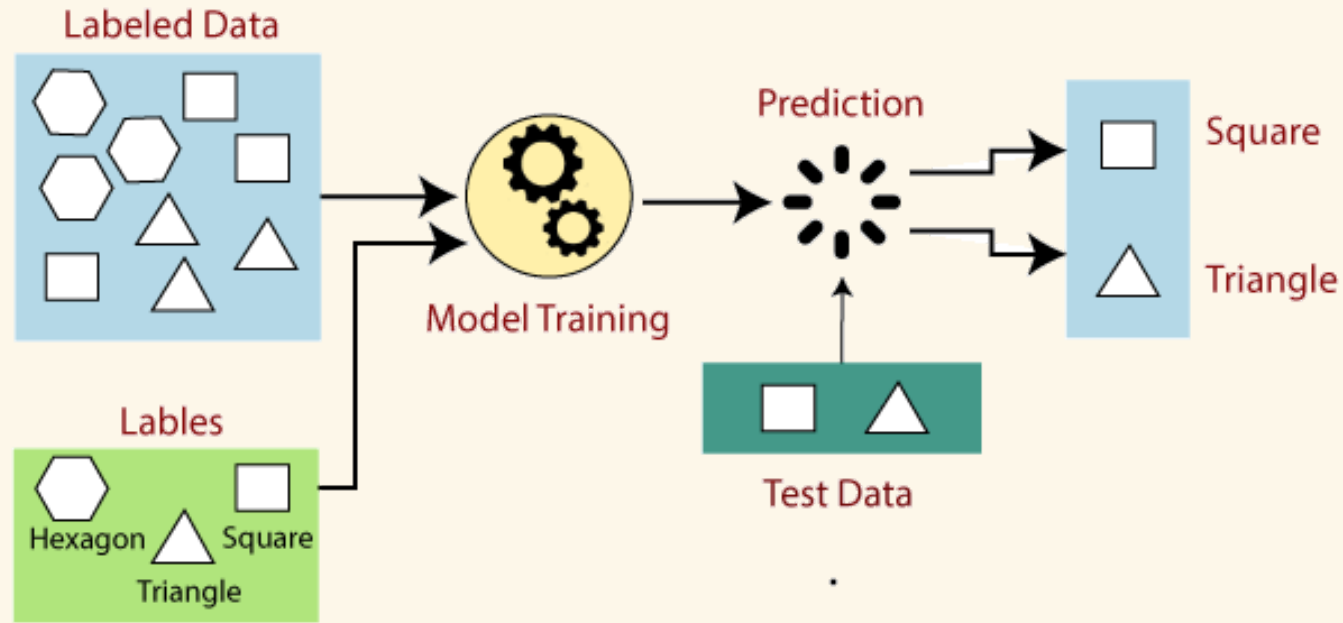


Intro to Clustering

Fall 2022

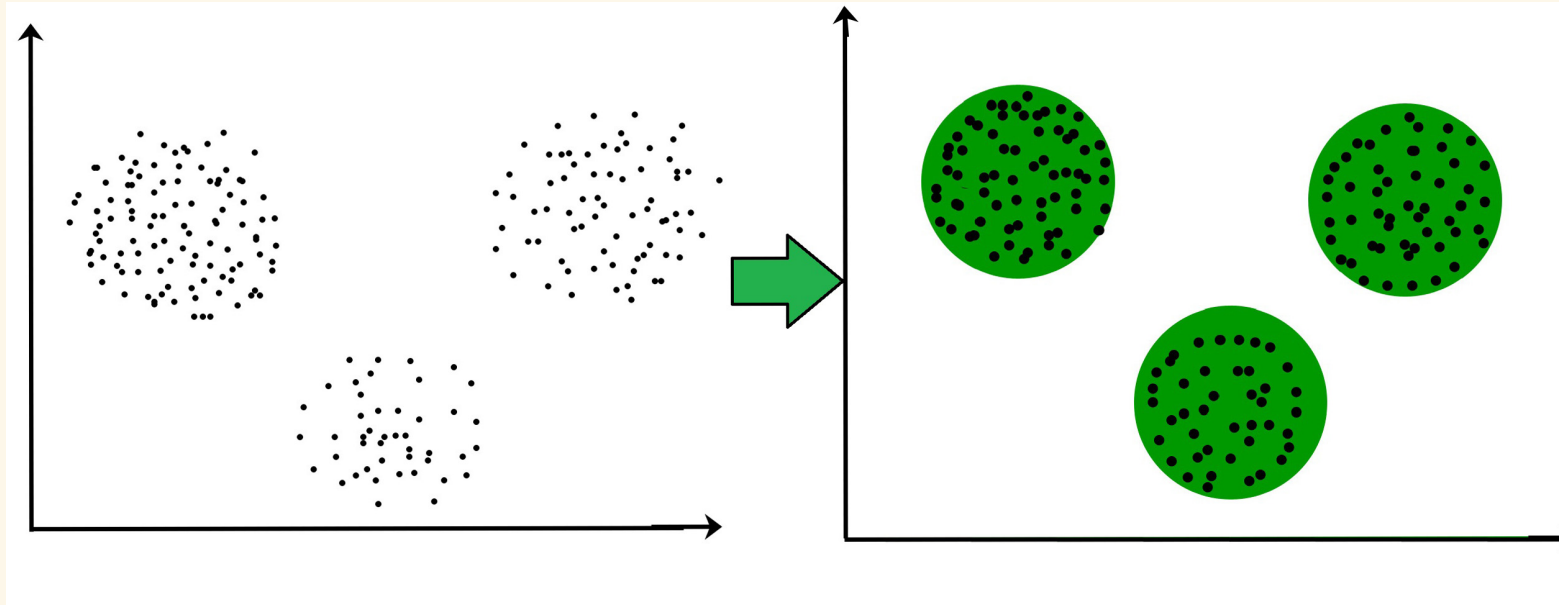
November 09 2022

Supervised learning



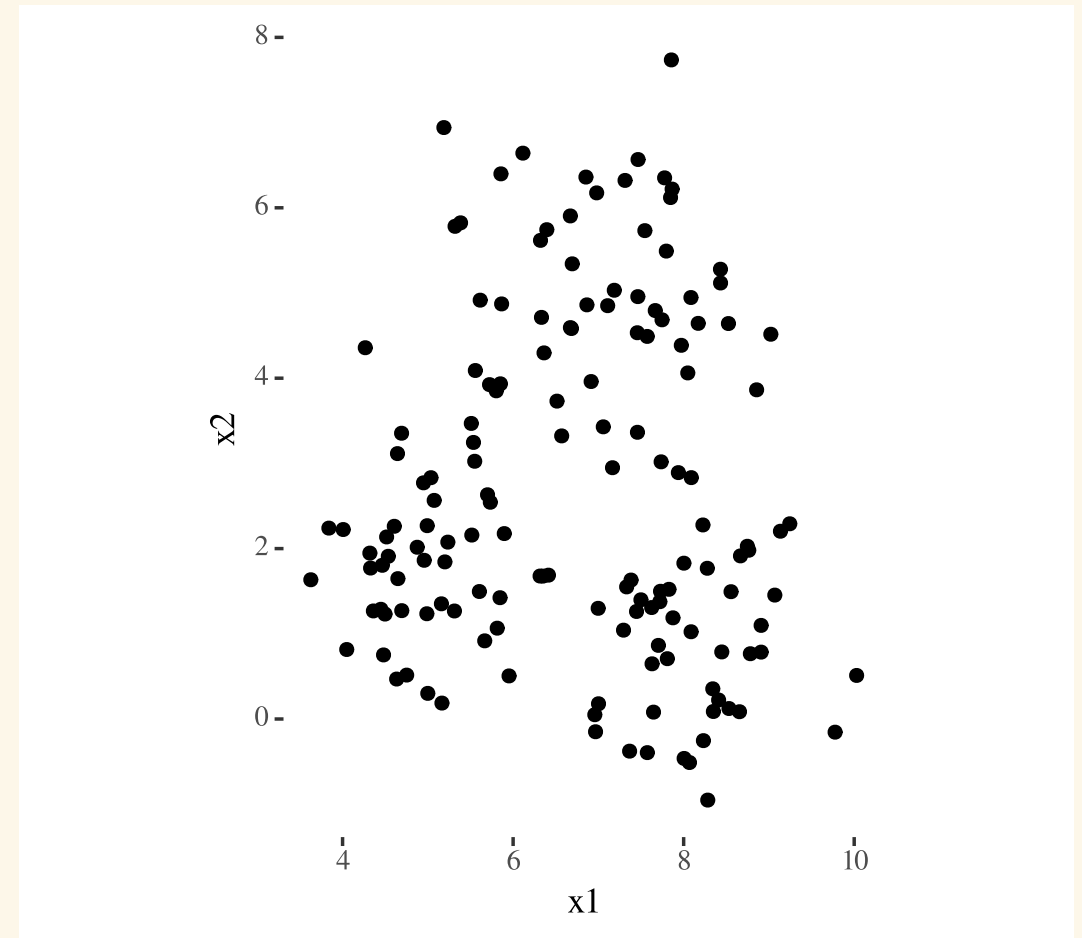
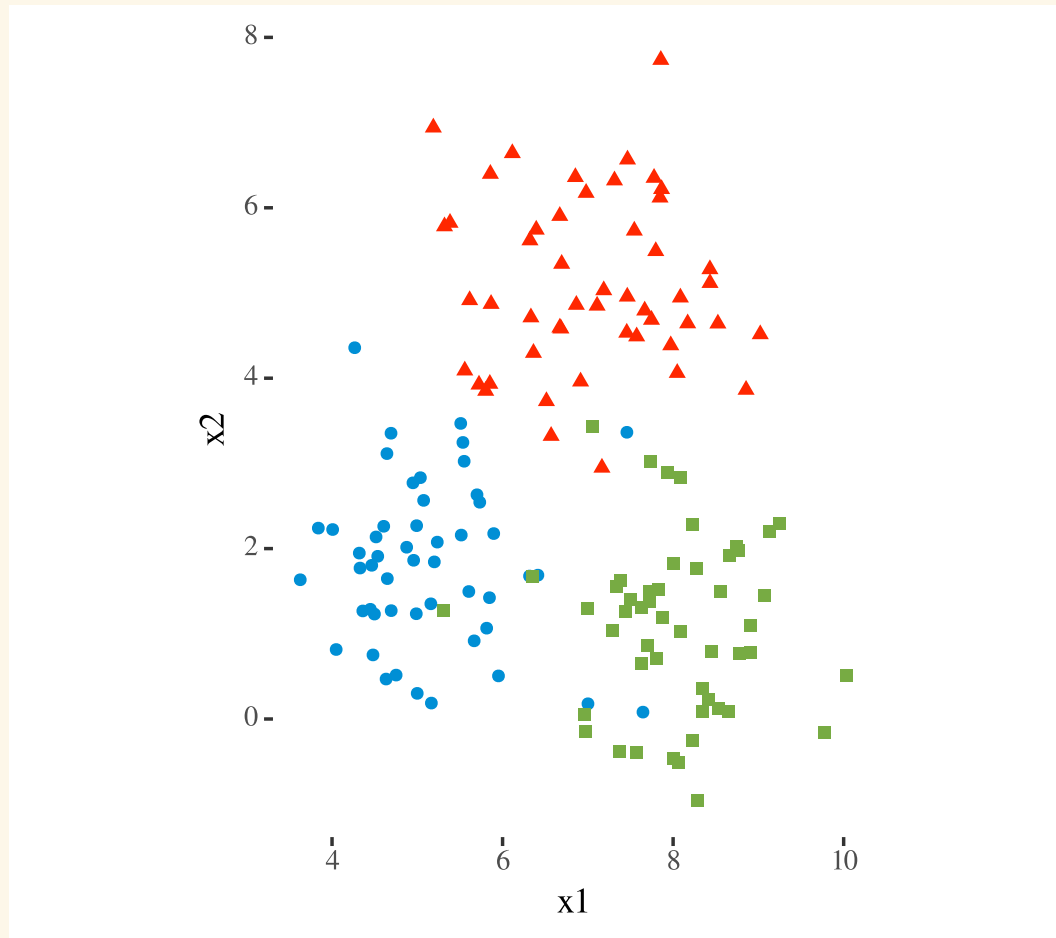
- train or "supervise" algorithms to use labels to classify data or predict outcomes
- use labeled inputs and outputs to measure model accuracy

