

K-Means Clustering

Mr. Gulati

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```
library("NbClust")
library("ggplot2")
library("factoextra")
```

```
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
```

```
library("dplyr")
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
##   filter, lag
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##   intersect, setdiff, setequal, union
```

Reading the data

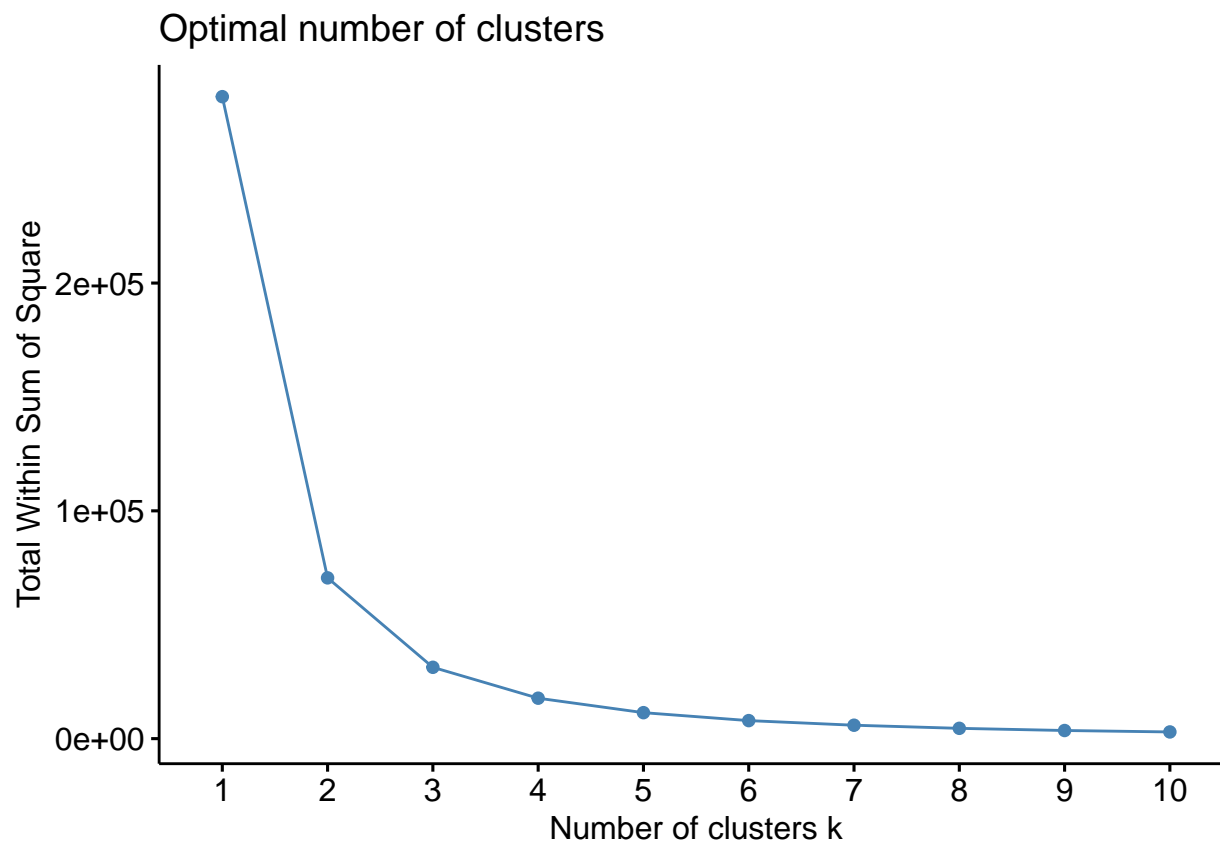
```
dataset <- read.csv("D:/Internship/Task 2/Iris.csv", sep = ",", header = TRUE)
dataset_df <- data.frame(dataset)
dataset_df <- na.omit(dataset_df)
head(dataset_df)
```

```
##   Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm   Species
## 1  1           5.1           3.5           1.4           0.2 Iris-setosa
## 2  2           4.9           3.0           1.4           0.2 Iris-setosa
## 3  3           4.7           3.2           1.3           0.2 Iris-setosa
## 4  4           4.6           3.1           1.5           0.2 Iris-setosa
## 5  5           5.0           3.6           1.4           0.2 Iris-setosa
## 6  6           5.4           3.9           1.7           0.4 Iris-setosa
```

Finding optimum number of clusters

```
set.seed(100)

fviz_nbclust(dataset_df[,c(1,2,3,4)], kmeans, method = "wss")
```



The results suggests the bend appears at $k=3$

Applying kmeans

```
model <- kmeans(dataset_df[,c(1,2,3,4)], 3, nstart = 25)
model
```

```
## K-means clustering with 3 clusters of sizes 50, 50, 50
```

```
##
```

```
## Cluster means:
```

```
##      Id SepalLengthCm SepalWidthCm PetalLengthCm
```

```
## 1  25.5          5.006          3.418          1.464
```

```
## 2  75.5          5.936          2.770          4.260
```

```
## 3 125.5          6.588          2.974          5.552
```

```
##
```

```
## Clustering vector:
```

```
##  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20
```

```
##  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1
```

```
## 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
```

```
##  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1
```

```
## 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60
```

```
##  1  1  1  1  1  1  1  1  1  1  2  2  2  2  2  2  2  2  2  2
```

```
## 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80
```

```
##  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2
```

```
## 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100
```

```
##  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2
```

```
## 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120
```

```
## 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
## 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140
## 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
## 141 142 143 144 145 146 147 148 149 150
## 3 3 3 3 3 3 3 3 3 3
##
## Within cluster sum of squares by cluster:
## [1] 10427.18 10441.20 10452.33
## (between_SS / total_SS = 88.9 %)
##
## Available components:
##
## [1] "cluster" "centers" "totss" "withinss" "tot.withinss"
## [6] "betweenss" "size" "iter" "ifault"
```

Visualising Clusters

```
iris_clustered <- data.frame(dataset_df, cluster=factor(model$cluster))

centroids <- data.frame(model$centers, cluster=factor(1:3))

ggplot(iris_clustered, aes(x=SepalLengthCm, y=SepalWidthCm, color=cluster, shape=Species)) + geom_point
# individual points from the 'iris_clustered' data frame
geom_point(data=centroids, aes(fill=cluster), shape=21, color="black", size=3, stroke=1) # centroids
```

