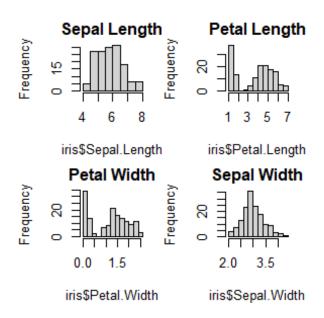
4. Create Histogram that plot multiple features at once for any dataset.

```
par(mfrow=c(2, 2), mar=c(4, 4, 2, 1))
hist(iris$Sepal.Length, main="Sepal Length")
hist(iris$Petal.Length, main="Petal Length")
hist(iris$Petal.Width, main="Petal Width")
hist(iris$Sepal.Width, main="Sepal Width")
```



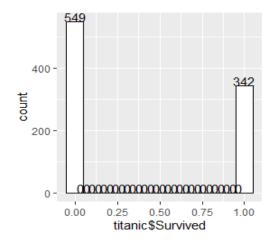
5. Plot every feature from the iris dataset in a histogram.

```
par(mfrow=c(2, 2), mar=c(4, 4, 2, 1))
hist(iris$Sepal.Length, main="Sepal Length")
hist(iris$Petal.Length, main="Petal Length")
hist(iris$Petal.Width, main="Petal Width")
hist(iris$Sepal.Width, main="Sepal Width")
```



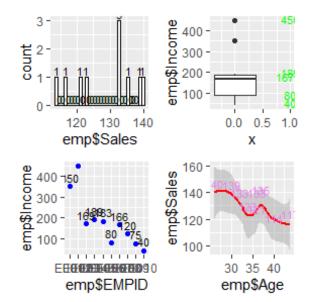
6. Consider a Titanic dataset and compare metric value across different subgroups of the data. Also assume you have a greater number of groups, which visualization method do you prefer over a column chart?

```
titanic<-read.csv("C:/Users/DSL-A-B1/Downloads/Titanic.csv")
ggplot(titanic,aes(x=titanic$Survived))+geom_histogram(binwidth =
0.1,color="black",fill="white")+stat_bin(aes(label = ..count..), geom =
"text", vjust = 0)</pre>
```

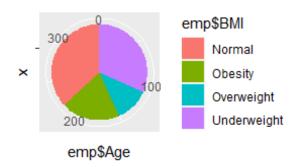


7. For the given dataset and plot different charts.

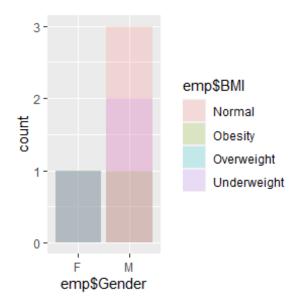
```
emp<-
data.frame(EMPID=c('E001','E002','E003','E004','E005','E006','E007','E008','E
,30,44,36,32,26,32,36),Sales=c(123,114,135,139,117,121,133,140,133,133),BMI=c
('Normal','Overweight','Obesity','Underweight','Underweight','Normal','Obesit
y','Normal','Normal','Underweight'),Income=c(350,450,169,189,183,80,166,120,7
5,40))
head(emp,3)
    EMPID Gender Age Sales
                                 BMI Income
               M 34
                                        350
## 1 E001
                      123
                              Normal
## 2 E002
               F 40
                      114 Overweight
                                        450
## 3 E003
               F 37
                      135
                             Obesity 0
                                        169
library(gridExtra)
ggplot(emp,aes(x=emp$Sales))+geom_histogram(color="black",fill="white")+stat_
bin(geom = "text", aes(label = ..count..), vjust=-0.2, size=3)
f2 <- ggplot(emp,aes(,y=emp$Income))+geom_boxplot()+annotate("text", x = 1, y</pre>
= fivenum(emp$Income), label =
fivenum(emp$Income), vjust=0.3, color="green", size=3)
f3 <- ggplot(emp,aes(x=emp$EMPID,y=emp$Income)) + geom_point(color="blue")</pre>
+geom text(aes(label = Income), vjust = -0.5, hjust = 0.5, color =
"black", size=3)
f4 <- ggplot(emp,aes(x=emp$Age,y=emp$Sales)) +
geom_smooth(color="red")+geom_text(aes(label = Sales), color = "violet",
vjust = -0.5, hjust = 0.5, size=3)
grid.arrange(f1,f2,f3,f4,ncol=2)
```



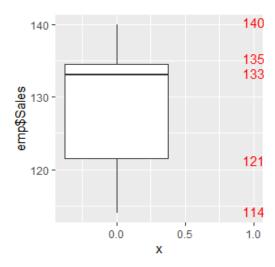
ggplot(emp,aes(x="",y=emp\$Age,fill=emp\$BMI)) + geom_bar(stat="identity",
width=1) + coord_polar("y", start=0)



