DWDM R PROGRAMMING-PRACTICALS

1.List of Programs:

1The intervals and corresponding frequencies are as follows. age frequency

1-5. 200

5-15 450

15-20 300

20-50 1500

50-80 700

80-110 44

Compute an approximate median value for the data

CODING-

```
class_interval<-c("1-5","5-15","15-20","20-50","50-80","80-110")
data<-c(200,450,300,1500,700,44)
data.frame(class_interval,data)
median(data)
```

OUTPUT-

> data.frame(class_interval,data)

class_interval data

- 1 1-5 200
- 2 5-15 450
- 3 15-20 300
- 4 20-50 1500
- 5 50-80 700
- 6 80-110 44

> median(data)

[1] 375

- 2. Suppose that the data for analysis includes the attribute age. The age values for the data tuples are (in increasing order) 13, 15, 16, 16, 19, 20, 20, 21, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 36, 40, 45, 46, 52, 70.
- (a) What is the mean of the data? What is the median?
- (b) What is the mode of the data? Comment on the data's modality (i.e., bimodal, trimodal, etc.).
- (c) What is the midrange of the data?
- (d) Can you find (roughly) the first quartile (Q1) and the third quartile (Q3) of the data?

```
Coding:
#2a
x<-c(13,15,16,16,19,20,20,21,22,22,25,25,25,25,30,33,33,35,35,35,35,36,40,45,46,52,70)
#mean
mean(x)
#median
median(x)
output:
mean(x)
[1] 29.96296
> #median
> median(x)
[1] 25
CODING FOR 2b-
#2b
#mode
MultipleModes <- function(x) {
uniqx <- unique(x)
freq_table <- tabulate(match(x, uniqx))</pre>
 modes <- uniqx[freq_table == max(freq_table)]
modes
}
```

```
age_values <- c(13, 15, 16, 16, 19, 20, 20, 21, 22, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 35, 36, 40,
45, 46, 52, 70)
multiple_modes <- MultipleModes(age_values)</pre>
print(multiple_modes)
output:
25 35
CODING FOR 2c-
#midrange
c) age_values <- c(13, 15, 16, 16, 19, 20, 20, 21, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 35, 36,
40, 45, 46, 52, 70)
median(age_values)
OUTPUT-
25
CODING FOR 2d-
d) #quartile
age_values <- c(13, 15, 16, 16, 19, 20, 20, 21, 22, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 35, 36, 40,
45, 46, 52, 70)
quantile(age_values)
output: 0% 25% 50% 75% 100%
```

3.Data Preprocessing :Reduction and Transformation

Use the two methods below to normalize the following group of data: 200, 300, 400, 600, 1000 (a) min-max normalization by setting min = 0 and max = 1 (b) z-score normalization

Coding:

```
#3a
```

```
data <- c(200, 300, 400, 600, 1000)
min<-min(data)
max<-max(data)
```

13.0 20.5 25.0 35.0 70.0

```
for (i in data)
{
result1=i-min
result2=max-min
result3=result1/result2
print(result3)
}
OUTPUT:
[1] 0
[1] 0.125
[1] 0.25
[1] 0.5
[1] 1
#3b
data <- c(200, 300, 400, 600, 1000)
mean1<-mean(data)
deviation<-sd(data)
for (i in data)
{
result1=i-mean1
result2=result1/deviation
print(result2)
}
OUTPUT:
[1] -0.9486833
[1] -0.6324555
[1] -0.3162278
[1] 0.3162278
[1] 1.581139
4.Data:11,13,13,15,15,16,19,20,20,20,21,21,22,23,24,30,40,45,45,45,71,
```

a) Smoothing by bin mean

b) Smoothing by bin median

c) Smoothing by bin boundaries

CODING-

```
#binning
data <- c(11, 13, 13, 15, 15, 16, 19, 20, 20, 20, 21, 21, 22, 23, 24, 30, 40, 45, 45, 45, 71, 72, 73, 75)
range=6
bin1=c()
bin2=c()
bin3=c()
bin4=c()
for(i in data[1:range]){
bin1=append(bin1,i)
}
range1=range+1
range2=range*2
for(j in data[range1:range2])
{
bin2=append(bin2,j)
}
range3=range2+1
range4=range*3
for(k in data[range3:range4])
{
bin3=append(bin3,k)
}
range5=range4+1
range6=range*4
```

```
for(I in data[range5:range6]){
bin4=append(bin4,l)
}
#4a
mean(bin1)
mean(bin2)
mean(bin3)
mean(bin4)
#4b
median(bin1)
median(bin2)
median(bin3)
median(bin4)
OUTPUT:
#4a
> mean(bin1)
[1] 13.83333
> mean(bin2)
[1] 20.16667
> mean(bin3)
[1] 30.66667
> mean(bin4)
[1] 63.5
#4b
> median(bin1)
[1] 14
> median(bin2)
[1] 20
> median(bin3)
```

