# Multivariate Analysis (slides 9)

- Today we consider k-means clustering.
- We will address the question of selecting the appropriate number of clusters.
- Properties and limitations of the algorithm will be explored.

### k-means clustering

- The aim is to divide the data into k distinct groups so that observations within a group are similar, whilst observations between groups are different.
- k-means clustering is an iterative, rather than a hierarchical, clustering algorithm.
- This means that at each stage of the algorithm data points will be assigned to a fixed number of clusters (contrast with hierarchical clustering where the number of clusters ranges from the number of data points down to a single cluster).
- We will discuss ways of selecting an appropriate k from a statistical viewpoint, but there may be expert knowledge as to the appropriate number of clusters.
- Alternatively, there may be previous results from preliminary data exploration, *i.e.*, we could start the k-means algorithm at the result of a hierarchical clustering.

### k-means clustering

- It is simple and computationally efficient, but can sometimes be sensitive to the selection of starting points.
- Running the k-means algorithm several times for different starting values can help check whether results are robust.
- We will see an example of the problems this can cause.

### Pseudo code

- 1. Choose the number of clusters k and designate cluster centers.
- 2. Assign each data point to the cluster whose center is closest.
- 3. For cluster i, calculate its centroid  $C_i^T = (C(i)_1, C(i)_2, \ldots, C(i)_m)$ , where m denotes the number of variables in an observation (these are found by averaging variables scores for data points within the cluster).
- 4. Calculate the sum of squared distances of each object to its cluster centroid:

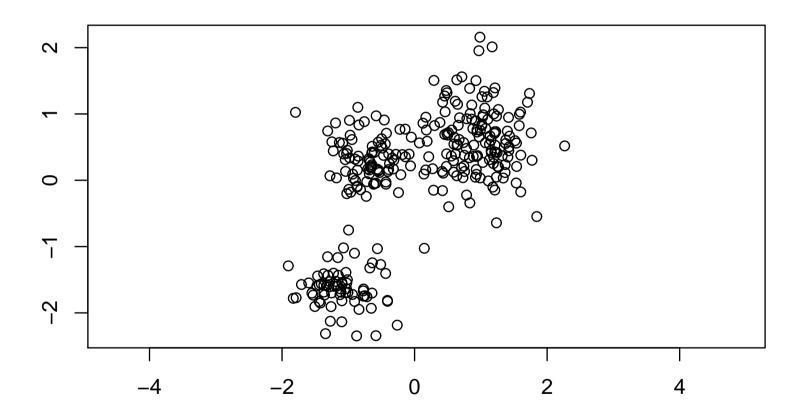
SS = 
$$\sum_{i=1}^{n} \sum_{j=1}^{m} (x_{ij} - C(i)_j)^2$$

Here we assume a total of n observations. We want the SS value to be as small as possible.

- 5. Re-assign each observation to the cluster whose centroid is closest.
- 6. Repeat (3)-(5) until convergence.

### Simulated Data

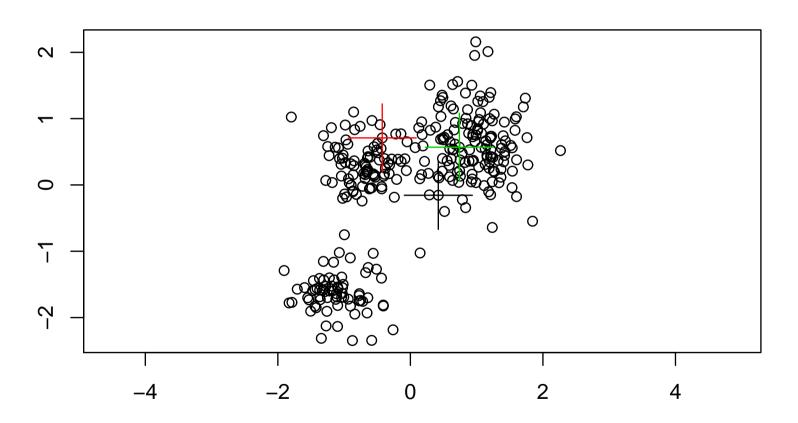
• Consider the following simulated data.



• We want to cluster the data into, say, three groups.

# k-Means Clustering: Iteration 0a

• We start by randomly generating three centers (prototypes).



