

# DSC\_520\_week10\_Assignment01

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## Load libraries as needed

```
if(!require('factoextra')) {  
  install.packages("factoextra", repos="http://cran.us.r-project.org")  
  library('factoextra')  
}
```

## Loading required package: factoextra

## Warning: package 'factoextra' was built under R version 4.2.1

## Loading required package: ggplot2

## Warning: package 'ggplot2' was built under R version 4.2.1

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

```
if(!require('cluster')) {  
  install.packages("cluster", repos="http://cran.us.r-project.org")  
  library('cluster')  
}
```

## Loading required package: cluster

## Warning: package 'cluster' was built under R version 4.2.1

```
if(!require('NbClust')) {  
  install.packages("NbClust", repos="http://cran.us.r-project.org")  
  library('NbClust')  
}
```

## Loading required package: NbClust

```
if(!require('e1071')) {  
  install.packages("e1071", repos="http://cran.us.r-project.org")  
  library('e1071')  
}
```

## Loading required package: e1071

## Warning: package 'e1071' was built under R version 4.2.1

```
if(!require('caTools')) {  
  install.packages("caTools", repos="http://cran.us.r-project.org")  
  library('caTools')  
}
```

## Loading required package: caTools

## Warning: package 'caTools' was built under R version 4.2.1

```
if(!require('class')) {
  install.packages("class", repos="http://cran.us.r-project.org")
  library('class')
}
```

```
## Loading required package: class
```

```
## Warning: package 'class' was built under R version 4.2.1
```

## Load data

```
## Set the working directory to the root of your DSC 520 directory
setwd("C:/Users/chris/dsc520/data")
## Load the `data/data/clustering-data.csv` to
df <- read.csv("C:/Users/chris/dsc520/data/clustering-data.csv")
head(df)
```

```
##      x    y
## 1  46 236
## 2  69 236
## 3 144 236
## 4 171 236
## 5 194 236
## 6 195 236
```

## Scale the value of the dataframe

```
df_scale <- scale(df) # scale the value of the dataframe
head(df_scale)
```

```
##           x           y
## [1,] -0.8482235 1.561107
## [2,] -0.5415045 1.561107
## [3,]  0.4586659 1.561107
## [4,]  0.8187273 1.561107
## [5,]  1.1254462 1.561107
## [6,]  1.1387818 1.561107
```

## Displays the number of rows and columns

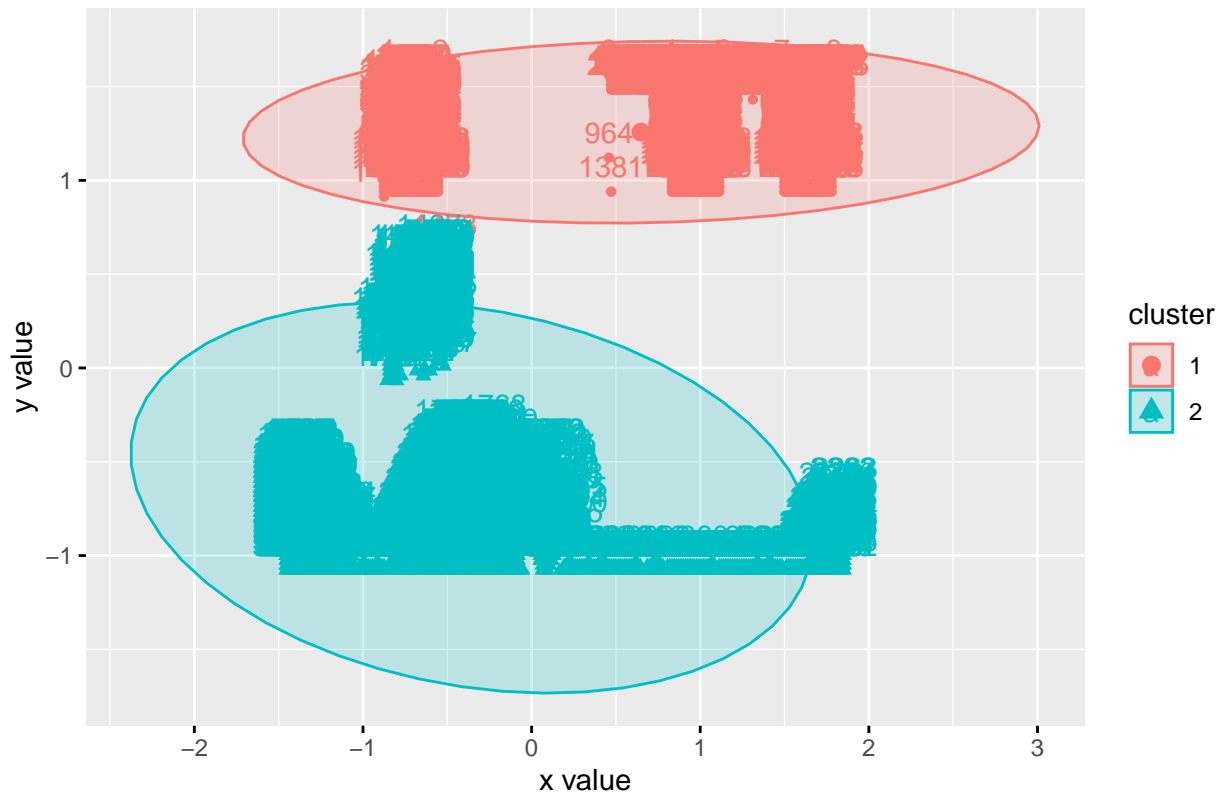
```
dim(df_scale) # displays the number of rows and columns
```

```
## [1] 4022    2
```