[1] 27

[1] 71.5

5) 5. Suppose that a hospital tested the age and body fat data for 18 randomly selected adults with the following results:

age	23	23	27	27	39	41	47	49	50
%fat	9.5	26.5	7.8	17.8	31.4	25.9	27.4	27.2	31.2
age	52	54	54	56	57	58	58	60	61

CODING-

#box plot

boxplot(df)

#scatter plot

plot(df)

df<-data.frame(age,body_fat_percent)</pre>

```
age <- c(23,23,27,27,39,41,47,49,50,52,54,54,56,57,58,58,60,61)

body_fat_percent <-
c(9.5,26.5,7.8,17.8,31.4,25.9,27.4,27.2,31.2,34.6,42.5,28.8,33.4,30.2,34.1,32.9,41.2,35.7)

#5.a

mean(age)

mean(body_fat_percent)

median(age)

median(body_fat_percent)

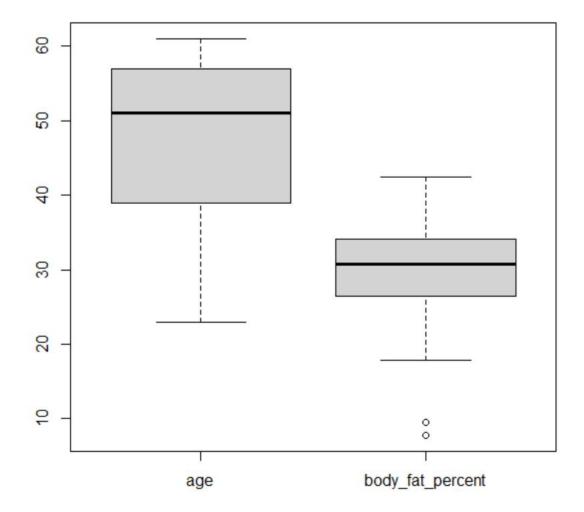
sd(age)

sd(body_fat_percent)

#5.b

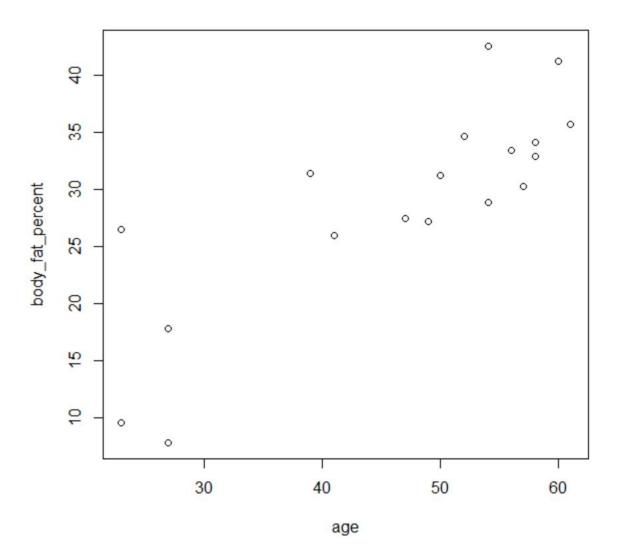
#create dataframe
```

```
#qq plot
qqnorm(age)
qqline(age)
qqnorm(body_fat_percent)
qqline(body_fat_percent)
OUTPUT-
#5a
> mean(age)
[1] 46.44444
> mean(body_fat_percent)
[1] 28.78333
> median(age)
[1] 51
> median(body_fat_percent)
[1] 30.7
> sd(age)
[1] 13.21862
> sd(body_fat_percent)
[1] 9.254395
#5.b
BOXPLOT-
```