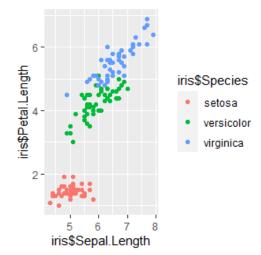
```
ggplot(emp,aes(x=emp$Gender,fill=emp$BMI)) + geom_bar(position="identity",
alpha=1/5)
```



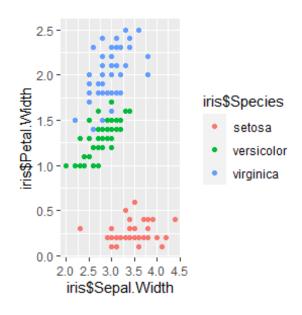
8. Consider the above dataset and draw the boxplot for the statistical data based on the minimum, first quartile, median, third quartile and maximum. ggplot(emp,aes(y=emp\$Sales))+geom_boxplot()+annotate("text", x = 1, y = fivenum(emp\$Sales), label = fivenum(emp\$Sales), vjust=0.3, color="red")



9. Make a scatterplot for the features in the Iris dataset.
ggplot(iris,aes(x=iris\$Sepal.Length,y=iris\$Petal.Length,color=iris\$Species))
+ geom point()

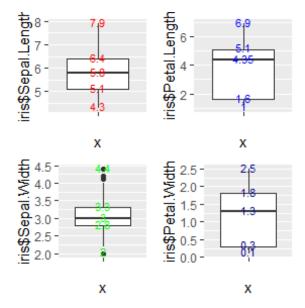


ggplot(iris,aes(x=iris\$Sepal.Width,y=iris\$Petal.Width,color=iris\$Species)) +
geom_point()



10. Look at an individual feature through a boxplot.

```
p1 <- ggplot(iris,mapping = aes(x="",y=iris$Sepal.Length)) +
geom_boxplot()+annotate("text", x = 1, y = fivenum(iris$Sepal.Length), label
= fivenum(iris$Sepal.Length),vjust=0.3,color="red",size=3)
p2 <- ggplot(iris,mapping = aes(x="",y=iris$Petal.Length)) +
geom_boxplot()+annotate("text", x = 1, y = fivenum(iris$Petal.Length), label
= fivenum(iris$Petal.Length),vjust=0.3,color="blue",size=3)
p3 <- ggplot(iris,mapping = aes(x="",y=iris$Sepal.Width)) +
geom_boxplot()+annotate("text", x = 1, y = fivenum(iris$Sepal.Width), label =
fivenum(iris$Sepal.Width),vjust=0.3,color="green",size=3)
p4 <- ggplot(iris,mapping = aes(x="",y=iris$Petal.Width)) +
geom_boxplot()+annotate("text", x = 1, y = fivenum(iris$Petal.Width), label =
fivenum(iris$Petal.Width),vjust=0.3,color="darkblue",size=3)
grid.arrange(p1,p2,p3,p4,ncol = 2)</pre>
```



Exercise - 3

1. Create a synthetic dataset of 50 entries (randomly generated) with the following fields and store it in a file

```
stud <-
data.frame(StudentId=sample(1:50,50,replace=FALSE),Dept=sample(c('CSE','ECE',
'EEE','IT','Civil','Mech'),50,replace=TRUE),Year1=sample(125:130,50,replace=T
RUE), Year2=sample(135:140,50, replace=TRUE), Year3=sample(145:150,50, replace=TR
UE), Year4=sample(155:160,50, replace=TRUE))
head(stud)
##
     StudentId Dept Year1 Year2 Year3 Year4
## 1
            47
                 CSE
                       125
                              140
                                    149
                                          158
## 2
            30
                CSE
                       127
                              140
                                    147
                                          156
## 3
            16
                EEE
                       125
                              138
                                    148
                                          156
write_delim(stud, "Student.txt", delim = "\t")
studtab <- read_delim("C:/Users/DSL-A-</pre>
B1/Downloads/RLAB/Student.txt",delim="\t")
head(studtab)
## # A tibble: 6 × 6
     StudentId Dept Year1 Year2 Year3 Year4
##
         <dbl> <chr> <dbl> <dbl> <dbl> <dbl> <dbl> <
## 1
            46 Mech
                        129
                                     148
                                           160
                               137
            21 ECE
                        126
                               137
                                     146
                                           159
## 2
## 3
             1 IT
                        126
                               138
                                     148
                                           156
```