Find the optimal cluster without any method randomly from  $k = 2$  to  $k = 12$

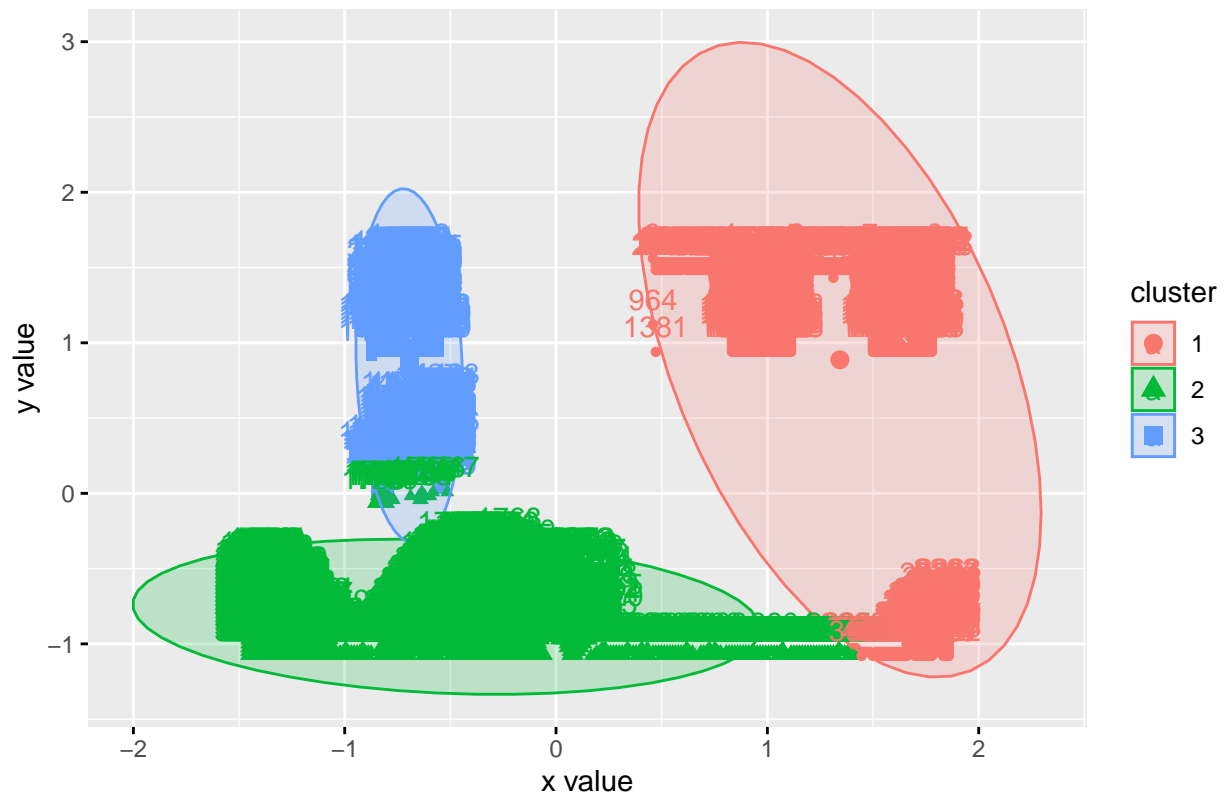
```
# Compute k-means with k = 2
set.seed(123)
km.res <- kmeans(df_scale, 2, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



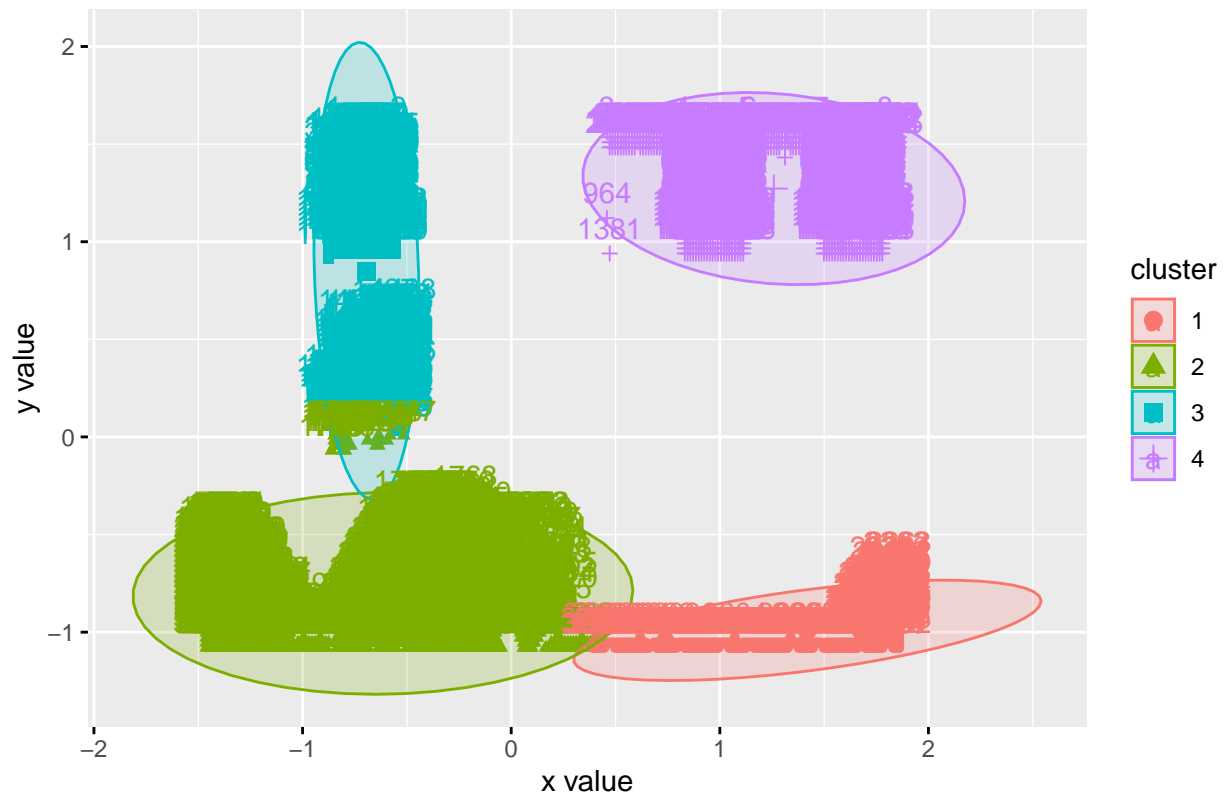
```
# Compute k-means with k = 3
set.seed(123)
km.res <- kmeans(df_scale, 3, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