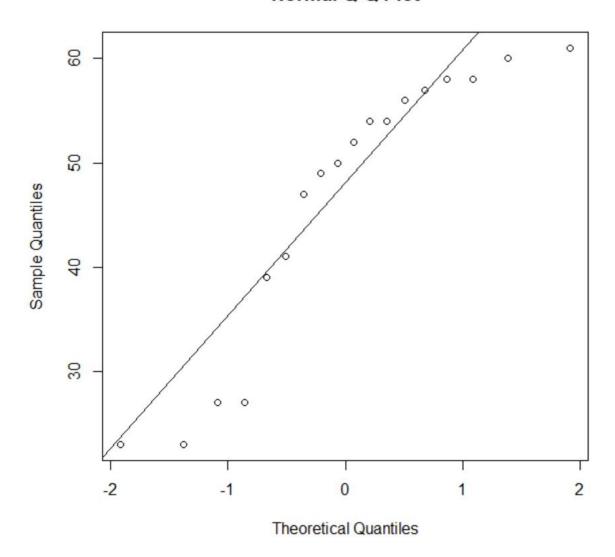
SCATTER PLOT-

#5c



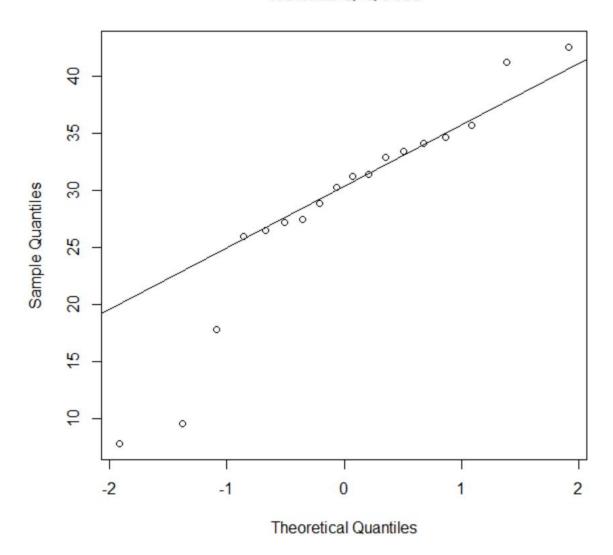
QQ PLOT FOR AGE-

Normal Q-Q Plot



QQPLOT FOR BODY FAT PERCENT-

Normal Q-Q Plot



6. Suppose that a hospital tested the age and body fat data for 18 randomly selected adults with the following results:

- (i) Use min-max normalization to transform the value 35 for age onto the range [0.0, 1.0].
- (ii) Use z-score normalization to transform the value 35 for age, where the standard deviation of age is 12.94 years.
- (iii) Use normalization by decimal scaling to transform the value 35 for age. Perform the above functions using R-tool

CODING-

```
age <- c(23,23,27,27,39,41,47,49,50,52,54,54,56,57,58,58,60,61)

new_age<-c()

for(i in age){
```

```
if(i<=35){
  new_age=append(new_age,i)
}
}
print(new_age)
#6a
#min max normalization
min<-min(new_age)
max<-max(new_age)
for (i in new_age)
{
result1=i-min
result2=max-min
result3=result1/result2
print(result3)
}
#6b
#z score normalization
mean1<-mean(new_age)</pre>
for (i in new_age)
{
result1=i-mean1
result2=result1/12.94
print(result2)
}
#6c
#decimal scaling
n=200
j=nchar(y)
scaling=n/10^j
print(scaling)
```

6.a MIN MAX NORMALIZATION [1] 0 [1] 0 [1] 1 [1] 1 [1] 1 6.b Z SCORE NORMALIZATION [1] -0.8660254 [1] -0.8660254 [1] 0.8660254 [1] 0.8660254 [1] 0.8660254

OUTPUT-

7. The following values are the number of pencils available in the different boxes. Create a vector and find out the mean, median and mode values of set of pencils in the given data.