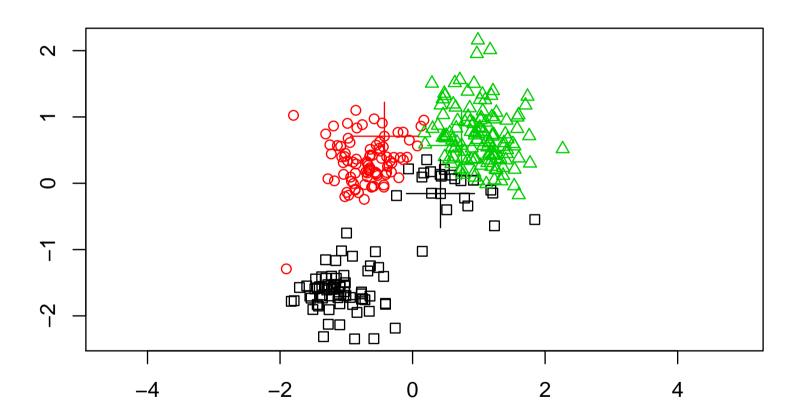
# k-Means Clustering: Iteration 0a

The initial partition can be constructed in several ways, e.g.,

- 1. A random selection of k observations.
- 2. Specify selection based on prior knowledge.
- 3. By using results from an exploratory hierarchical clustering algorithm.

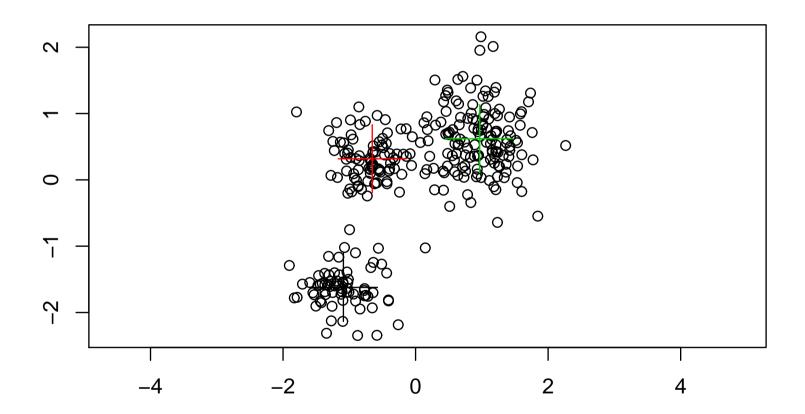
# k-Means Clustering: Iteration 0b

• Label points according to which center is closest.



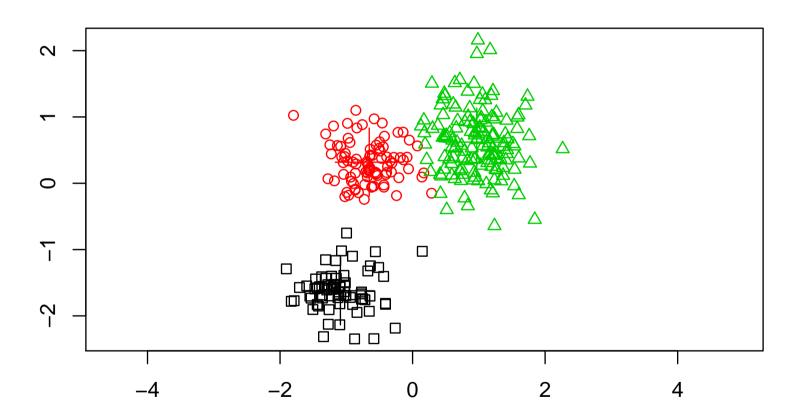
# k-Means Clustering: Iteration 1a

• Update the values for the three centers (prototypes).



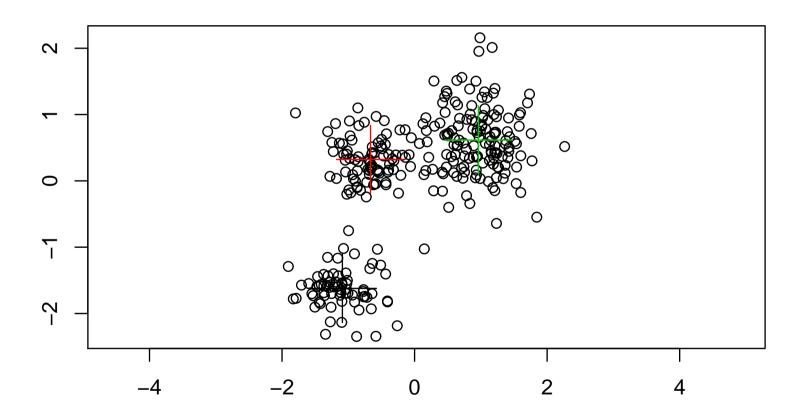
# k-Means Clustering: Iteration 1b

• Label points according to which center is closest.



