## R PROGRAMMING

Ragu.v REG NO: 191921031

1. 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. What is the median?

Solution

> age <- 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)

> median age <- median (age)

> print (median age)

Output

[1] 25

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, 25, 30, 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?

Solution

```
> age <- 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)
```

> g1 <- quantile (age, 0.25)

> g3 <- quantile (age, 0.75)

> print (g1)

25%

20.5

> print (g3)

75%

35

3. Load iris Dataset which is inbuilt in R .explore the dataset in terms of dimension and summary statistics

```
Solution
> library (datasets)
> data (iris)
> head (iris)
Sepal. Length Sepal.Width Petal.Width Species
1
5.1
3.5
1.4
2
4.9
3.0
1.4
3
4.7
3.2
1.3
4
4.6
3.1
1.5
5
5.0
3.6
1.4
6
5.4
3.9
1.7
0.2
setosa
0.2
setosa
0.2
setosa
0.2 setosa
0.2
setosa
0.4 setosa
> summary (iris)
Sepal.Length
Min.
:4.300
1st Qu.:5.100
Median :5.800
Mean
:5.843
```

3rd Qu.: 6.400

```
Max.
    :7.900
    Sepal.Width
    Min.:2.000
    1st Qu.:2.800
    Median :3.000
    Mean
    :3.057
    3rd Qu.: 3.300
    Max.
    4.400
    Petal.Length
    Petal.Width
    Min.
    :1.000
    Min.:0.100
    1st Qu.: 1.600
    1st Qu.: 0.300
    Median: 4.350
    Median: 1.300
    Mean
    :3.758
    Mean
    :1.199
    3rd Qu.: 5.100
    3rd Qu.: 1.800
    Max.
    :6.900
    Max.
    :2.500
    Species
    setosa
    :50
    versicolor: 50
    virginica:50
4. Find the categorical column data and convert that to factor form, also find the number of
    rows for each factors in dataset.
    Solution
    > df <- data.frame(
    +name=c("John", "Jane", "Bob", "Sara", "Mike", "Emma", "Tom"),
    gender = c("Male", "Female", "Male", "Female", "Male", "Female", "Male"), + marital status =
   c("Married", "Single", "Married", "Single", "Divorced", "Single", "Widowed")
```

```
age = c(25, 30, 35, 20, 45, 28, 33), city= c("Kolkata", "Delhi", "Chennai", "Hubali", "Mysore",
    "Bombay", "Surat"),
    > dfsgender <- factor (dfsgender)
    > df marital status <- factor (dfSmarital status) > table (dfsgender)
    Female Male
    3
    4
    > table (df@marital status)
    Divorced Married
    1
    2
    Single
    3
    Widowed
    1
5. Find mean of numeric
6. Find mean of numeric data in dataset based on Species group. and plot Bar chart (use ggplot )
    to interpret same
            Species
                            Sepal Length
                                            Sepal Width
                                                            Petal Length
                                                                            Petal Width
                               5.006
            Setosa
                                               3.428
                                                               1.462
                                                                              0.246
            Versicolour
                               5.936
                                               2.770
                                                                4.250
                                                                              1.326
            Virginica
                               6.588
                                               2.974
                                                                5.552
                                                                              2.026
      ١.
            Draw a suitable plot which summaries statistical parameter of Sepal. Width based on
            Species group
            > library(dplyr)
            > library(ggplot2)
```

```
> df <- data.frame( Species c("Setosa", "Versicolour", "Virginia"),
Sepal.Length (5.006, 5.936, 6.500),
Sepal.Width (3.428, 2.77, 2.974), Petal.Length c(1.462, 4.25, 5.552),