Unsupervised learning



- uses statistical learning algorithms to analyze and cluster unlabeled data sets
- discover hidden patterns in data without human intervention, so "unsupervised"
 - group unlabeled data based on their similarities or differences

Example: get cluster association from unlabeled data



Can use an unsupervised algorithm called k-means to achieve this!

K-means Basics

- Algorithm to group data into K clusters
- Starts with an initial clustering of data
- Iteratively improves the cluster assignments
- Stops until the assignments cannot be improved further

Algorithm

1. Randomly assign a number, from 1 to K , to each of the observations
2. Compute the centroid of each of the K clusters
3. Assign each point to the nearest centroid and redefine the cluster
4. Repeat steps 2 and 3 until no point change clusters

Main Idea

To minimize the total within cluster variation

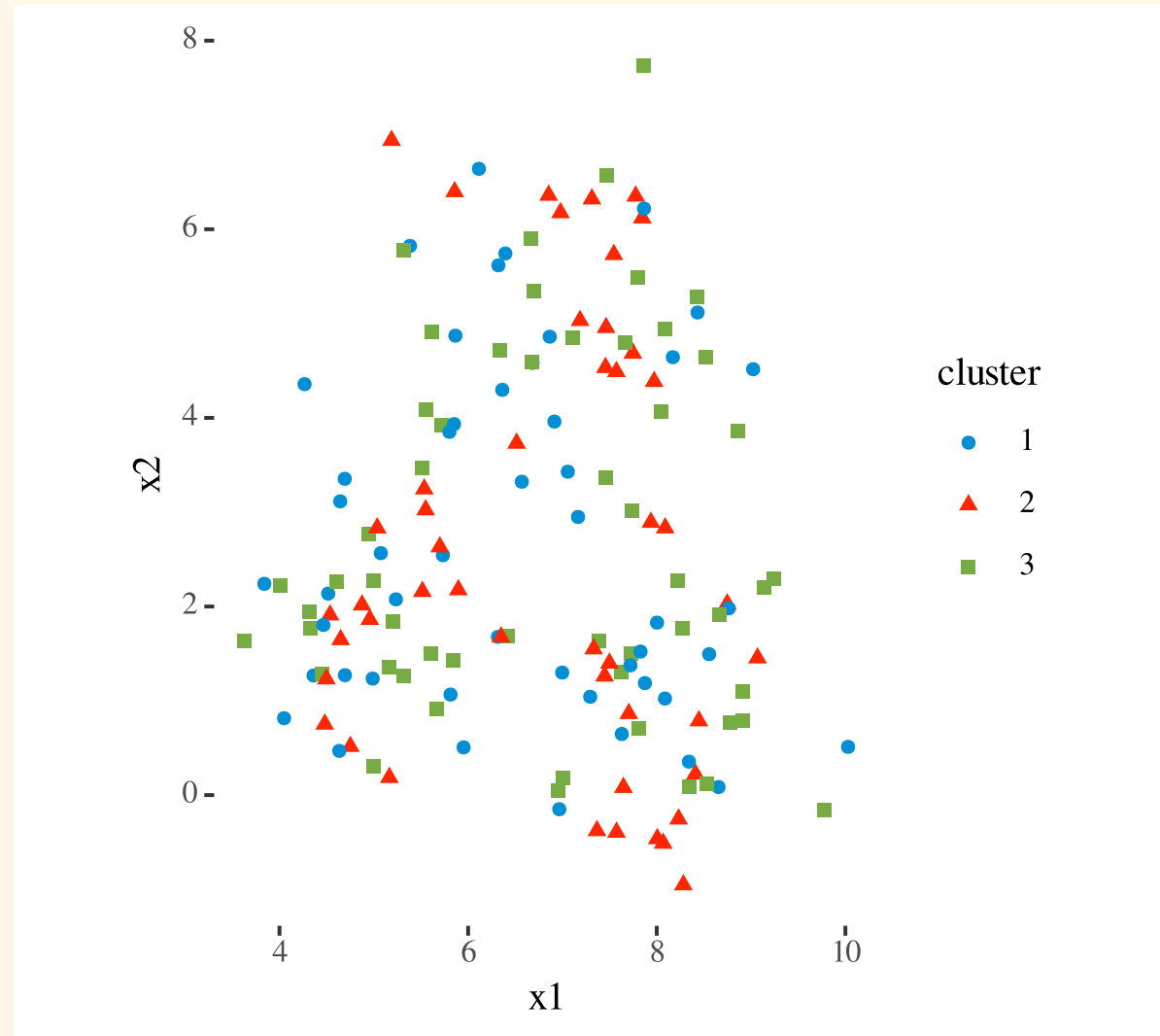
The total within-cluster variation is the sum of squared Euclidean distances between items and the corresponding centroid:

$$WSS = \sum_{k=1}^K WSS(C_k) = \sum_{k=1}^K \sum_{x_i \in C_k} (x_i - \mu_k)^2$$

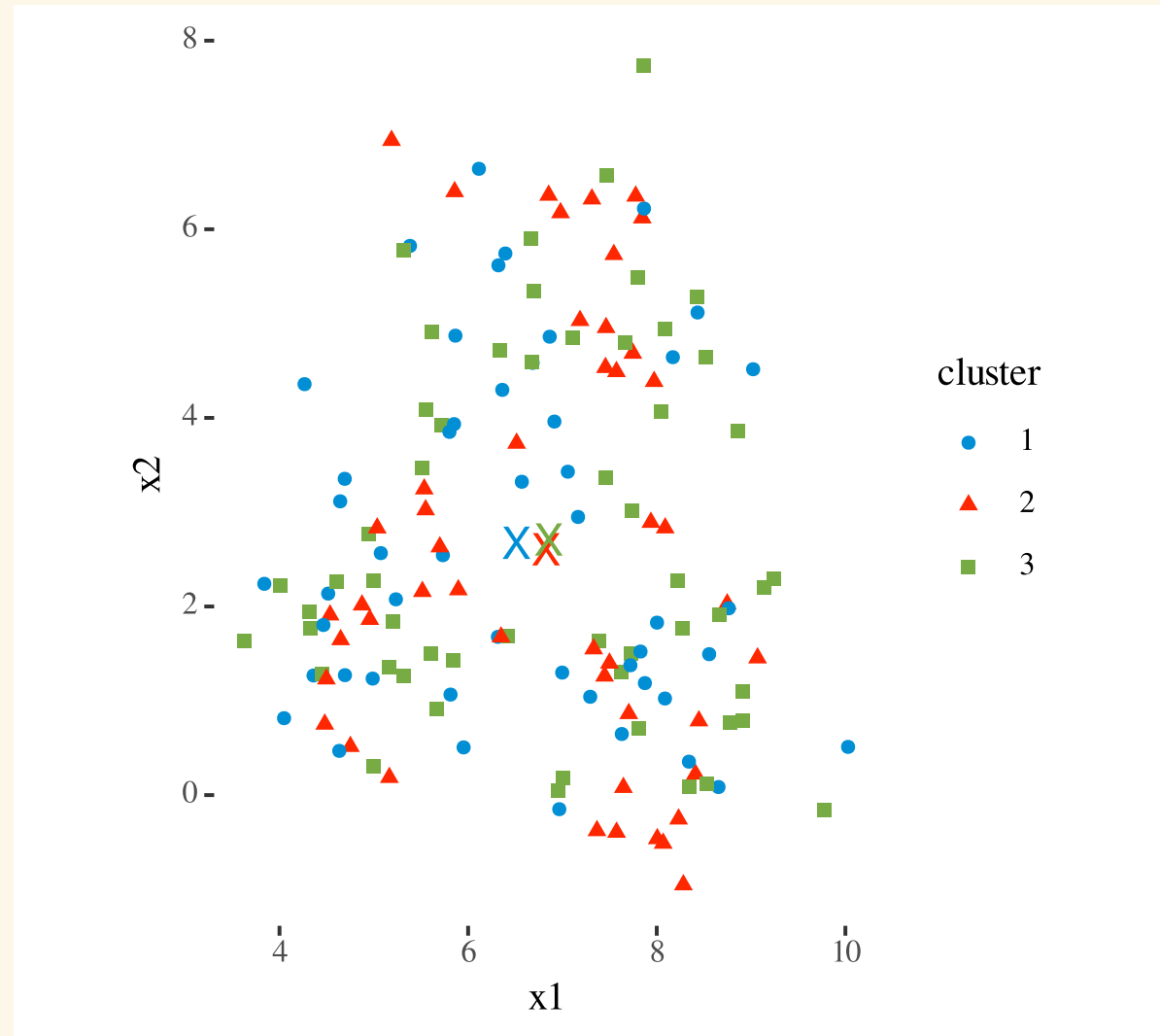
where:

- WSS is the Within Cluster Sum of Squared Errors
- x_i is a data point in the cluster C_k
- μ_k is the mean value of the points assigned to the cluster C_k

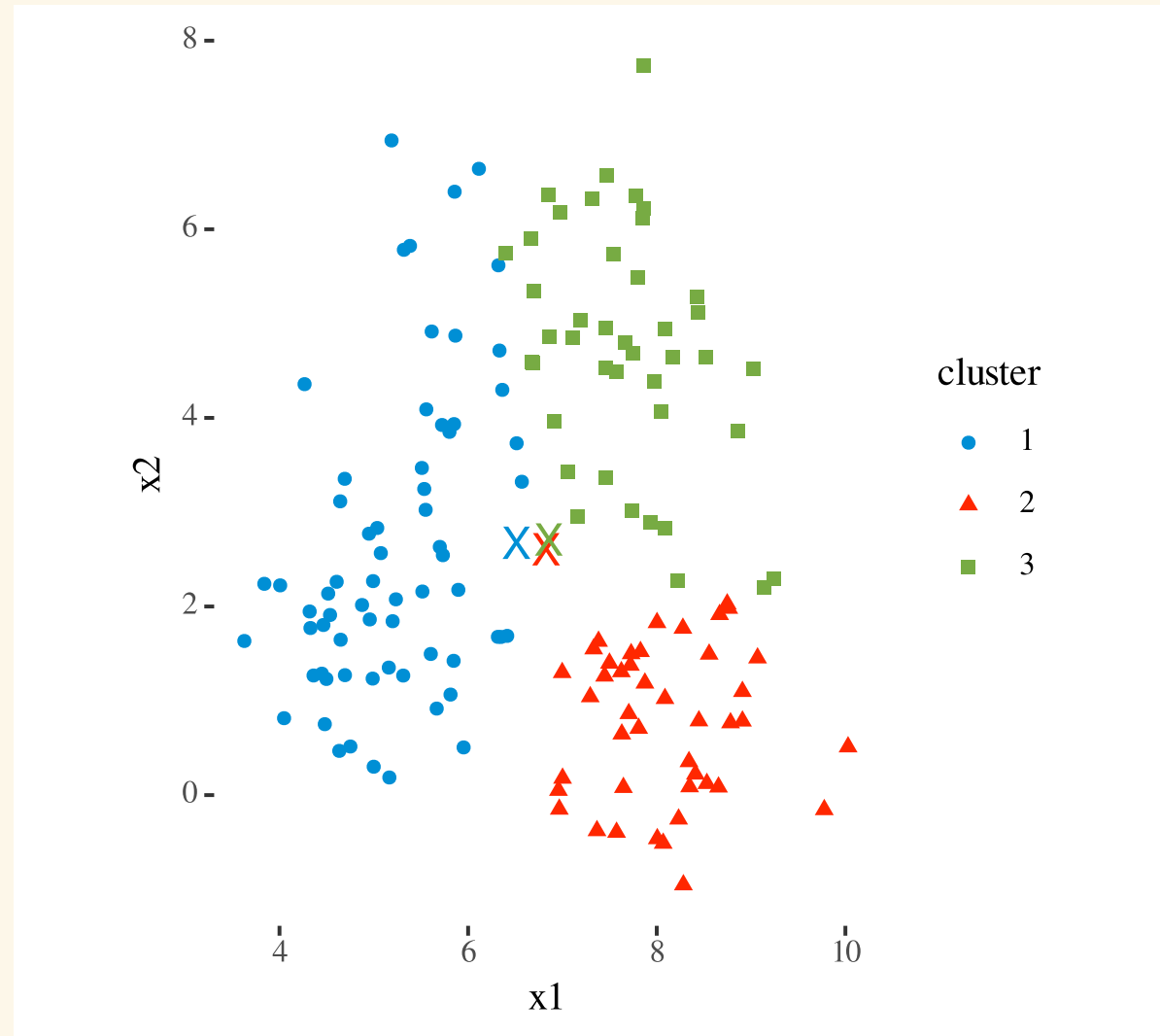
(1). Randomly assign a number, from 1 to K , to each of the observations



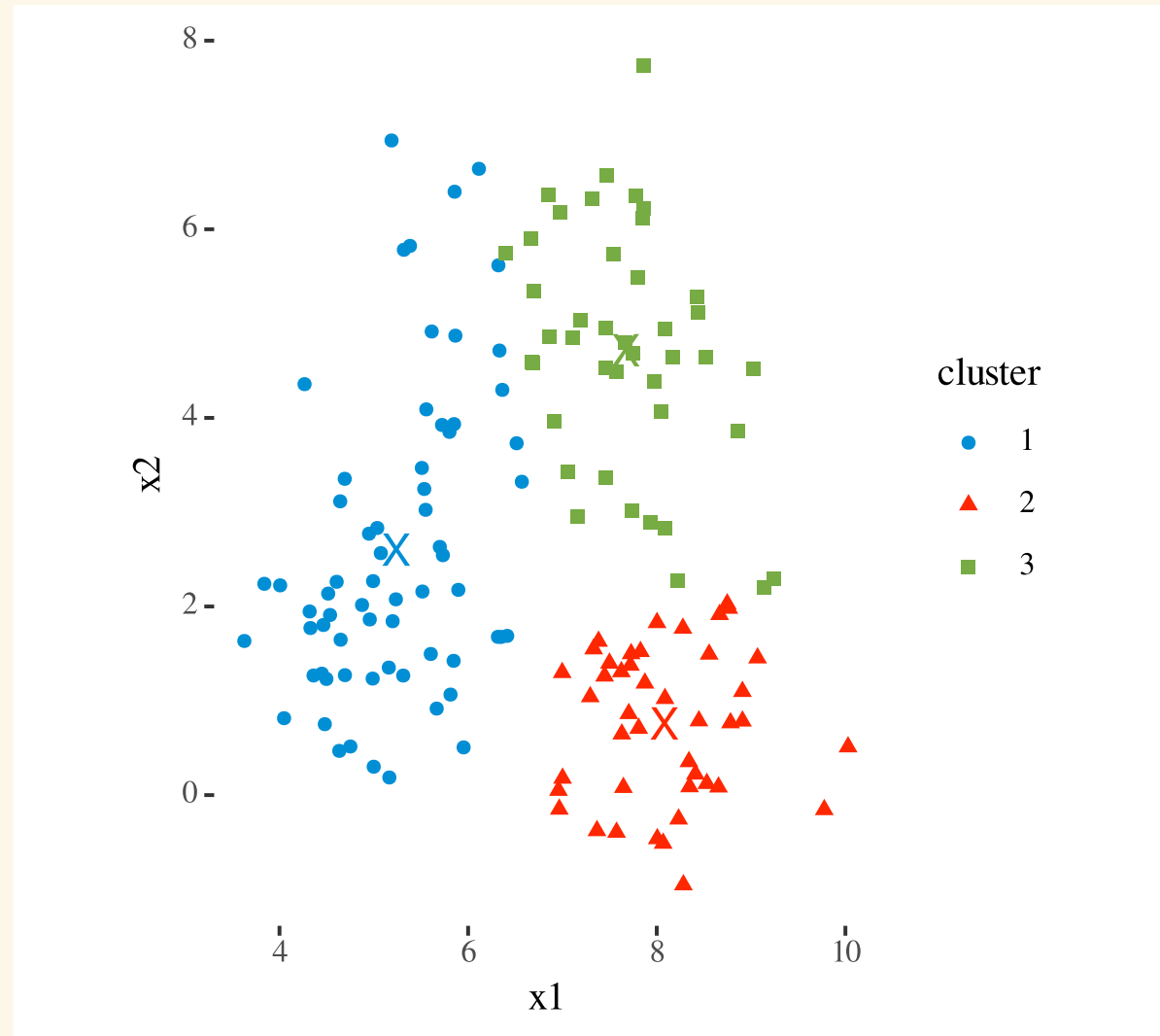
(2). Compute the centroid of each cluster



