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, 22, 25, 25, 25, 25, 30, 33,

33, 35, 35, 35, 35, 36, 40, 45, 46, 52, 70. What is the median?

code :

```
ages <- 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)
sorted_ages <- sort(ages)
median_age <- median(sorted_ages)
```

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, 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?

code:

```
ages <- 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)
Q1 <- quantile(ages, 0.25)
Q3 <- quantile(ages, 0.75)
> Q1
 25%
 20
```

```
> Q3
 75%
 35
```

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

summary statistics (2M)

code:

```
data(iris)
dim(iris)
[1] 150   5
head(iris)
summary(iris)
```

```
 Sepal.Length    Sepal.Width     Petal.Length    Petal.Width        Species
 Min.   :4.300   Min.   :2.000   Min.   :1.000   Min.   :0.100   setosa    :50
 1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300   versicolor:50
 Median :5.800   Median :3.000   Median :4.350   Median :1.300   virginica :50
 Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199
 3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
 Max.   :7.900   Max.   :4.400   Max.   :6.900   Max.   :2.500
```

4.Find the categorical column data and convert that to factor form, also find the number of rows for each factors in dataset. (2)

```
iris$Species <- as.factor(iris$Species)
table(iris$Species)
setosa versicolor  virginica
   50      50       50
```

5. Find mean of numeric data in dataset based on Species group. and plot Bar chart (use ggplot) to interpret same (8m)

```
library(dplyr)
library(ggplot2)

dataset <- read.csv("my_dataset.csv")
```

```r
species_means <- dataset %>%
  group_by(Species) %>%
  summarize(mean = mean(NumericData))
ggplot(species_means, aes(x = Species, y = mean)) +
  geom_bar(stat = "identity") +
  labs(title = "Mean Numeric Data by Species",
       x = "Species",
       y = "Mean Numeric Data")


library(ggplot2)
data(iris)
```

6.Draw a suitable plot which summaries statistical parameter of Sepal.Width based on Species group(6m)

```r
ggplot(iris, aes(x = Species, y = Sepal.Width, fill = Species)) +
  geom_boxplot() +
  labs(x = "Species", y = "Sepal Width", title = "Box plot of Sepal Width by Species")
```

7. Draw a suitable plot to find the skewness of the data for Sepal.Width and print the comment about skewness. (6m)

```r
library(ggplot2)


data(iris)


ggplot(iris, aes(x = Sepal.Width)) +
  geom_histogram(aes(y = ..density..), bins = 20, color = "black
```

8.Draw ggplot2 scatterplot showing the variables Sepal.Length and Petal.Length grouped by the three-level factor "Species". (6m)

```
library(ggplot2)

data(iris)

ggplot(iris, aes(x = Sepal.Length, y = Petal.Length, color = Species)) +

 geom_point() +

 labs(x = "Sepal Length", y = "Petal Length", color = "Species")
```