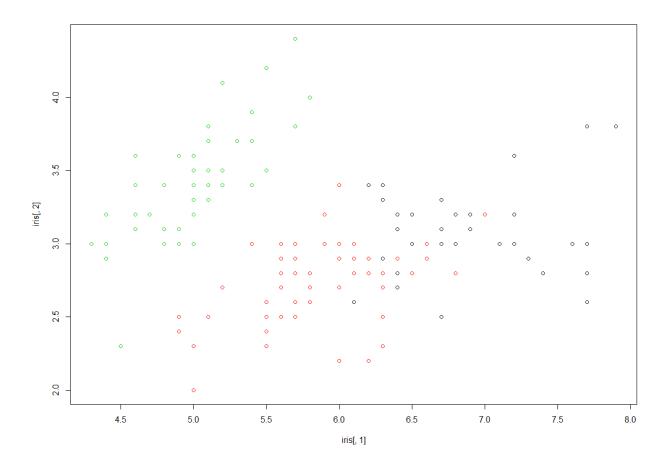
Unsupervised Learning - K-mean clustering

Code and Output

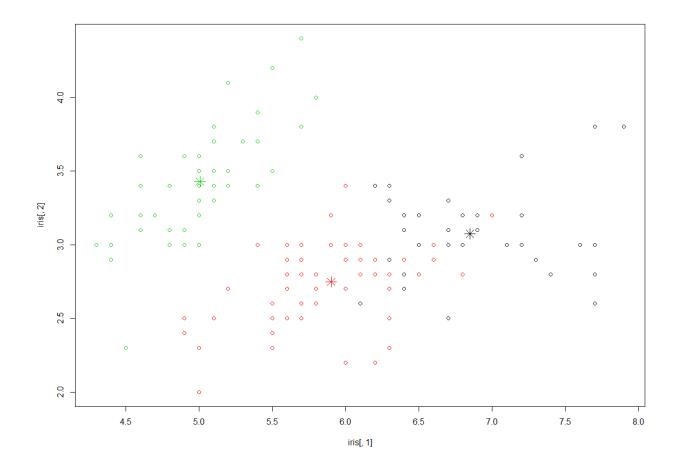
```
> rm(list=ls()) #Clear the environment
> setwd("E:/Dropbox/RU DataScience/MSDS650/Week7/Assignment") #Set working directory for the assignment
> getwd() #Check working directory
[1] "E:/Dropbox/RU DataScience/MSDS650/Week7/Assignment"
> data(iris) #include iris dataset
> head(iris) #view the first few rows of data
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
                                   1.4
1
           5.1
                      3.5
                                              0.2 setosa
2
           4.9
                        3.0
                                     1.4
                                                 0.2 setosa
3
           4.7
                       3.2
                                    1.3
                                                 0.2 setosa
4
           4.6
                       3.1
                                    1.5
                                                 0.2 setosa
           5.0
                       3.6
                                     1.4
                                                 0.2 setosa
5
                       3.9
6
           5.4
                                   1.7
                                                 0.4 setosa
> set.seed(42) #set seed for the algorithm to ensure reproducible results
> km <-kmeans(iris[,1:4], 3, nstart=25) #chose 3 clusters as there are three species in the dataset
> km #output results K-means clustering with 3 clusters of sizes 38, 62, 50 \,
cluster means:
 Sepal.Length Sepal.width Petal.Length Petal.width
6.850000 3.073684 5.742105 2.071053
5.901613 2.748387 4.393548 1.433871
    5.006000
            3.428000
clustering vector:
within cluster sum of squares by cluster:
[1] 23.87947 39.82097 15.15100
(between_SS / total_SS = 88.4 %)
Available components:
[1] "cluster"
             "centers"
                         "totss"
                                               "tot.withinss" "betweenss"
                                                                     "size"
                                                                                 "iter"
                                                                                            "ifault"
                                    "withinss"
> rm(list=ls()) #Clear the environment
> setwd("E:/Dropbox/RU DataScience/MSDS650/Week7/Assignment") #Set working directory for the assignment
> getwd() #Check working directory
 [1] "E:/Dropbox/RU DataScience/MSDS650/Week7/Assignment"
> data(iris) #include iris dataset
> head(iris) #view the first few rows of data
   Sepal.Length Sepal.width Petal.Length Petal.width Species
 1
                                                  1.4
                                                                   0.2
                5.1
                                3.5
                4.9
                                3.0
                                                  1.4
                                                                   0.2
                                                                         setosa
 3
                4.7
                                3.2
                                                  1.3
                                                                   0.2
                                                                          setosa
 4
                                                                   0.2
                4.6
                                3.1
                                                   1.5
                                                                          setosa
 5
                5.0
                                3.6
                                                  1.4
                                                                   0.2
                                                                          setosa
                5.4
                                3.9
                                                                   0.4
 6
                                                  1.7
                                                                         setosa
   set.seed(42) #set seed for the algorithm to ensure reproducible results
>
> km <-kmeans(iris[,1:4], 3, nstart=25) #chose 3 clusters as there are three
species in the dataset
> km #output results
K-means clustering with 3 clusters of sizes 38, 62, 50
Cluster means:
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width
      6.850000
                  3.073684
                               5.742105
                                           2.071053
23
      5.901613
                  2.748387
                               4.393548
                                           1.433871
                  3.428000
                               1.462000
      5.006000
                                           0.246000
Within cluster sum of squares by cluster: [1] 23.87947 39.82097 15.15100
 (between_SS / total_SS = 88.4 %)
Available components:
[1] "cluster"
s" "betweenss"
                   "centers"
                                  "totss"
                                                "withinss"
                                                               "tot.withins
                                                "ifault"
                  "size"
                                 "iter"
> #Compare the clusters with the species and plot results
> table(km$cluster, iris$Species)
    setosa versicolor virginica
 1
        0
                   2
                            36
 2
        0
                  48
                            14
 3
       50
                   0
                             0
> table(km$cluster, iris$Species)
    setosa versicolor virginica
  1
         0
                    2
                             36
  2
         0
                             14
                   48
   3
        50
                    0
                              0
>
> #Plot the results
> plot(iris[,1], iris[,2], col=km$cluster) #Plot cluster centers by Sepal Le ngth by Sepal width
>
```



```
> #add center points
> points(km$centers[,c(1,2)], col=1:3, pch=8, cex=2)
```

>



```
#Plot Petal length and width
> plot(iris[,3], iris[,4], col=km$cluster)
> #add center points
> points(km$centers[,c(3,4)], col=1:3, pch=8, cex=2)
```