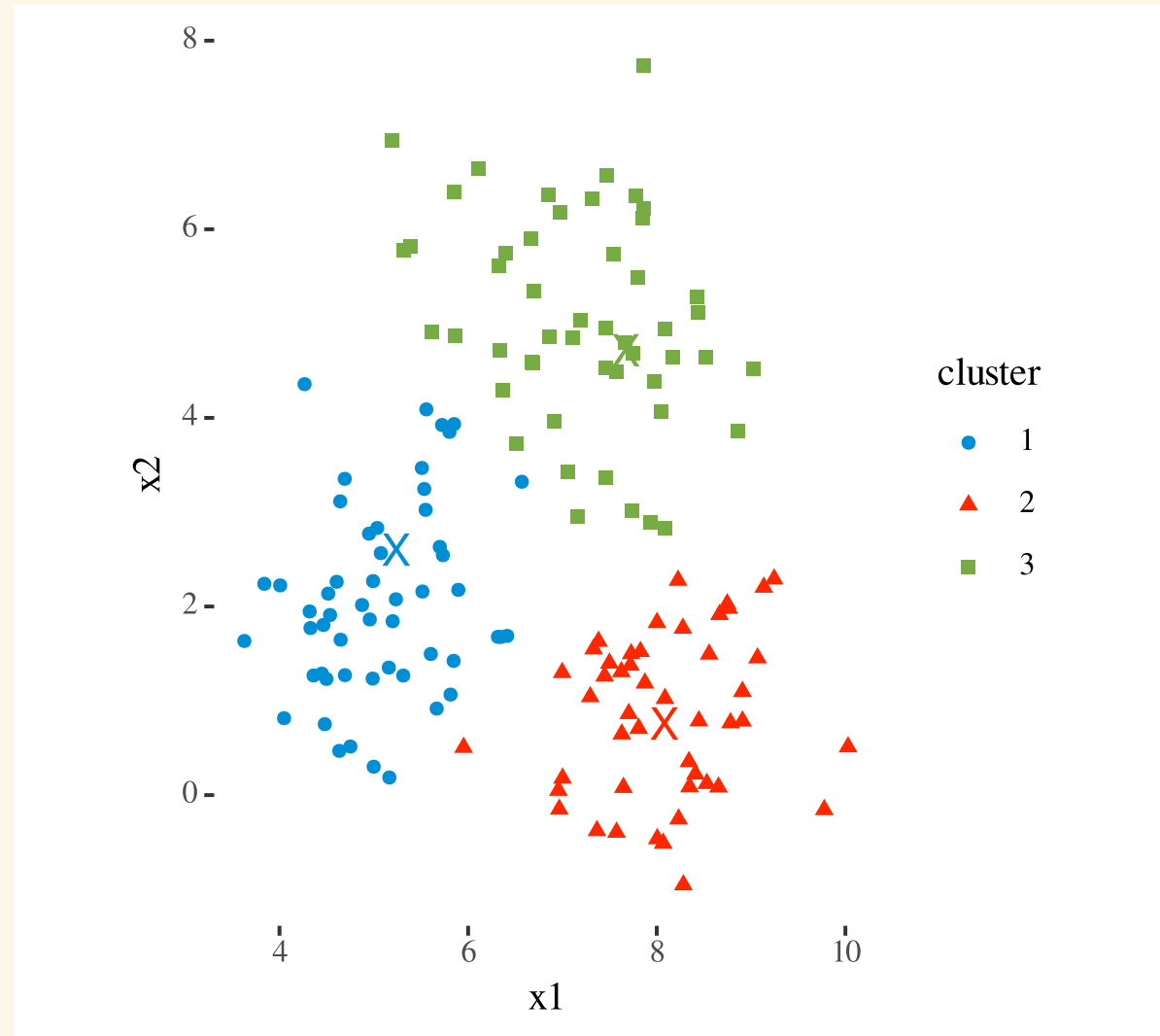
(3). Re-assign each observation to the cluster whose centroid is closest



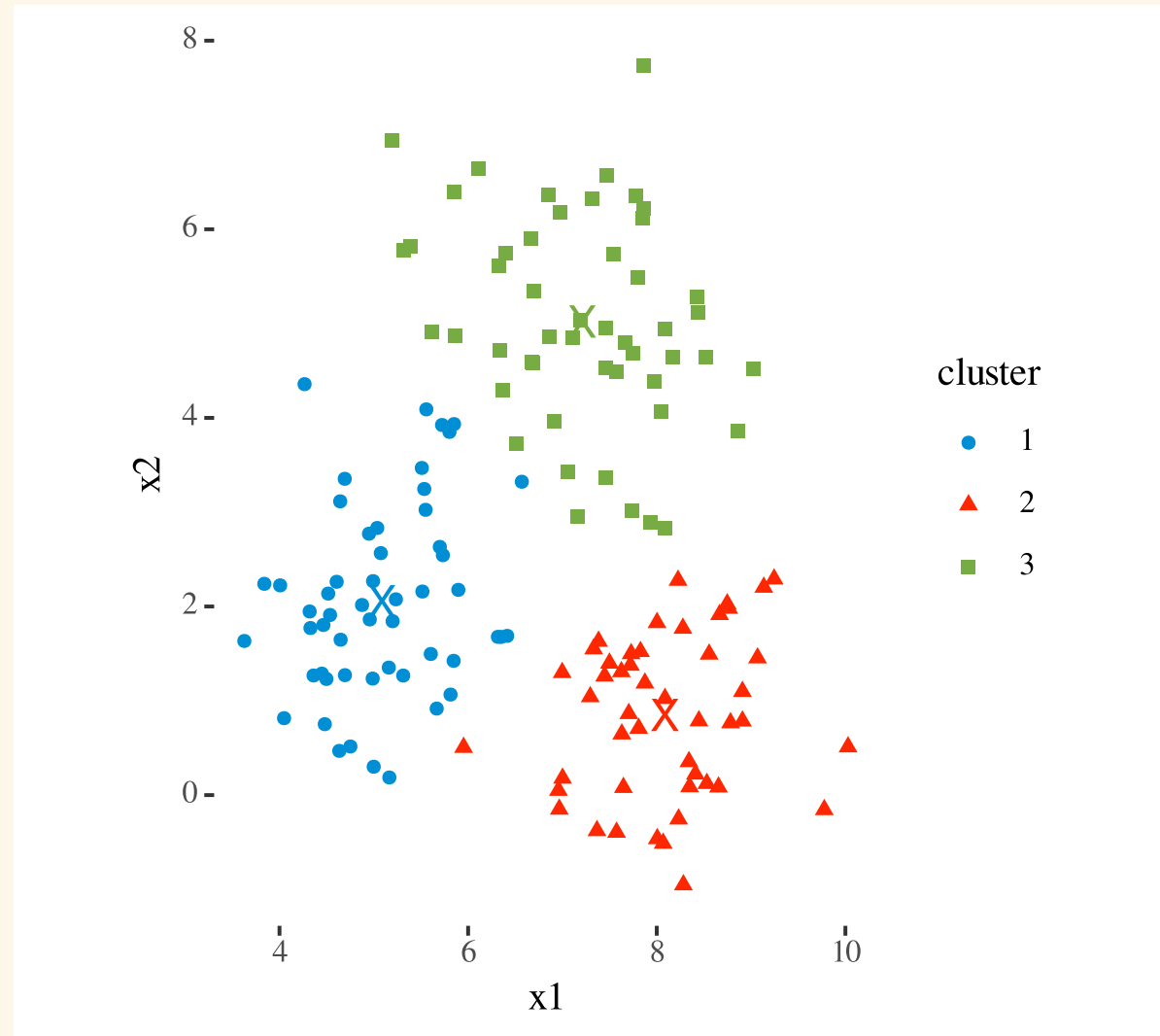
(4). Re-compute the centroid of each cluster



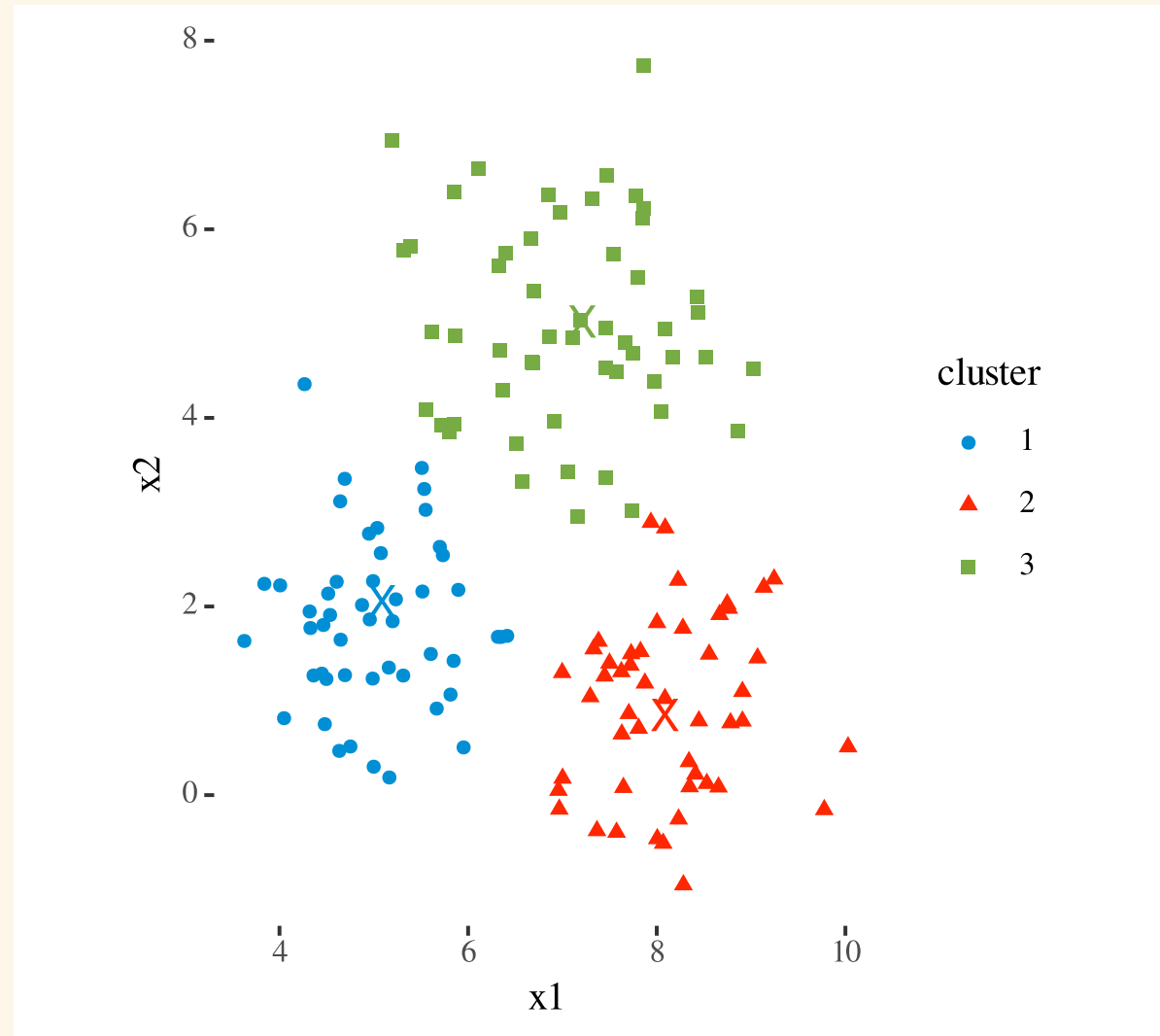
(5). Re-assign each observation to the cluster whose centroid is closest