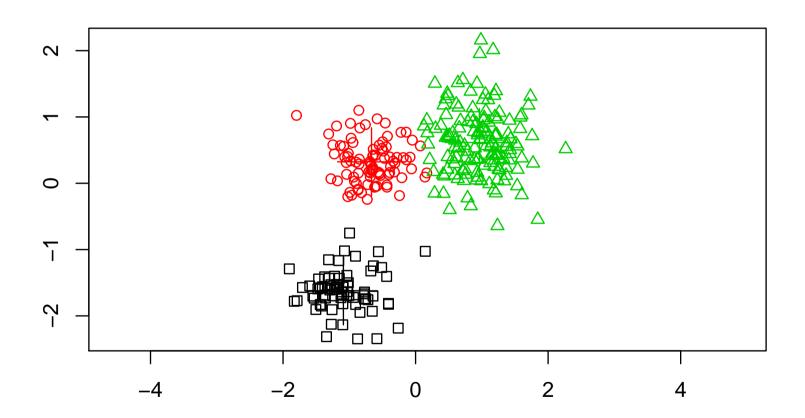
# k-Means Clustering: Iteration 2a

• Update the values for the three centers (prototypes).



# k-Means Clustering: Iteration 2b

• We label points according to which center is closest.



### Convergence

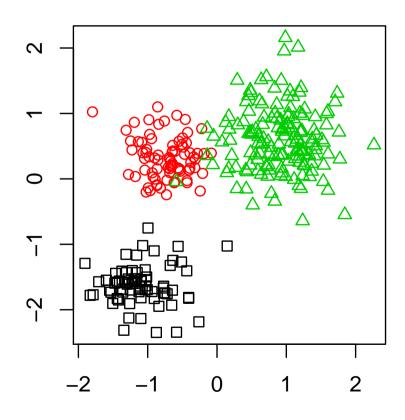
- The k-Means algorithm has converged when no points are moved between groups on an iteration.
- Once this happens, the estimates of the centers will no longer change, nor will the allocation of points to groups thereafter.
- This convergence criteria might not be suitable in some cases, e.g., if n is very large, and alternatives are possible, e.g., within cluster sum of squares does not change over 3 iterations etc.

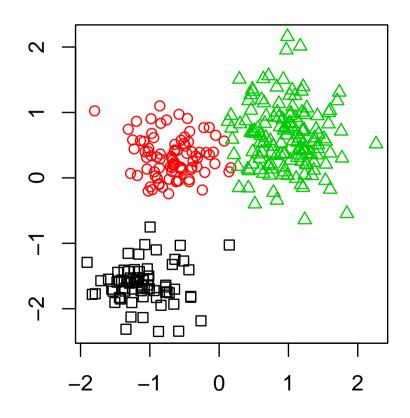
### How Did It Do?

- The data from the last example had been simulated, so that there were actually three groups in the data.
- $\bullet$  How well did k-means perform at finding these groups?

### **True Groups**

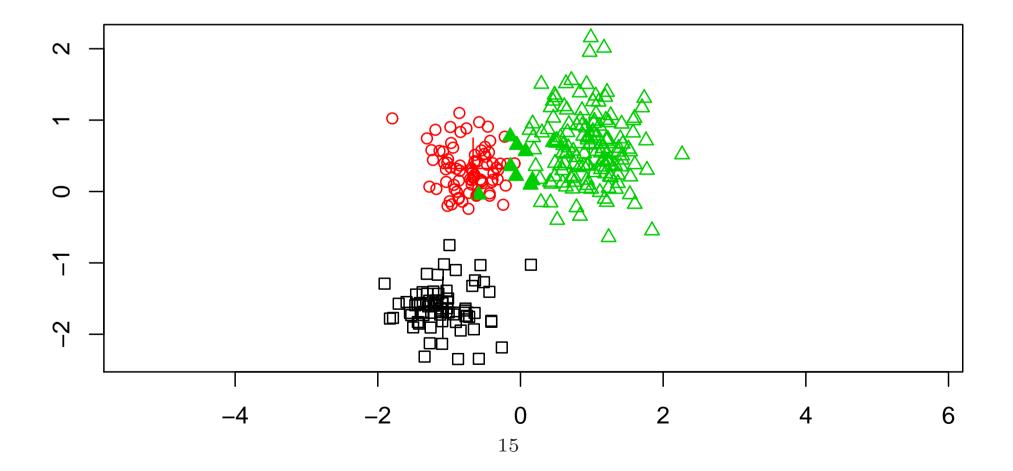
#### Classification





# Any errors?

- The coloured in points were misclassified.
- $\bullet$  Only 8/300 were misclassified.