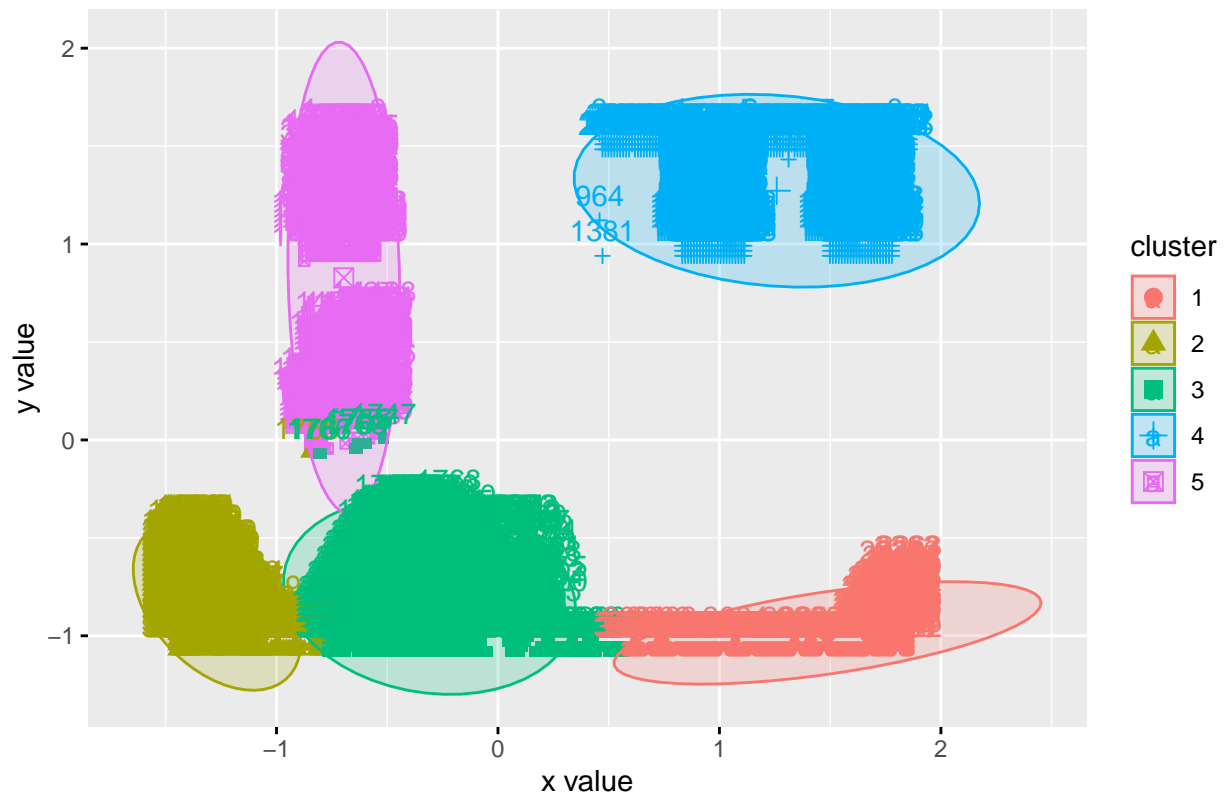
```
# Compute k-means with k = 4
set.seed(123)
km.res <- kmeans(df_scale, 4, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



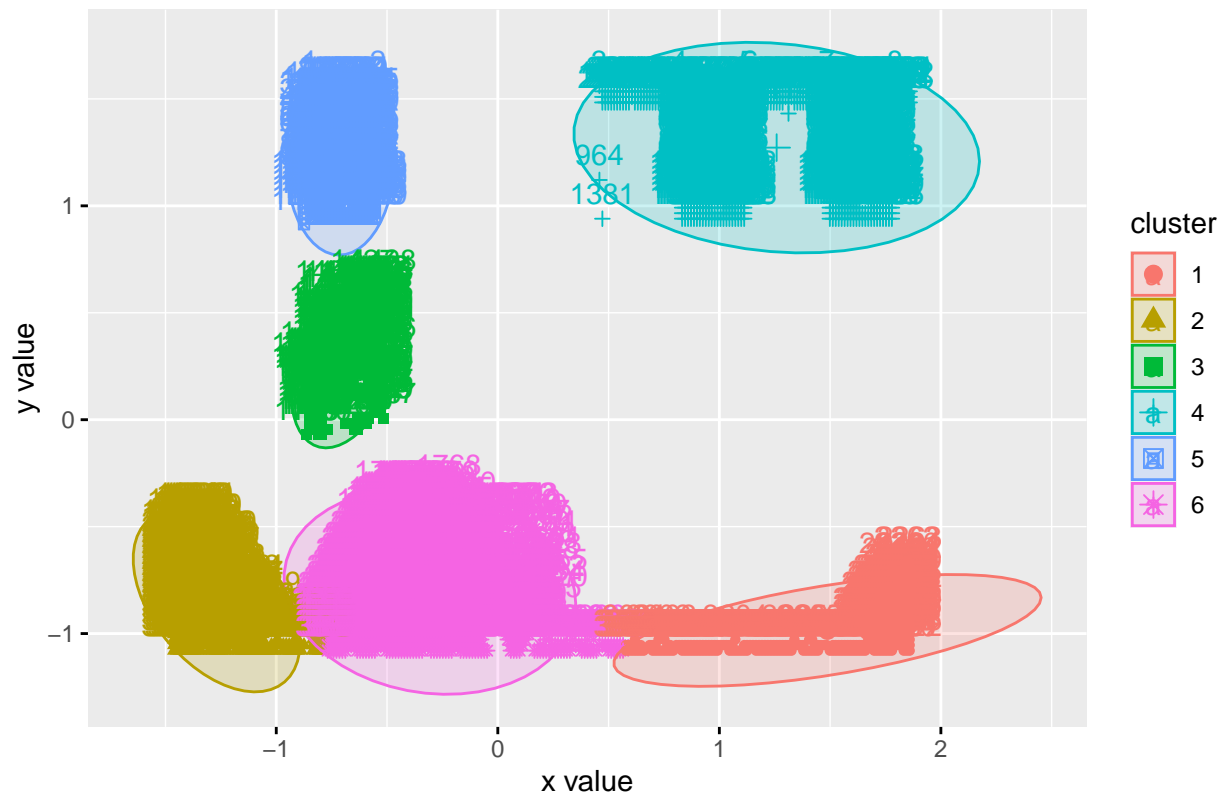
```
# Compute k-means with k = 5
set.seed(123)