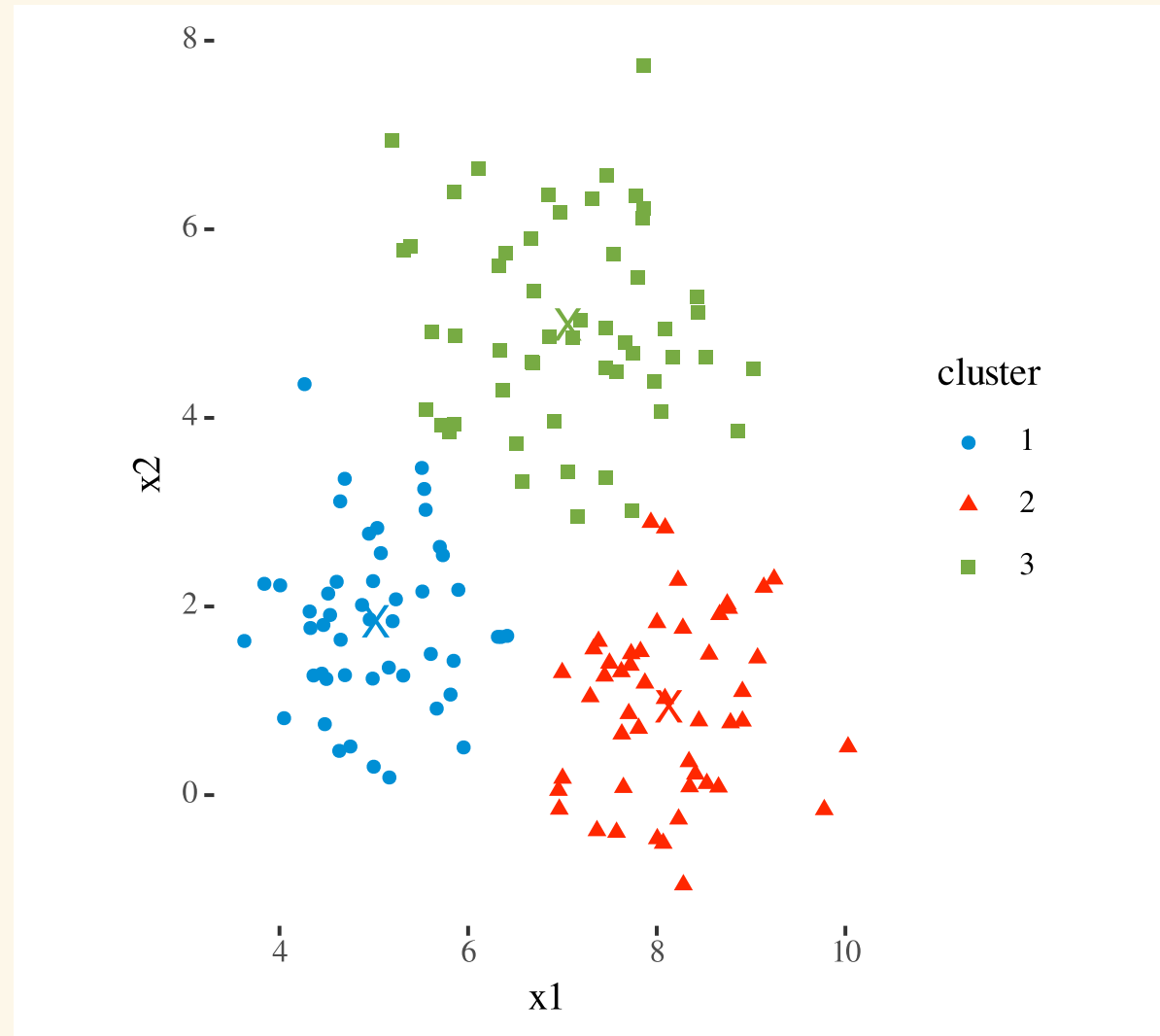
(6). Re-compute the centroid of each cluster



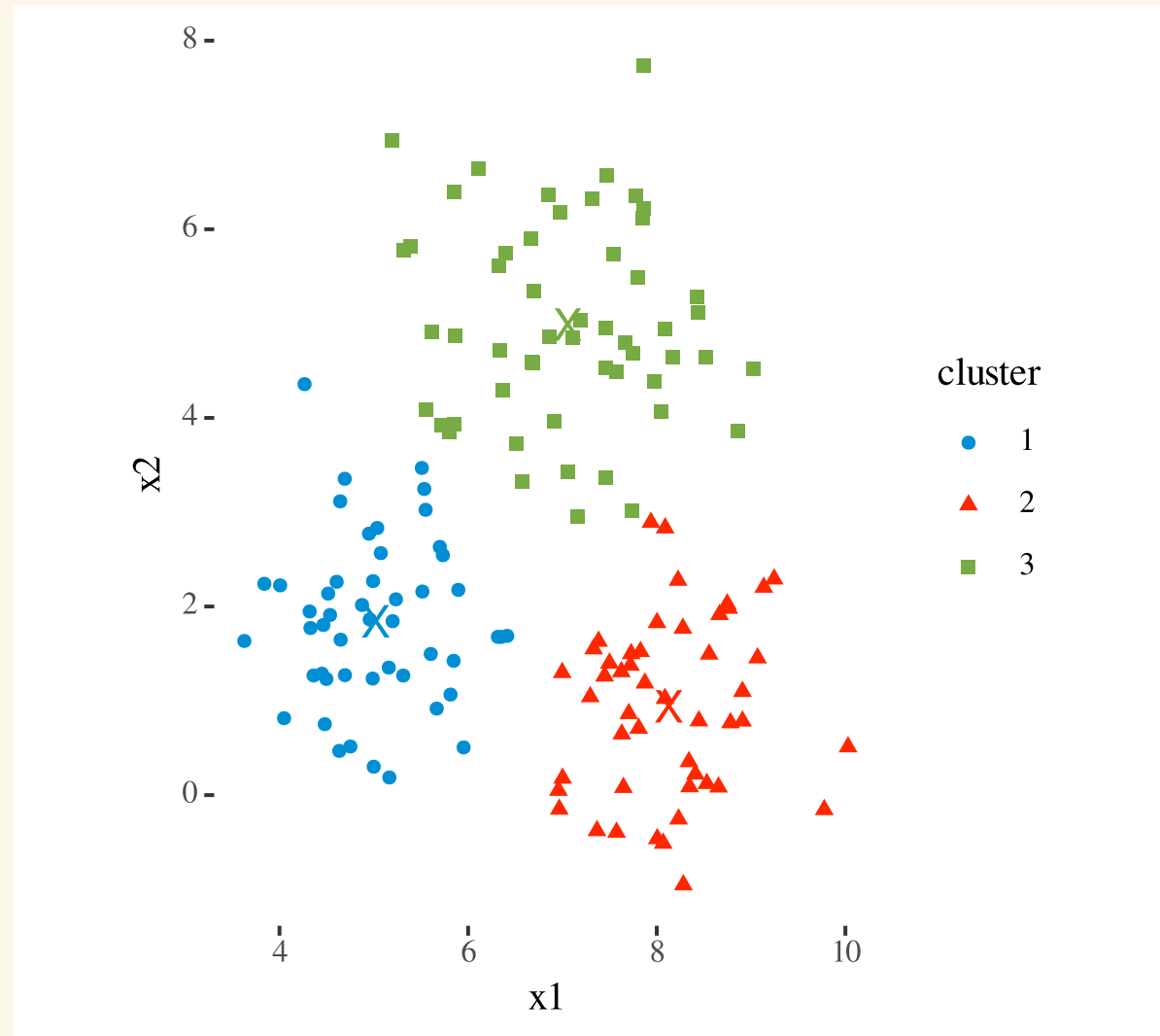
(7). Re-assign each observation to the cluster whose centroid is closest