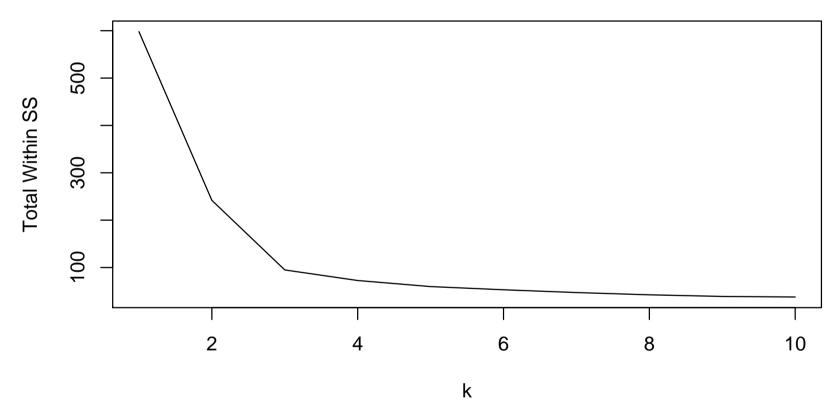


### Choosing The Value For k

- This is not an exact science, but there are guidelines.
- Generally we should run the k-means algorithm for a differing number of values for k, e.g., k = 1, ..., 10.
- When running k-means the aim is to minimize the SS, so why not choose k to minimize the SS?
- However, the more clusters that are fitted the smaller the SS (think of what would happen if we selected k = n).
- A general rule is to plot k against SS and look for a 'kink' in the curve. If there is no kink then there is a trade-off between additional complexity by increasing k and better fit by reducing the SS.

### Choosing The Value For k

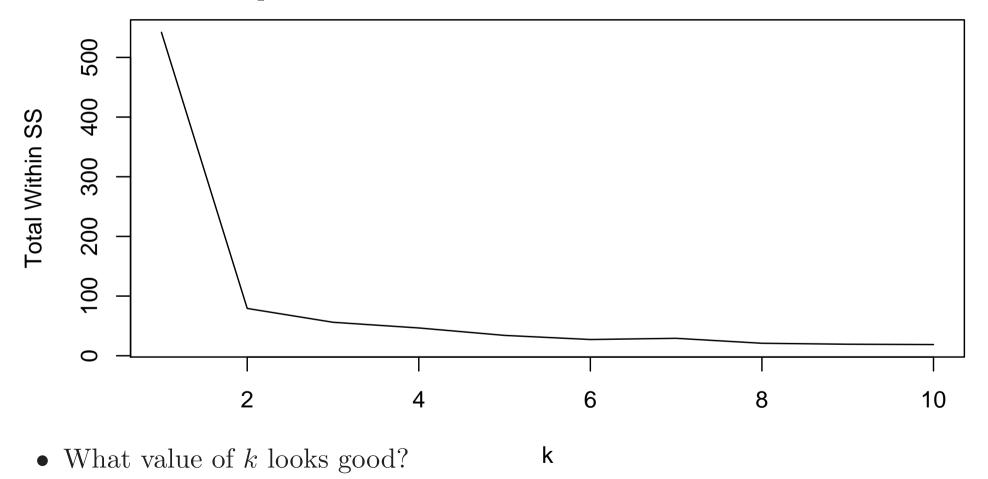
• If we plot the total of the within sum of squares values versus k, then we get the following:



• Notice that the graph flattens very quickly. What k would you use?

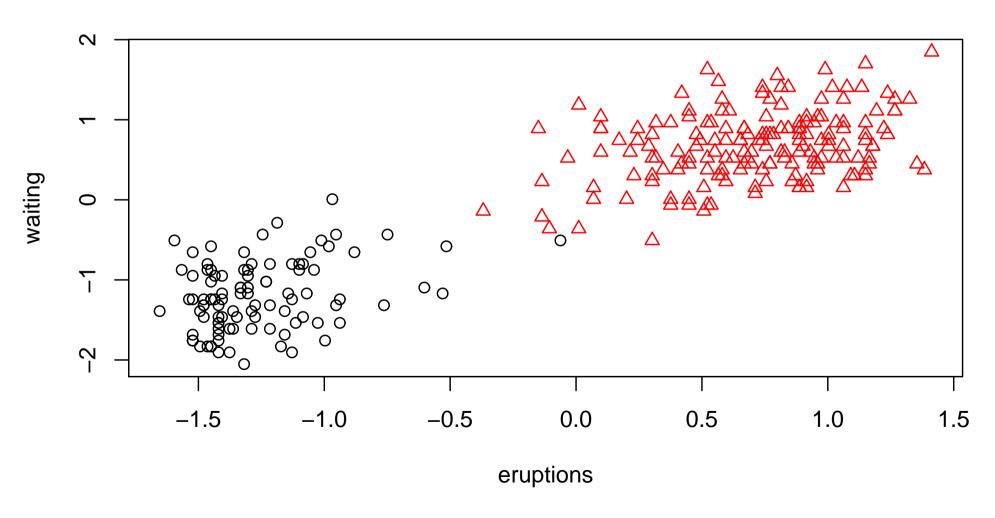
### Old Faithful Data

• Running k-means on the standardized Old Faithful data allows a plot of the within sum of squares values versus k:



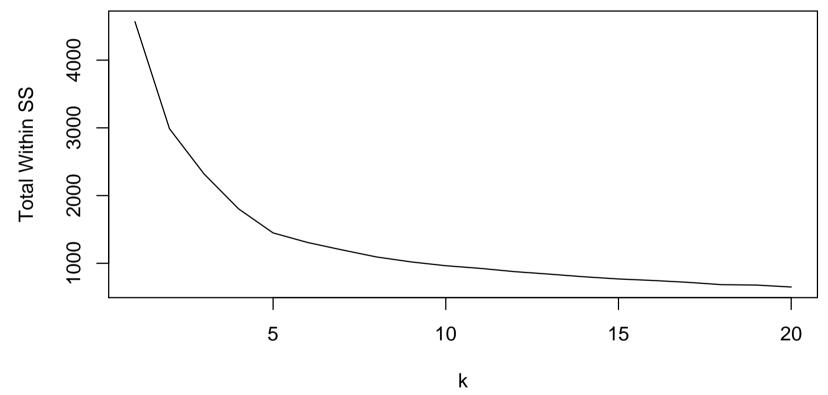
### Old Faithful Data

• This provides the following clustering of the data:



### Olive Oil Data

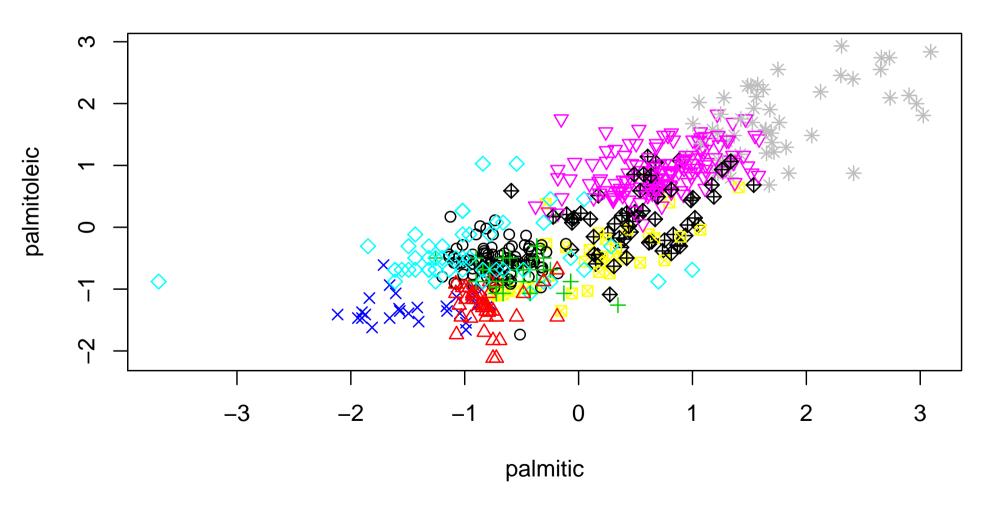
• Running k-means on the standardized olive oil data data allows a plot of the within sum of squares values versus k:



• What value of k looks good? Let's look at k = 9.

### Olive Oil Data

• This provides the following clustering of the data:



### Cross Tabulation

• A cross tabulation of the olive oil regions (rows) and the clusters (columns) shows some agreement:

```
1 22 2
       0 0 0 0
       0 23 0 0 1 0
2 0 32
 0 12 144 1 0 49 0 0
  6 16
       0 12 0
       0 0 65
5
  0 0
              0 0 0
  0 0
       0 0 33
              0
                 0
       0 0 0 0 33 10
  0 0
  0 0
       0 0 0 0
                 0 50
       0
          0
  0
   0
            0
             0
                 1
                   0 50
```

### Faithful Data k=2

• Consider the k=2 solution for the Faithful data:

Number of clusters: 2

```
      Number of Obs.
      WSS
      Avg. Dist. to Centroid

      Cluster 1
      100
      3456.2
      4.9

      Cluster 2
      172
      5445.6
      4.6

      Sum
      272
      8901.8
```