Box1 Box2 Box3 Box4 Box5 Box6 Box7 Box8 Box9 Box 10

9 25 23 12 11 6 7 8 9 10

CODING-

[1] 0.2

```
box_no=c("box1","box2","box3","box4","box5","box6","box7","box8","box9","box10")
pencil=c(9,25,23,12,11,6,7,8,9,10)
df<-data.frame(box_no,pencil)
#dataframe
print(df)
#mean
mean(pencil)</pre>
```

```
#median
median(pencil)
#mode
mode=names(which.max(table(pencil)))
print(mode)
OUTPUT-
> data.frame(box_NO,pencil)
 box_NO pencil
1 box1 9
2 box2 25
3 box3 23
4 box4 12
5 box5 11
6 box6 6
7 box7
        7
8 box8 8
9 box9 9
10 box10 10
> mean(pencil)
[1] 12
> median(pencil)
[1] 9.5
> print(mode)
[1] "9"
```

8. the following table would be plotted as (x,y) points, with the first column being the x values as number of mobile phones sold and the second column being the y values as money. To use the scatter plot for how many mobile phones sold.

x :4 1 5 7 10 2 50 25 90 36

y:12 5 13 19 31 7 153 72 275 110

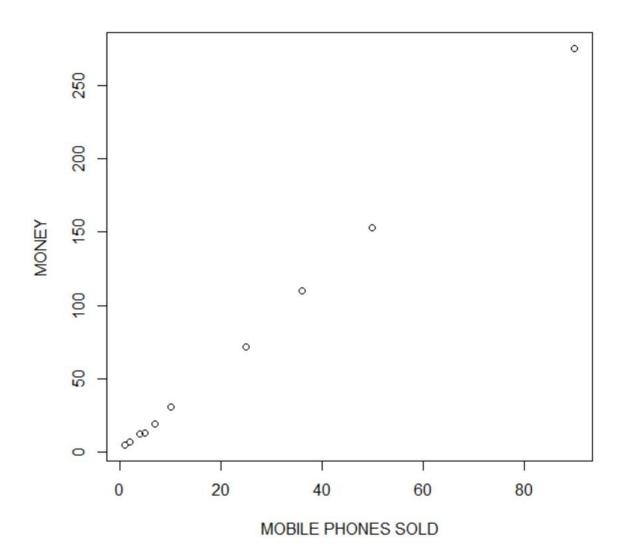
CODING-

x<-c(4, 1, 5, 7, 10, 2, 50, 25, 90, 36)

y<-c(12,5, 13, 19, 31, 7, 153, 72, 275, 110)

plot(x,y,xlab='MOBILE PHONES SOLD',ylab='MONEY')

OUTPUT-



9. Implement of the R script using marks scored by a student in his model exam has been sorted as follows: 55, 60, 71, 63, 55, 65, 50, 55,58,59,61,63,65,67,71,72,75. Partition them into three bins by each of the following methods. Plot the data points using histogram.

(a) equal-frequency (equi-depth) partitioning (b) equal-width partitioning

CODING-

```
marks<-c(55, 60, 71, 63, 55, 65, 50, 55, 58, 59, 61, 63, 65, 67, 71, 72, 75)
binning1=c()
binning2=c()
binning3=c()
class=6
#binning partition
for(a in marks[1:class]){
 binning1=append(binning1,a)
}
range1=range+1
range2=range*2
for(b in marks[range1:range2])
{
 binning2=append(binning2,b)
}
range3=range2+1
range4=range*3
for(c in marks[range3:range4])
{
```

```
binning3=append(binning3,c)
}
print(binning1)
print(binning2)
print(binning3)
#histogram
hist(binning1)
hist(binning2)
hist(binning3)
#9a
#equal-frequency
freq=length(marks)/range
print(freq)
#9b
#equal-width
min<-min(marks)
max<-max(marks)
result<-max-min
width<-result/range
cat("width is",width)
bin1=width+min
print(bin1)
bin2=2*width+min
print(bin2)
```

bin3=3*width+min
print(bin3)
OUTPUT-

> print(binning1)

[1] 55 60 71 63 55 65

> print(binning2)

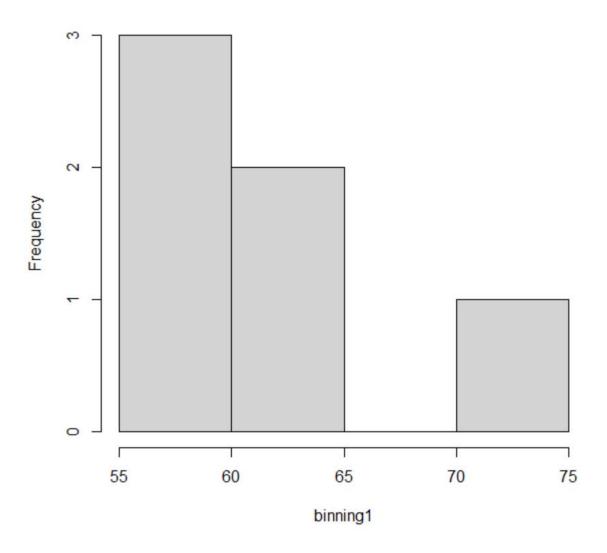
[1] 50 55 58 59 61 63

> print(binning3)