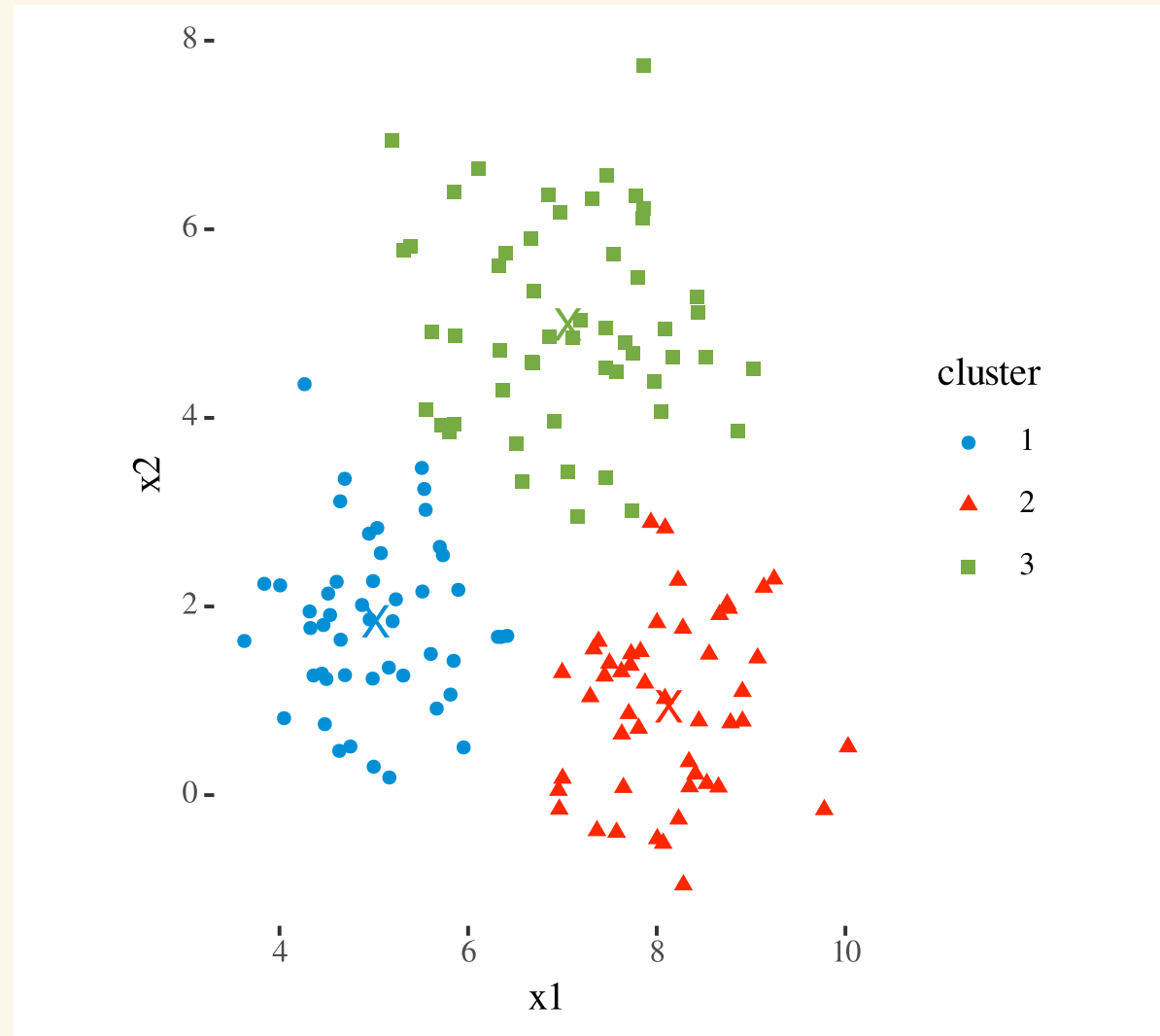
(8). Re-compute the centroid of each cluster



(9). Re-assign each observation to the cluster whose centroid is closest



(10). Re-compute the centroid of each cluster



USArrests

```
USData <- as_tibble(USArrests, rownames = "state") %>% drop_na() %>%  
  column_to_rownames("state") %>%  
  select(Murder, UrbanPop)
```

```
head(USData, 10)
```

	Murder	UrbanPop
Alabama	13.2	58
Alaska	10.0	48
Arizona	8.1	80
Arkansas	8.8	50
California	9.0	91
Colorado	7.9	78
Connecticut	3.3	77
Delaware	5.9	72
Florida	15.4	80
Georgia	17.4	60

Means and standard deviations

```
USAData %>%  
  map_dfr(mean)  
# A tibble: 1 × 2  
  Murder UrbanPop  
  <dbl>    <dbl>  
1    7.79    65.5
```

```
USAData %>%  
  map_dfr(sd)  
# A tibble: 1 × 2  
  Murder UrbanPop  
  <dbl>    <dbl>  
1    4.36    14.5
```

Standardize the data

```
USData <- USData %>% mutate(across(where(is.numeric), standardize))
```

```
head(USData,10)
```

	Murder	UrbanPop
Alabama	1.24256408	-0.5209066
Alaska	0.50786248	-1.2117642
Arizona	0.07163341	0.9989801
Arkansas	0.23234938	-1.0735927
California	0.27826823	1.7589234
Colorado	0.02571456	0.8608085
Connecticut	-1.03041900	0.7917228
Delaware	-0.43347395	0.4462940
Florida	1.74767144	0.9989801
Georgia	2.20685994	-0.3827351

So, how do we fit all of this in R?

kmeans()

- `kmeans()` function takes a matrix or data-frame or tibble and the number of centers/clusters we want to find.
- We also set `nstart = 20-25` to have multiple initial starting positions in the hope of finding global optimal solution instead of local optimal solution
- Use `set.seed()` for reproducibility

Within Cluster Sum of Squared Errors (WSS)

- Calculate WSS for different values of K.
- Choose K for which WSS first starts to diminish.
- Visually deciphered with an elbow graph.
- The number of clusters is taken at the elbow joint point.

K-means

```
set.seed(1234)  
k.means <- kmeans(USADData, centers = 2, nstart = 25)
```



```
k.means
```

```
K-means clustering with 2 clusters of sizes 23, 27
```

```
Cluster means:
```

```
      Murder   UrbanPop
1  0.8961762  0.1939808
2 -0.7634094 -0.1652429
```

```
Clustering vector:
```

Alabama	Alaska	Arizona	Arkansas	California
1	1	1	2	1
Colorado	Connecticut	Delaware	Florida	Georgia
1	2	2	1	1
Hawaii	Idaho	Illinois	Indiana	Iowa
2	2	1	2	2
Kansas	Kentucky	Louisiana	Maine	Maryland
2	1	1	2	1
Massachusetts	Michigan	Minnesota	Mississippi	Missouri
2	1	2	1	1
Montana	Nebraska	Nevada	New Hampshire	New Jersey
2	2	1	2	1
New Mexico	New York	North Carolina	North Dakota	Ohio
1	1	1	2	2
Oklahoma	Oregon	Pennsylvania	Rhode Island	South Carolina
2	2	2	2	1
South Dakota	Tennessee	Texas	Utah	Vermont
2	1	1	2	2
Virginia	Washington	West Virginia	Wisconsin	Wyoming
1	2	2	2	2

```
Within cluster sum of squares by cluster:
```

```
[1] 31.59219 30.59764
```

```
(between_SS / total_SS = 36.5 %)
```

Tidy the information

```
k.means %>% tidy()  
# A tibble: 2 × 5  
  Murder UrbanPop  size withinss cluster  
  <dbl>    <dbl> <int>    <dbl> <fct>  
1  0.896    0.194   23    31.6  1  
2 -0.763   -0.165   27    30.6  2
```

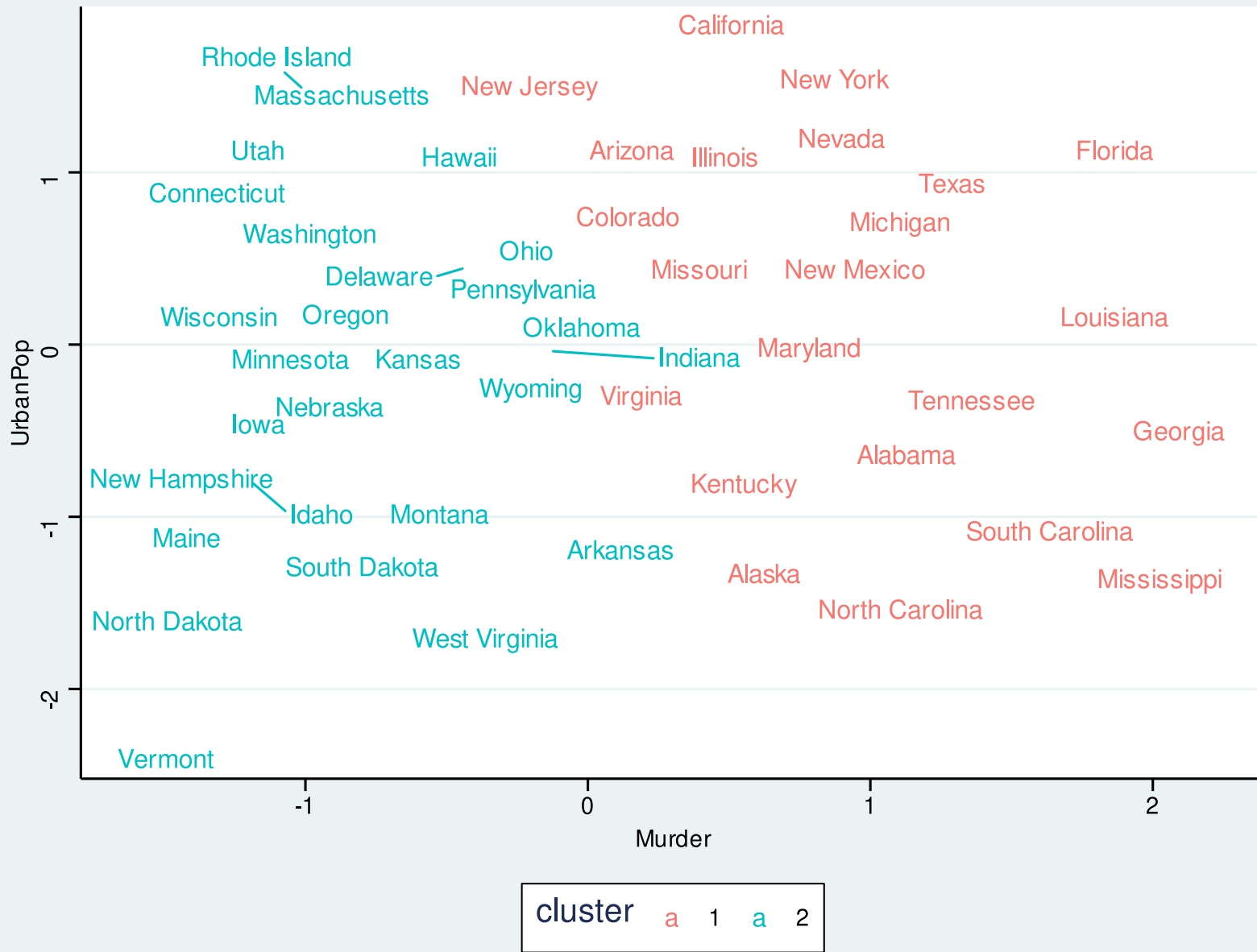
Glance at the sum of square decompositions

```
glance(k.means)
# A tibble: 1 × 4
  totss tot.withinss betweenss  iter
  <dbl>         <dbl>         <dbl> <int>
1     98          62.2          35.8     1
```