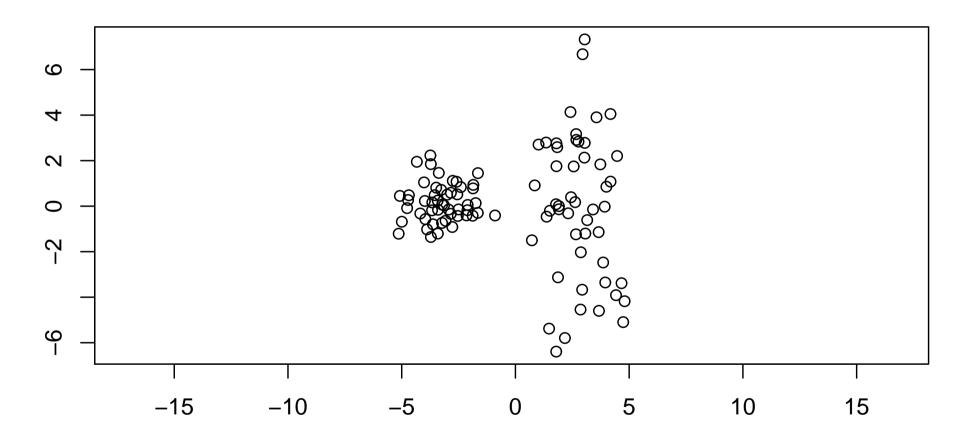
Cluster Centroids:

```
Cluster 1 Cluster 2 Total Data eruptions 2.1 4.3 3.5 waiting 54.8 80.3 70.9
```

Distance Between Cluster Centroids:

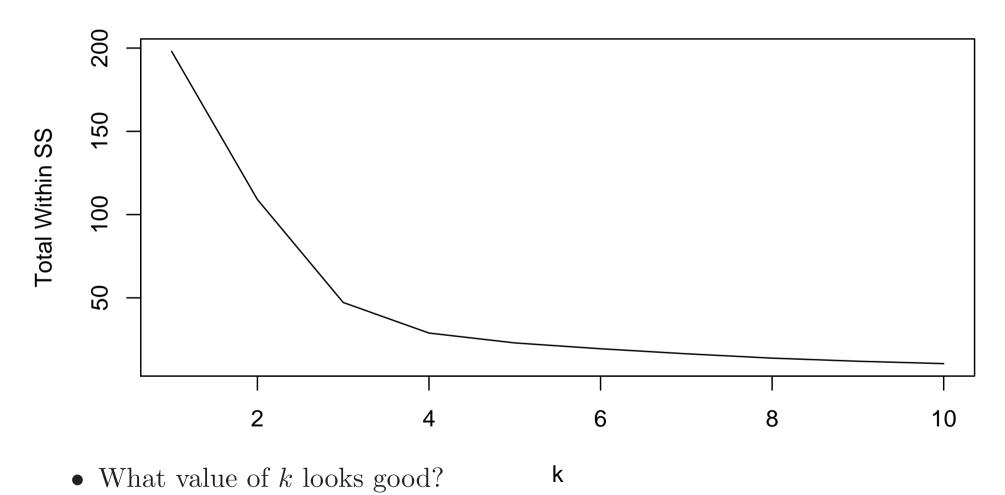
Cluster 1 Cluster 2
Cluster 1 0.0 25.6
Cluster 2 25.6 0.0

• What if we run k-means on the following, more tricky, standardized data:

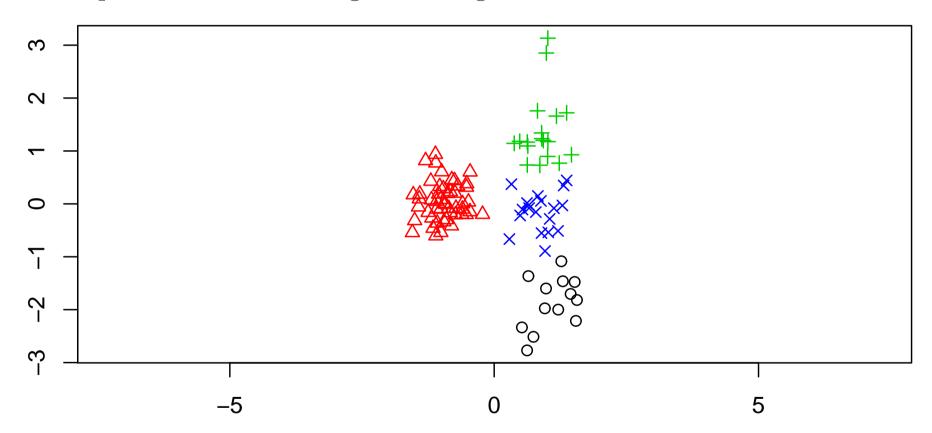


• It looks like there should be two groups.