km.res <- kmeans(df_scale, 5, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



```
# Compute k-means with k = 6
set.seed(123)
km.res <- kmeans(df_scale, 6, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

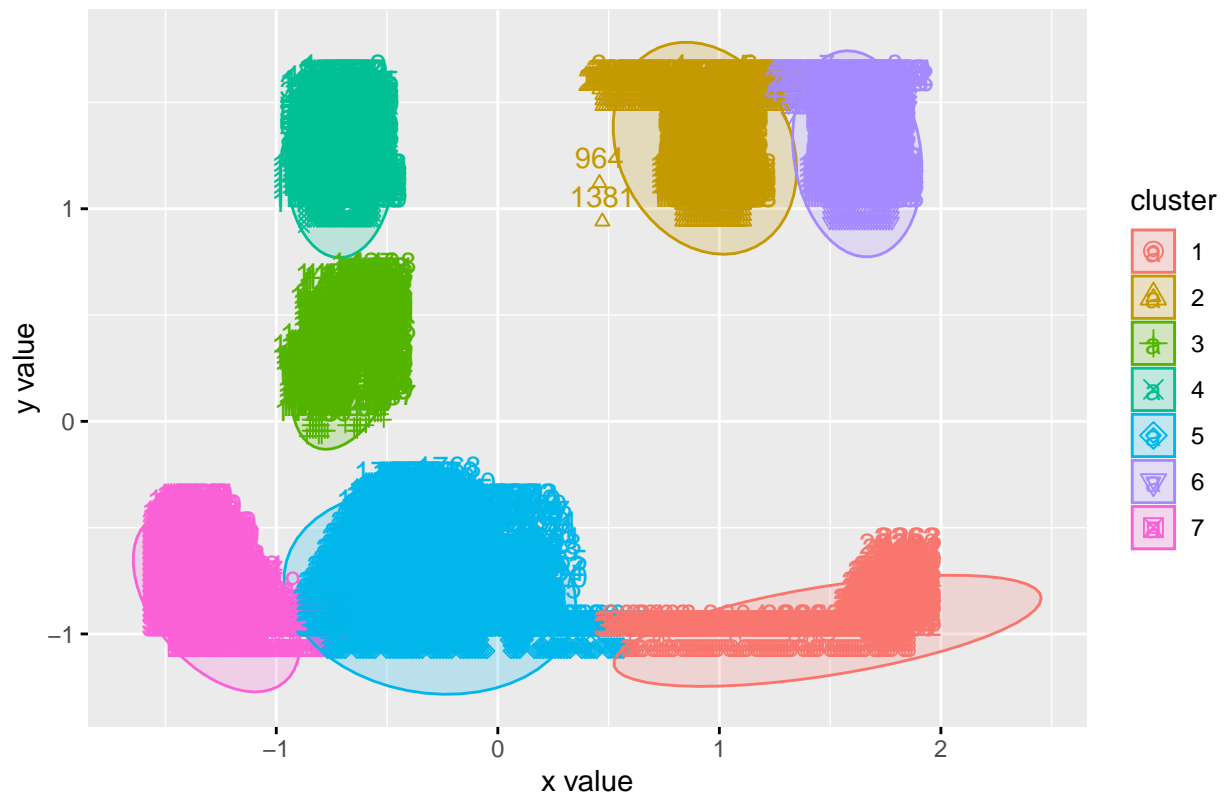
Cluster plot



