```
In [1]: ## Question 2
library(ggplot2)

data(iris)

head(iris)

x = iris[,-5]
y = iris$Species

kc <- kmeans(x,3)
kc

table(y,kc$cluster)

## Sepal Width vs Sepal Length
ggplot(x,aes(x = Sepal.Length, y= Sepal.Width,color = as.factor(kc$cluster))) + geom_point()

## Petal Width vs Petal Length
ggplot(x,aes(x = Petal.Length, y= Petal.Width,color = as.factor(kc$cluster))) + geom_point()</pre>
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

K-means clustering with 3 clusters of sizes 38, 50, 62

```
Cluster means:
```

```
Sepal.LengthSepal.WidthPetal.LengthPetal.Width16.8500003.0736845.7421052.07105325.0060003.4280001.4620000.24600035.9016132.7483874.3935481.433871
```

Clustering vector:

Within cluster sum of squares by cluster: [1] 23.87947 15.15100 39.82097

(between SS / total SS = 88.4 %)

Available components:

- [1] "cluster" "centers" "totss" "withinss" "tot
 .withinss"
 [6] "betweenss" "size" "iter" "ifault"
- y 1 2 3 setosa 0 50 0 versicolor 2 0 48 virginica 36 0 14