2. Create a new variable avg_height for each students and compute its value

```
studtab %>% group by(StudentId) %>%
summarize(Avg_Height=(Year1+Year2+Year3+Year4)/4)
## # A tibble: 50 × 2
     StudentId Avg Height
##
##
         <dbl>
                   <dbl>
## 1
             1
                     142
             2
                     142.
## 2
## 3
             3
                     144.
## 4
                     142
```

3. Compute dept wise count of students

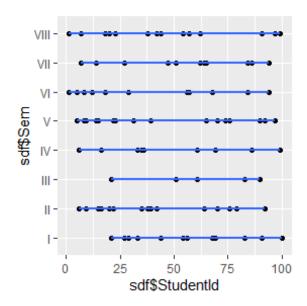
Exercise - 4

1. Create a synthetic dataset of 100 entries (randomly generated) with the following fields and store it in a .csv file:

```
stud1 <-
data.frame(StudentId=sample(1:100,50,replace=FALSE),Dept=sample(factor(c('CSE
,'ECE','EEE','IT','Civil','Mech')),100,replace=TRUE),Sem=sample(factor(c('I'
,'II','III','IV','V','VI','VII','VIII')),100,replace=TRUE),GPA=sample(seq(5,1)
0,0.1),100,replace=TRUE))
head(stud1)
##
    StudentId Dept Sem GPA
## 1
         44 EEE I 5.7
           29 EEE VI 9.1
## 2
## 3
           56 CSE I 8.9
write.csv(stud1, "Student.csv")
sdf<-read.csv("C:/Users/DSL-A-B1/Downloads/Student.csv")</pre>
head(sdf)
##
    X StudentId Dept Sem GPA
## 1 1
             44 EEE I 5.7
## 2 2
             29 FEE VI 9.1
```

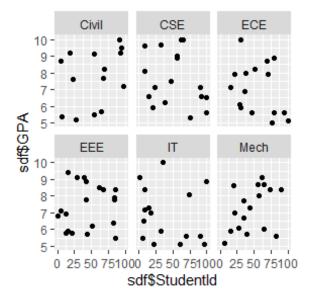
2. Scatterplot with smooth curve for every semester with different pattern in batch-wise mentioned

ggplot(sdf,aes(x=sdf\$StudentId,y=sdf\$Sem))+geom point()+geom smooth()



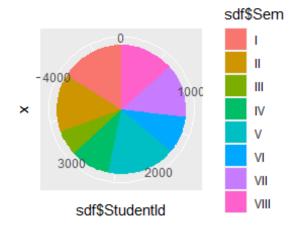
3. Draw subsets of scatterplot to plot GPA for each department.

ggplot(sdf)+geom_point(aes(x=sdf\$StudentId,y=sdf\$GPA))+facet_wrap(~
sdf\$Dept,nrow = 2)



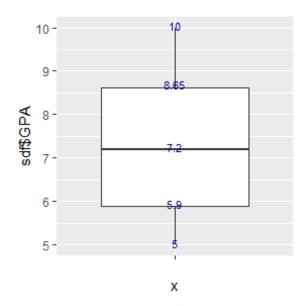
4. Bar chart specification mentioned in batch-wise

```
ggplot(sdf,aes(x="",y=sdf$StudentId,fill=sdf$Sem)) +
geom_bar(stat="identity", width=1) + coord_polar("y", start=0)
```



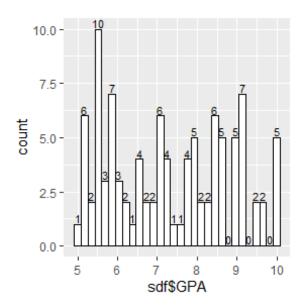
5. Identify the outlier's using boxplot.

```
ggplot(sdf,mapping = aes(x="",y=sdf$GPA)) + geom_boxplot()+annotate("text", x
= 1, y = fivenum(sdf$GPA), label =
fivenum(sdf$GPA),vjust=0.3,color="darkblue",size=3)
```



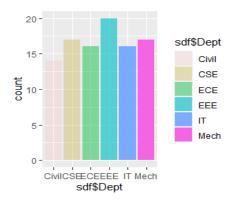
6. Draw histograms for count of GPA with different bin width & zoom to see any one region

```
ggplot(sdf,aes(x=sdf$GPA,bin_width=sdf$GPA))+geom_histogram(color="black",fil
l="white")+stat_bin(geom = "text", aes(label = ..count..),vjust=-0.2,size=3)
```



7. Transparency for every Department

ggplot(sdf,aes(x=sdf\$Dept,fill=sdf\$Dept)) + geom_bar(position="identity",
aes(alpha=sdf\$Dept))



8. Number of students in each department with different colours each department

ggplot(sdf,aes(x=sdf\$Dept,fill=sdf\$Dept)) + geom_bar(position="identity",
aes(alpha=sdf\$Dept))