Petal.Width c(0.246, 1.326, 2.026)
>df mean of 44
group by (Species)
summarise all (mean)
> ggplot (df_mean, aes(x Species, y Sepal.Width))+
gece bar (stat "identity")
labs (title="Mean Sepal Width based on Species group", x "Species", y "Mean Sepal
Width")
> ggplot (df, aes(x Species, y Sepal.Width)) +
geom boxplot() +
labs (title "Statistical parameter of Sepal Width based on Species group", x "Species",
y "Sepel Width")
> ggplot (df, aes (x Sepal.Width, fill Species))+
geom density (alpha 0.5)+
labs (title "Density plot of Sepal Width", "Sepal Width")
Warning messages:
1: Groups with fewer than two data points have been dropped.
2: Groups with fewer than two data points have been dropped.
3: Groups with fewer than two data points have been dropped.
4: In max(ids, na.rm TRUE):
no non-missing arguments to max: returning -Inf
5: In max(ids, na.rm=TRUE):
```

```
no non-missing arguments to max: returning -Inf
     6: In max(ids, na.zm TRUE):
     no non-missing arguments to max: returning -Inf
     > ggplot (df, nes (x- Sepal.Length, y Petal.Length, color Species)) +
     geom_point()+
     labs (title "Scatterplot of Sepal Length and Petal Length grouped by Species", "Sepal
     Length", "Fetal Length")
11.
     Draw a suitable plot to find the skewness of the data for Sepal. Width and print the
     comment about skewness.
     > library(dplyr)
     > library(ggplot2)
     > df <- data.frame( Species c("Setosa", "Versicolour", "Virginia"),
     Sepal.Length (5.006, 5.936, 6.500),
     Sepal.Width (3.428, 2.77, 2.974), Petal.Length c(1.462, 4.25, 5.552),
     Petal.Width c(0.246, 1.326, 2.026)
     >df mean of 44
     group by (Species)
     summarise all (mean)
     > ggplot (df_mean, aes(x Species, y Sepal.Width))+
     gece bar (stat "identity")
     labs (title="Mean Sepal Width based on Species group", x "Species", y "Mean Sepal
     Width")
     > ggplot (df, aes(x Species, y Sepal.Width)) +
     geom boxplot() +
     labs (title "Statistical parameter of Sepal Width based on Species group", x "Species",
     y "Sepel Width")
```

```
> ggplot (df, aes (x Sepal.Width, fill Species))+
        geom density (alpha 0.5)+
       labs (title "Density plot of Sepal Width", "Sepal Width")
        Warning messages:
        1: Groups with fewer than two data points have been dropped.
        2: Groups with fewer than two data points have been dropped.
        3: Groups with fewer than two data points have been dropped.
       4: In max(ids, na.rm TRUE):
        no non-missing arguments to max: returning -Inf
        5: In max(ids, na.rm=TRUE):
       no non-missing arguments to max: returning -Inf
        6: In max(ids, na.zm TRUE):
       no non-missing arguments to max: returning -Inf
       > ggplot (df, nes (x- Sepal.Length, y Petal.Length, color Species)) +
       geom_point()+
       labs (title "Scatterplot of Sepal Length and Petal Length grouped by Species", "Sepal
       Length", "Fetal Length")
 III.
       Draw ggplot2 scatterplot showing the variables Sepal.Length and Petal.Length
        grouped by the three-level factor "Species".
               > library(datasets) > data (izia)
> head(iris) Sepal.Length
Sepal.Width 3.5
Petal.Length
Petal.Width
Species 0.2 setosa
```

5.1 4.9

1.4

1.4

0.2

setosa

3.0

3.2 3.1

3

4.7

1.3

0.2

setosa

4.6

1.5

0.2

setosa

5.0

3.6

1.4

0.2

setosa

5.4

3.9

1.7

setosa

Petal.Width Min.: 0.100

Species

> summary(iris) Sepal.Length

Min. 14.300 1st Qu.:5.100

Median:5.800 Mean

setosa:50

Sepal.Width

Min. :2.000 1st

Qu.:2.800

Median: 3.000: 3.057

Mean

Petal.Length

Min. 1st Qu.:1.600

:1.000

Median :4.350 Mean

1st Qu.:0.300

Median :1.300

versicolor:50 virginica :50

:5.843

:3.758

Mean

1.199

3rd Qu.:1.800

3rd Qu.:6.400

Max.

:7.900

3rd Qu.:3.300

Max. :4.400

3rd Qu.:5.100

Max.

:6.900

Max.

:2.500