augment from broom package

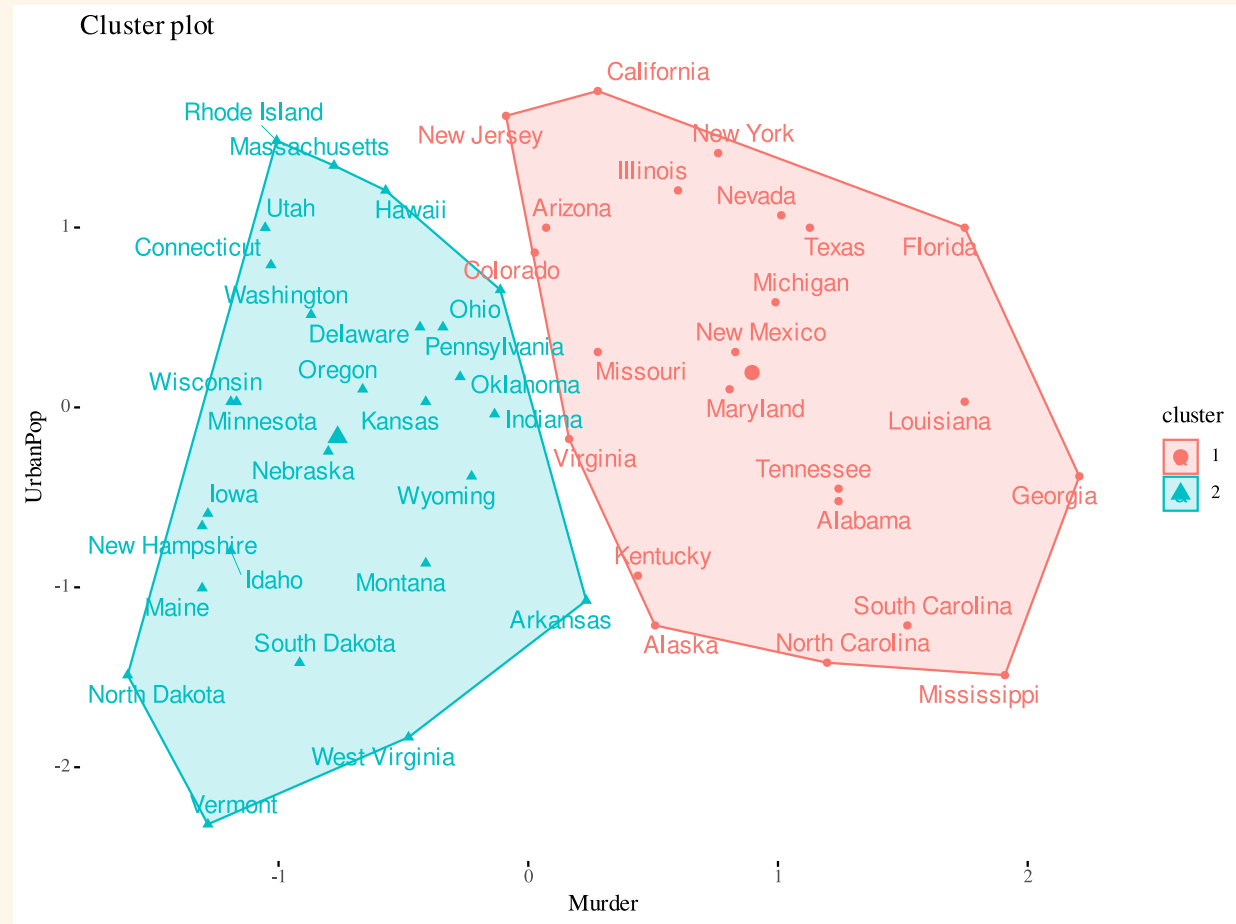
```
library(broom)
knitr::kable(augment(k.means, data = USAData))
```

.rownames	Murder	UrbanPop	.cluster
Alabama	1.2425641	-0.5209066	1
Alaska	0.5078625	-1.2117642	1
Arizona	0.0716334	0.9989801	1
Arkansas	0.2323494	-1.0735927	2
California	0.2782682	1.7589234	1
Colorado	0.0257146	0.8608085	1
Connecticut	-1.0304190	0.7917228	2

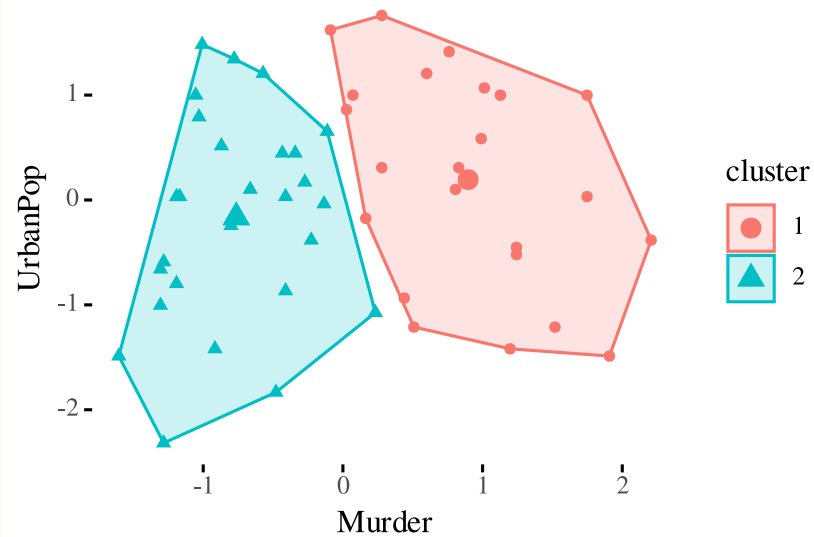


In-built function for visuals using factoextra

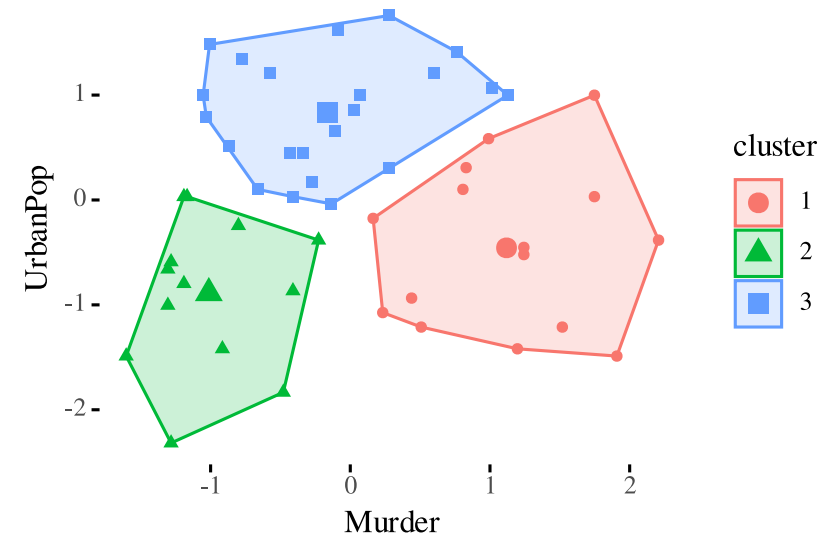
```
library(factoextra)
fviz_cluster(k.means, data = USADData, repel = TRUE, ggtheme = theme_tufte())
```



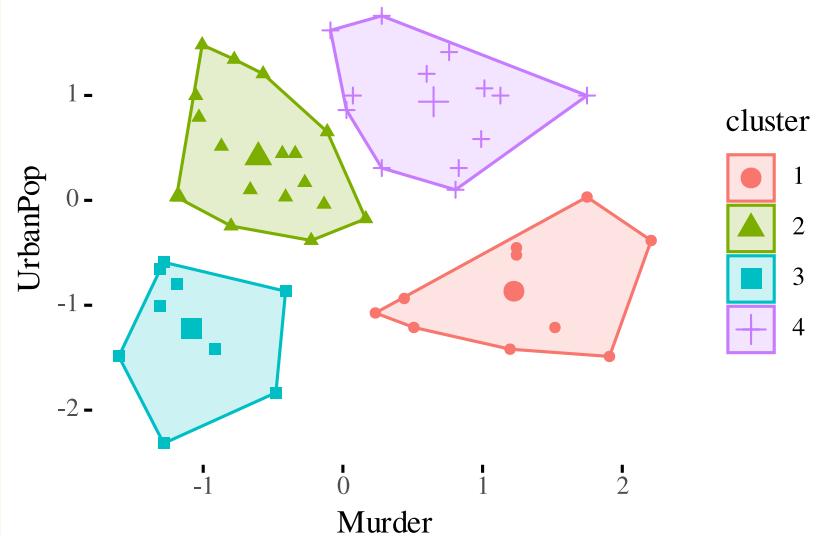
k = 2



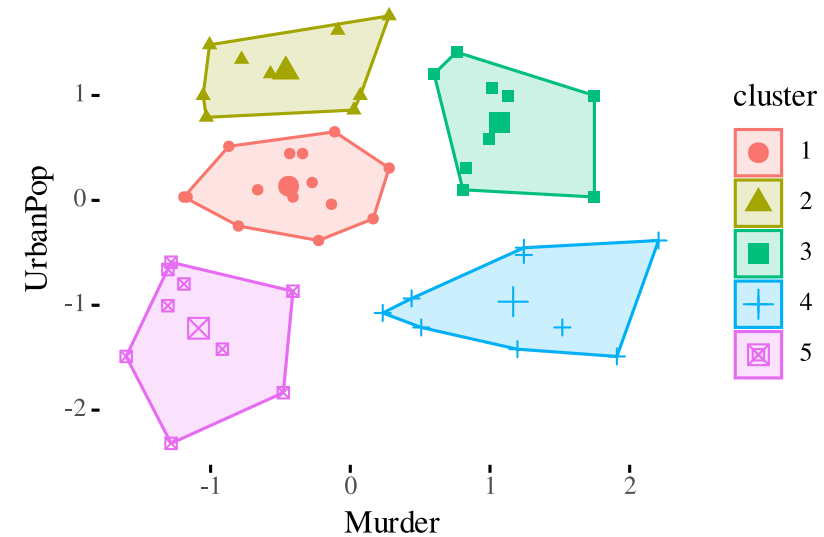
k = 3



k = 4



k = 5



Group Activity 1

10:00



- Get the class activity 26.Rmd file from [moodle](#)
- Let's work on group activity 1 together

Visuals do not tell all the story

Visuals tell us where the true delineations occur, but do not tell us what the optimal number of clusters is.