• Plotting the within sum of squares values versus k gives:



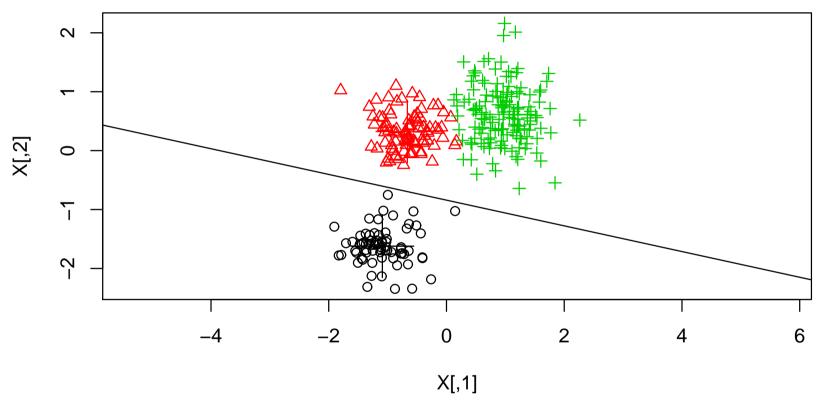
• This provides the following clustering of the data:



 $\bullet$  The elliptical group is broken into subgroups. This is because k-means clustering looks for circular clusters.

### Distance From Means

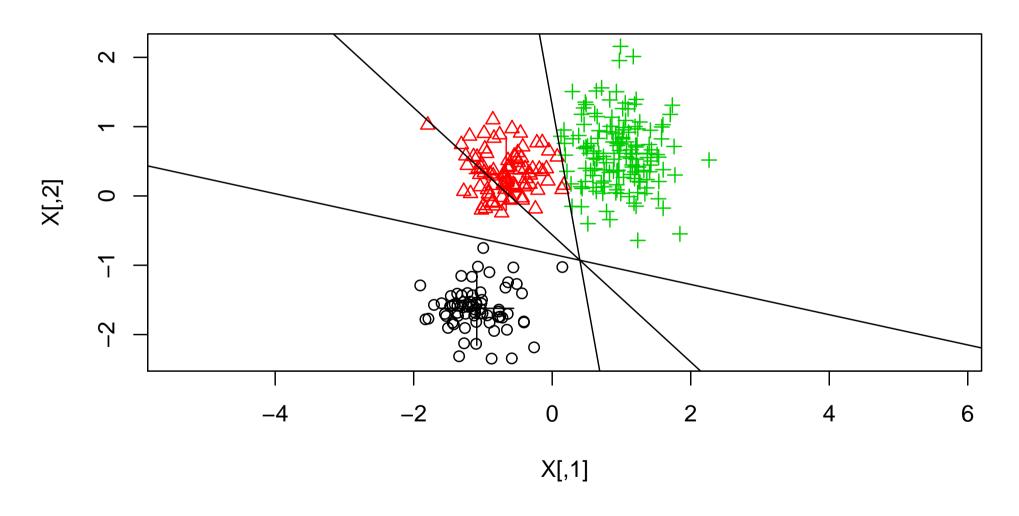
• Return to the first data set.



• Consider the line separating the points that are closest to the mean of the triangles ( $\triangle$ ) and the points closest to the mean of the circles ( $\circ$ ).

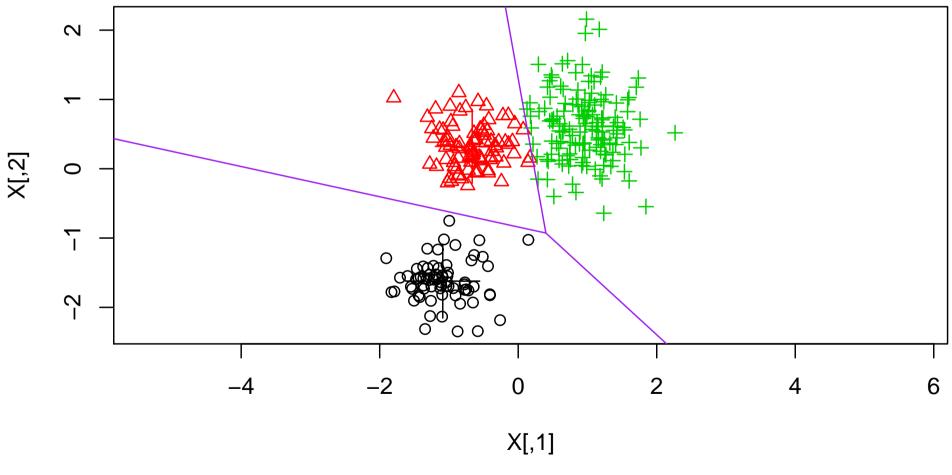
### Distance From Means

• Including such lines for each pair of means:

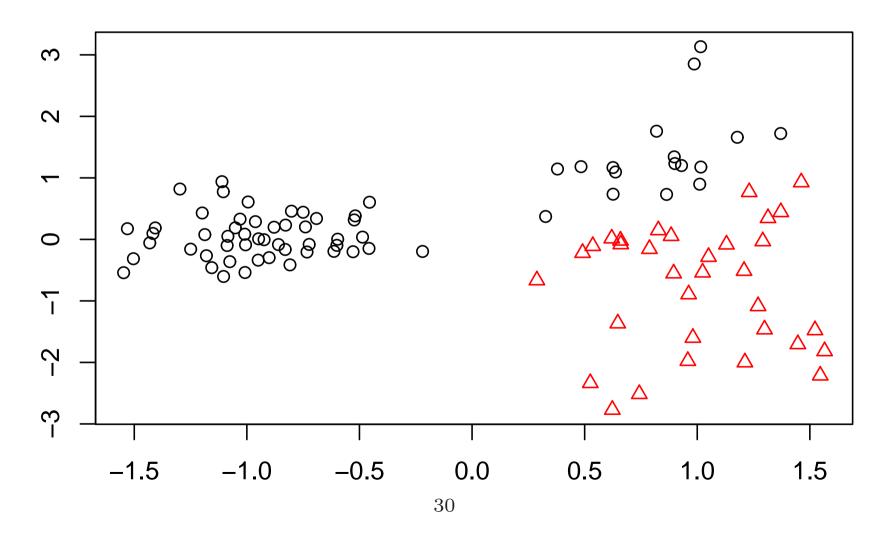


# Partitioning Of The Plane

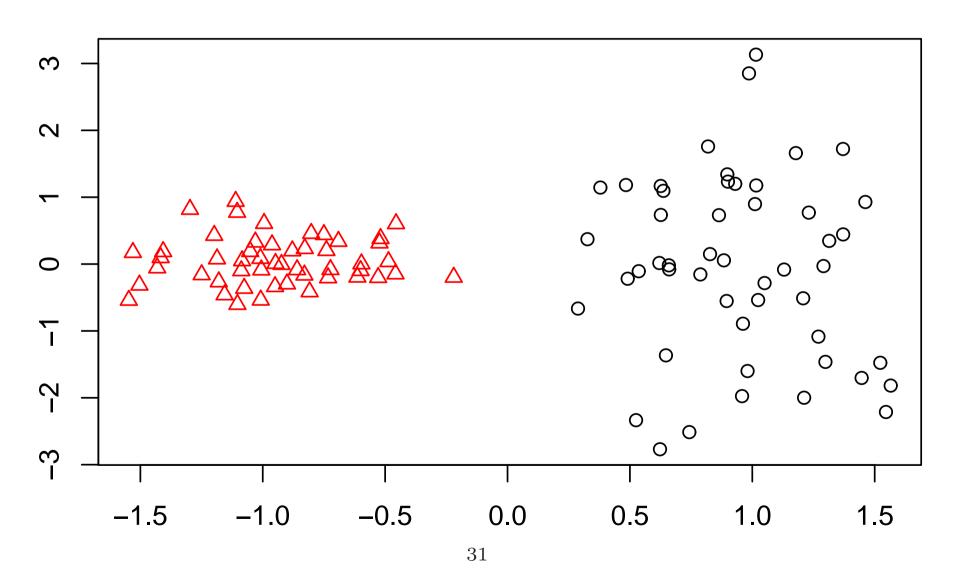
• The plane is partitioned into three polygonal regions depending on which mean is closest.



- Recall the tricky data from earlier.
- If we cluster it into two groups, we get:



• But if we ran k-means from a different and specific starting point:



### Local Minima

- The k-means algorithm can give different answers when initiated at different starting values.
- This means that the algorithm does not always find the minimum value for the Total Within Sum of Squares.
- The Total Within Sum of Squares for the first clustering is 82.4+36.3=118.7.
- The Total Within Sum of Squares for the second clustering is 11.0+98.1=109.1.
- Therefore, the second set of results is better.