#import dataset iris.txt as space as Seperator > data(iris) # Print value of iris > str(iris) 150 obs. of 5 variables: 'data.frame': \$ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ... \$ Sepal.Width: num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ... \$ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ... \$ Petal.Width: num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ... \$ Species : Factor w/ 3 levels "setosa", "versicolor", ..: 1111111111... > print(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 > # Identify the value of K > # make this example reproducible, use Seed > set.seed(1234) > # K max value to 10 to draw the plot between k and within class sum of square(wcss) > k.max <- 10

> wcss <- sapply(1:k.max,function(k){kmeans(data,k,nstart=6,iter.max=18)\$tot.withinss})

> data <- as.matrix(iris[,1:4])

> wcss

[1] 681.37060 152.34795 78.85144 57.26562 46.44618 41.70442 34.29823 29.98894 27.78609 26.41374

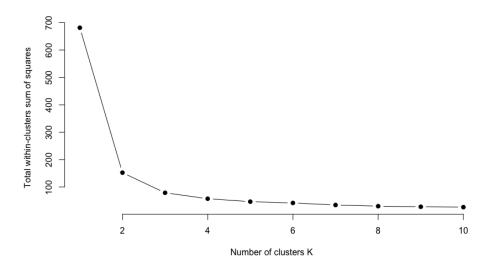
>

> # Plot wcss for k = 2 ..4..6..8

> plot(1:k.max, wcss,

- + type="b", pch = 19, frame = FALSE,
- + xlab="Number of clusters K",
- + ylab="Total within-clusters sum of squares")

>



> # List the within-class sum of squares for different sets of predictors

> # From the plot , ideal value would be 2 but from data, have noticed three classess of flowers so select K =3

> find_predictors = function(d){kmeans(d,3,nstart=6,iter.max=18)\$tot.withinss}

>

- > find_predictors(iris[,1]) #15.8
- [1] 15.81662
- > find_predictors(iris[,2]) # 5.34
- [1] 5.535155
- > find_predictors(iris[,3]) #24.66

```
[1] 24.51643
> find_predictors(iris[,4]) #4.91
[1] 4.913174
> find_predictors(iris[,1:2]) #37.05
[1] 37.0507
> find_predictors(iris[,2:3]) #40.73
[1] 40.73707
> find_predictors(iris[,3:4]) #31.37
[1] 31.37136
> find_predictors(iris[,c(1,3)]) #53.80
[1] 53.80998
> find_predictors(iris[,c(1,4)]) #32.73
[1] 32.73348
> find_predictors(iris[,c(2,3)]) #40.73
[1] 40.73707
> find_predictors(iris[,c(2,4)]) #20.60
[1] 20.6024
> find_predictors(iris[,1:3]) #69.42
[1] 69.42974
> find_predictors(iris[,2:4]) #47.86
[1] 47.86643
> find_predictors(iris[,1:4]) #78.85
[1] 78.85144
> find_predictors(iris[,c(1,2,4)])# 48.66
[1] 48.66078
```

```
> # Based on above predictions, best clustering value 78.85 found for 1:4, using this value for Best Model
> # Suggested K value is 3
> best_model <-kmeans(iris[,1:4],3,nstart=6,iter.max=18)
> table(iris[,5], best_model$cluster)
       1 2 3
 setosa 50 0 0
 versicolor 0 2 48
 virginica 0 36 14
> # using the best model , 1 , 2, 3
> #Setosa 50, 0, 0
> #VersiColor 0, 2, 48
> #Virginica 0 , 36, 14
R-SCript Only
#import dataset iris.txt as space as seperator
data(iris)
str(iris)
print(summary(iris))
```

>

```
# Identify the value of K
# make this example reproducible
set.seed(1234)
k.max <- 10
data <- as.matrix(iris[,1:4])</pre>
wcss <- sapply(1:k.max,function(k){kmeans(data,k,nstart=6,iter.max=18)$tot.withinss})
wcss
# Plot wcss for k = 2 ..4..6..8
plot(1:k.max, wcss,
  type="b", pch = 19, frame = FALSE,
  xlab="Number of clusters K",
  ylab="Total within-clusters sum of squares")
# List the within-class sum of squares for different sets of predictors
# From the plot , ideal value would be 2 but from data, have noticed three classess of flowers so select K
=3
find_predictors = function(d){kmeans(d,3,nstart=6,iter.max=18)$tot.withinss}
find_predictors(iris[,1]) #15.8
find_predictors(iris[,2]) # 5.34
find_predictors(iris[,3]) #24.66
find_predictors(iris[,4]) #4.91
find_predictors(iris[,1:2]) #37.05
find_predictors(iris[,2:3]) #40.73
find_predictors(iris[,3:4]) #31.37
```

```
find_predictors(iris[,c(1,3)]) #53.80
```

Best clustering value 78.85 found for 1:4, using this value for Best Model

Suggested K value is 3

best_model <-kmeans(iris[,1:4],3,nstart=6,iter.max=18)</pre>

table(iris[,5], best_model\$cluster)