Determine the optimal number of clusters

```
set.seed(1234)
multi_kmeans <- tibble(k = 1:10) %>%
  mutate(
    model = purrr::map(k, ~ kmeans(USAData, centers = .x, nstart = 25)),
    tot.withinss = purrr::map_dbl(model, ~ glance(.x)$tot.withinss)
  )
```

```
multi_kmeans
```

```
# A tibble: 10 × 3
```

	k	model	tot.withinss
	<int>	<list>	<dbl>
1	1	<kmeans>	98
2	2	<kmeans>	62.4
3	3	<kmeans>	36.6
4	4	<kmeans>	24.9
5	5	<kmeans>	19.6
6	6	<kmeans>	16.4
7	7	<kmeans>	13.7
8	8	<kmeans>	11.0
9	9	<kmeans>	9.85
10	10	<kmeans>	8.04

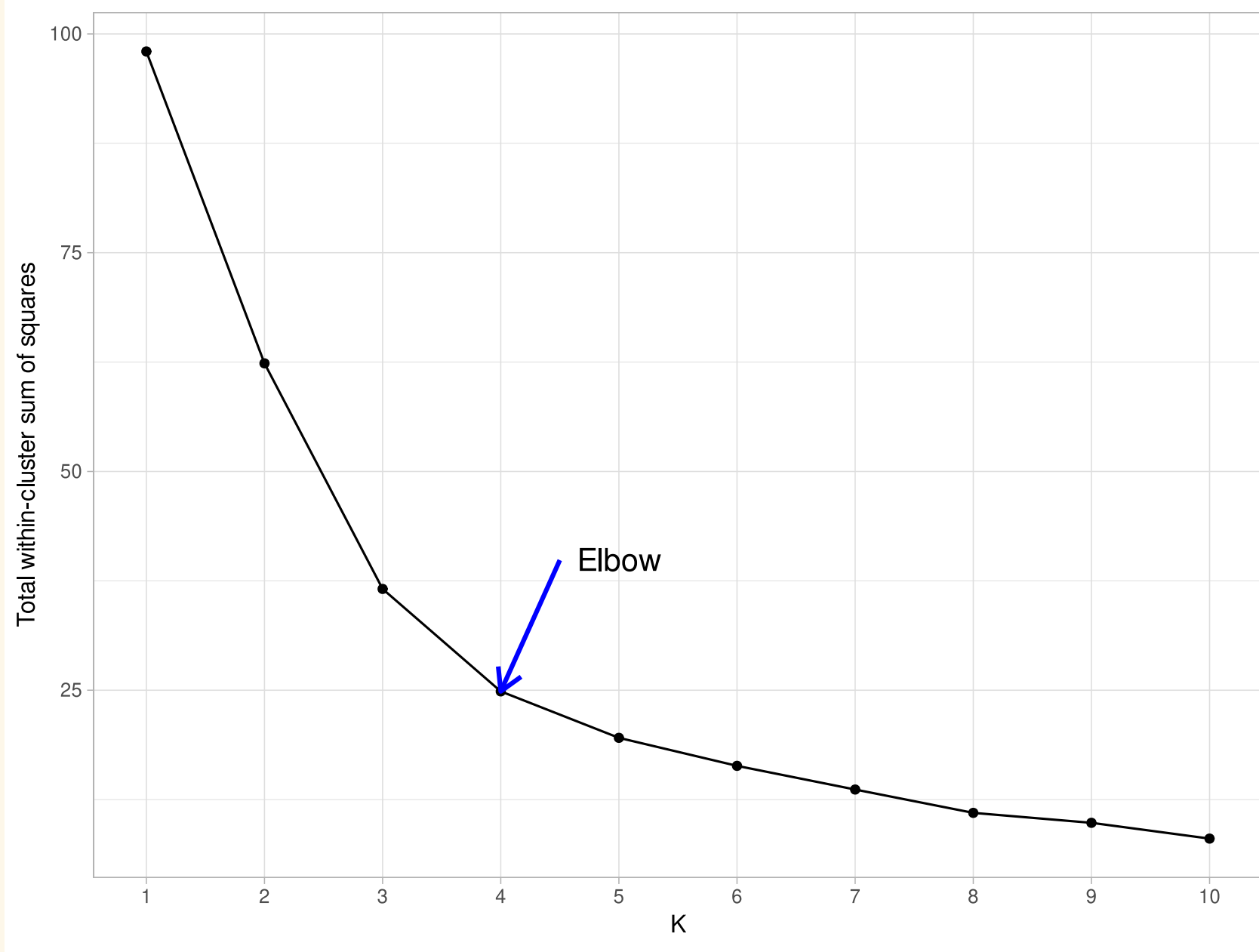
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  mutate(
    model = purrr::map(k, ~ kmeans(USAData, centers = .x, nstart = 25)),
    tot.withinss = purrr::map_dbl(model, ~ glance(.x)$tot.withinss)
  )
```

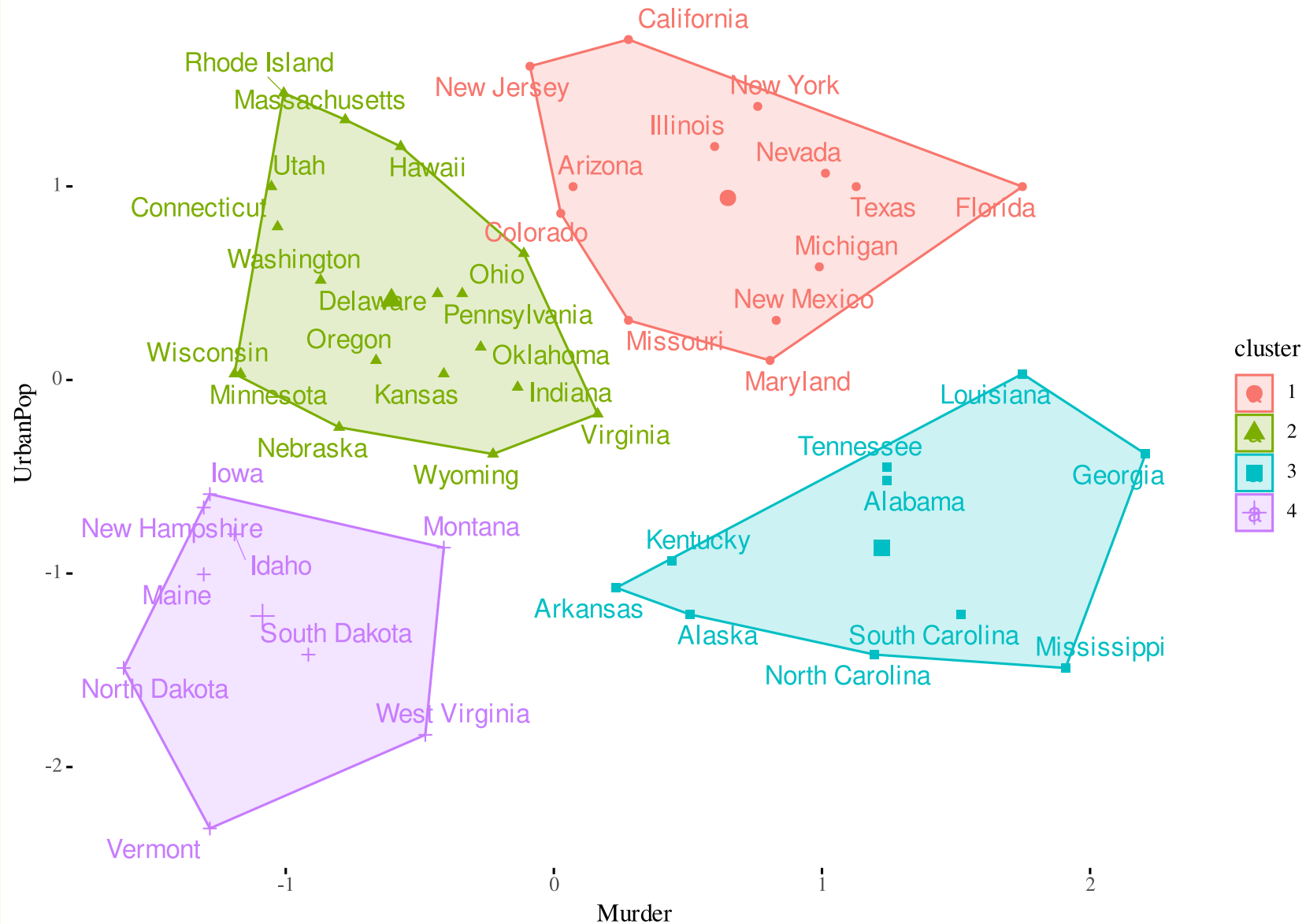
```
multi_kmeans
```

```
# A tibble: 10 × 3
```

	k	model	tot.withinss
	<int>	<list>	<dbl>
1	1	<kmeans>	98
2	2	<kmeans>	62.4
3	3	<kmeans>	36.6
4	4	<kmeans>	24.9
5	5	<kmeans>	19.6
6	6	<kmeans>	16.4
7	7	<kmeans>	13.7
8	8	<kmeans>	11.0
9	9	<kmeans>	9.85
10	10	<kmeans>	8.04



Cluster plot



Extract the centroids

```
USDataData %>%  
  mutate(Cluster = kmeans.final$cluster) %>%  
  group_by(Cluster) %>%  
  summarise_all("mean")  
# A tibble: 4 × 3  
  Cluster Murder UrbanPop  
    <int>   <dbl>    <dbl>  
1         1  0.649     0.941  
2         2 -0.606     0.412  
3         3  1.22     -0.866  
4         4 -1.09     -1.22
```

Group Activity 2

10:00



- Please continue working on the remainder of the group activities