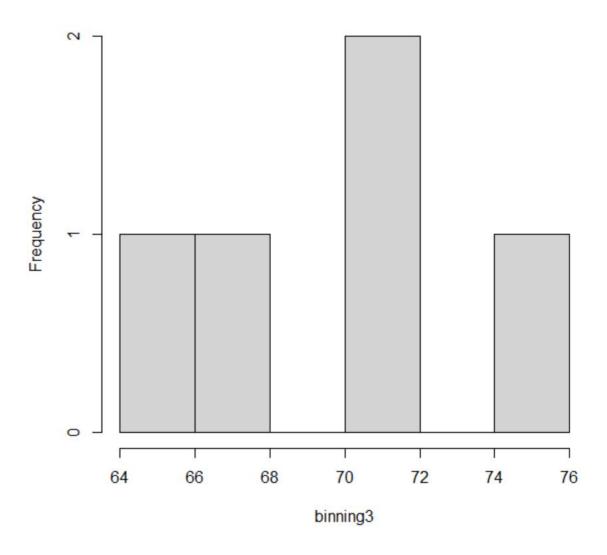
[1] 65 67 71 72 75 NA

HISTOGRAM-

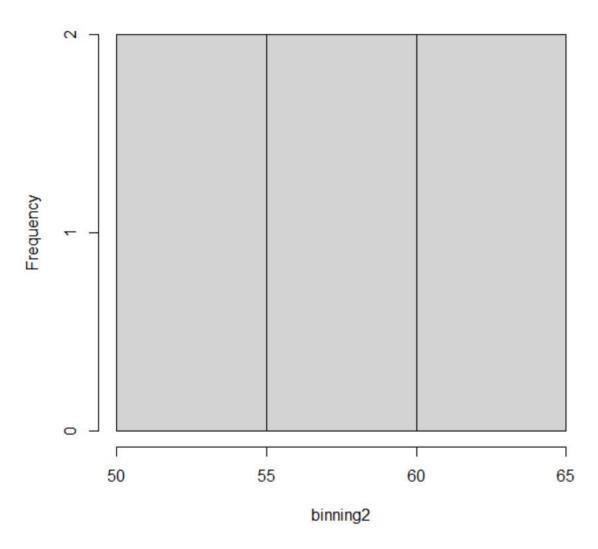
Histogram of binning1



Histogram of binning3



Histogram of binning2



#9a

#equal frequency

> print(freq)

[1] 2.833333

#9b

#equal width

width is 4.166667> bin1=width+min

> print(bin1)

[1] 54.16667

```
> bin2=2*width+min
> print(bin2)
[1] 58.33333
> bin3=3*width+min
> print(bin3)
[1] 62.5
10. Suppose that the speed car is mentioned in different driving
style.
Regular 78.3 81.8 82 74.2 83.4 84.5 82.9 77.5 80.9 70.6 Speed
Calculate the Inter quantile and standard deviation of the given data.
CODING-
speed<-c(78.3,81.8,82,74.2,83.4,84.5,82.9,77.5,80.9,70.6)
#interquartile
IQR(speed)
#standard deviation
sd(speed)
OUTPUT-
> IQR(speed)
[1] 4.975
> sd(speed)
[1] 4.445835
```

11. Suppose that the data for analysis includes the attribute age. The age values for the data tuples are (in increasing order) 13, 15, 16, 16, 19, 20, 20, 21, 22, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 35, 36, 40, 45, 46, 52, 70.

Can you find (roughly) the first quartile (Q1) and the third quartile

(Q3) of the data?

CODING-

marks<-c(13,15, 16, 16, 19, 20, 20, 21, 22, 22, 25, 25, 25, 25, 30, 33, 33, 35, 35, 35, 35, 36, 40, 45, 46, 52, 70)

quantile(marks)

OUTPUT-

> quantile(marks)

0% 25% 50% 75% 100%

13.0 20.5 25.0 35.0 70.0