```
# Compute k-means with k = 7
set.seed(123)
km.res <- kmeans(df_scale, 7, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

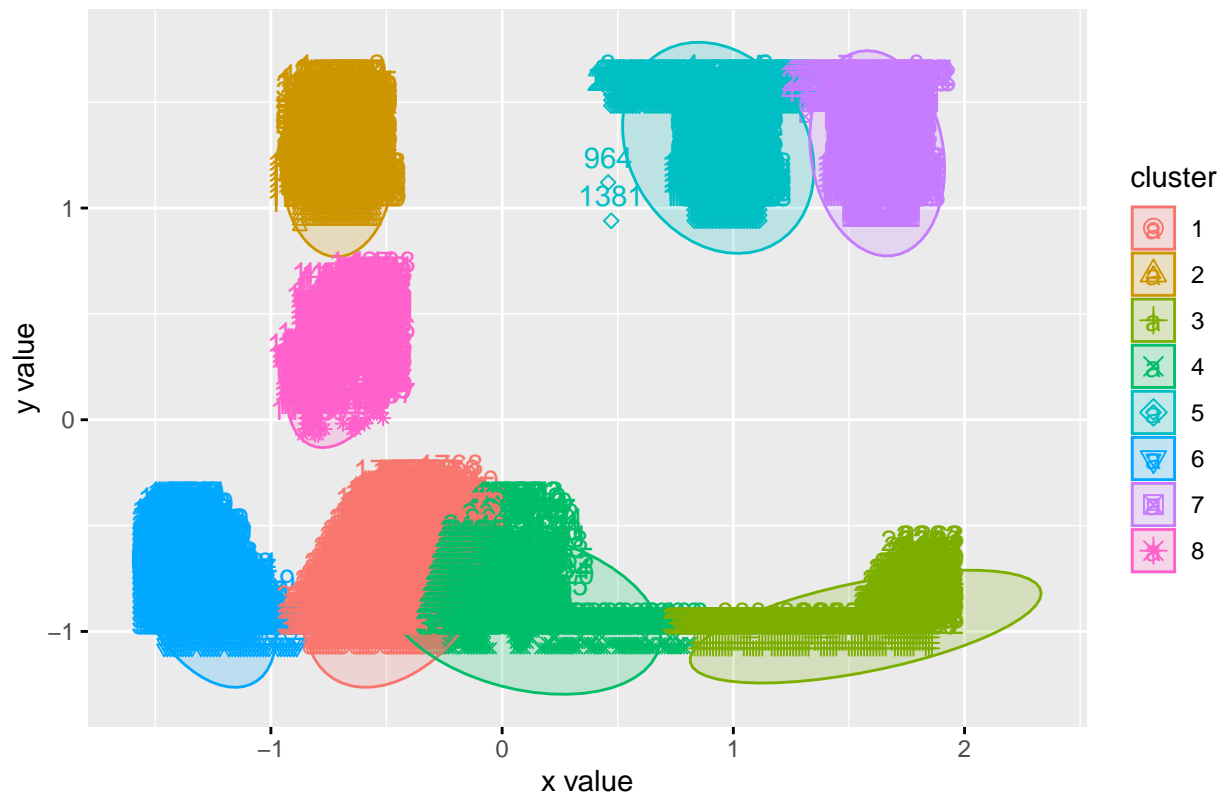


Cluster plot



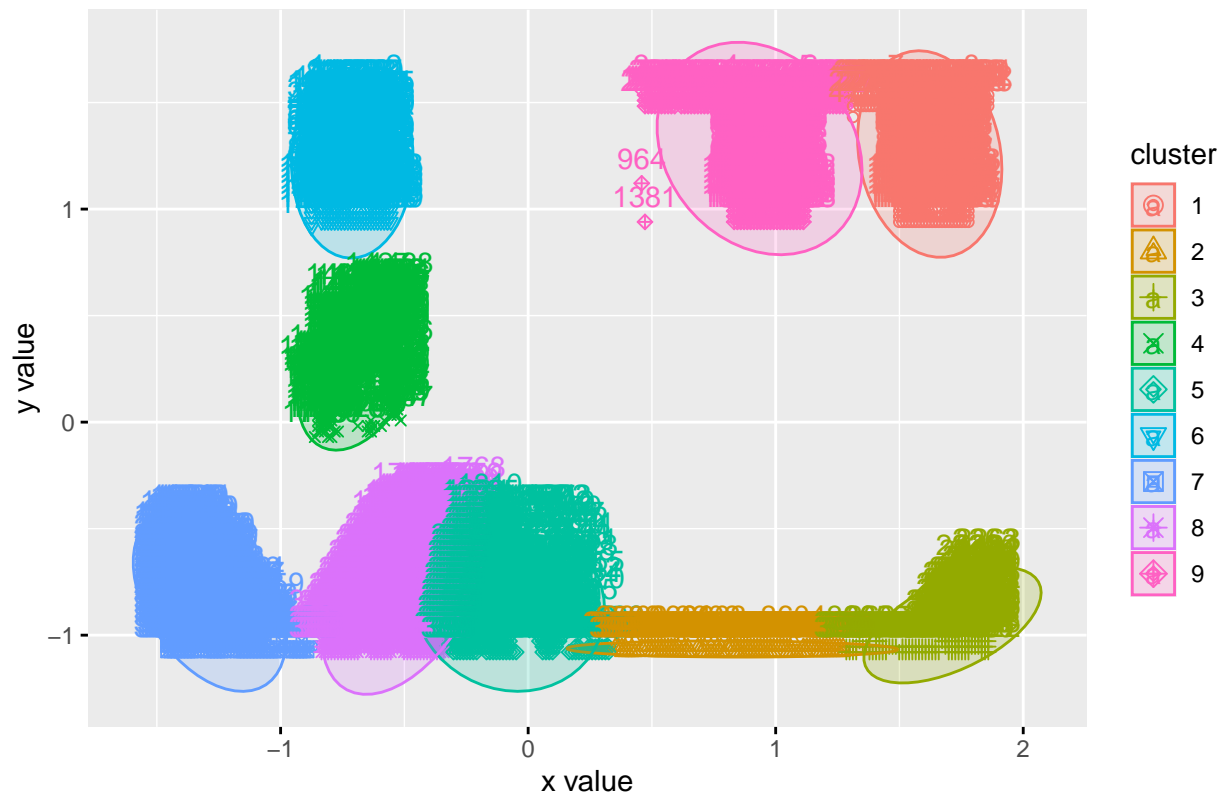
```
# Compute k-means with k = 8
set.seed(123)
km.res <- kmeans(df_scale, 8, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



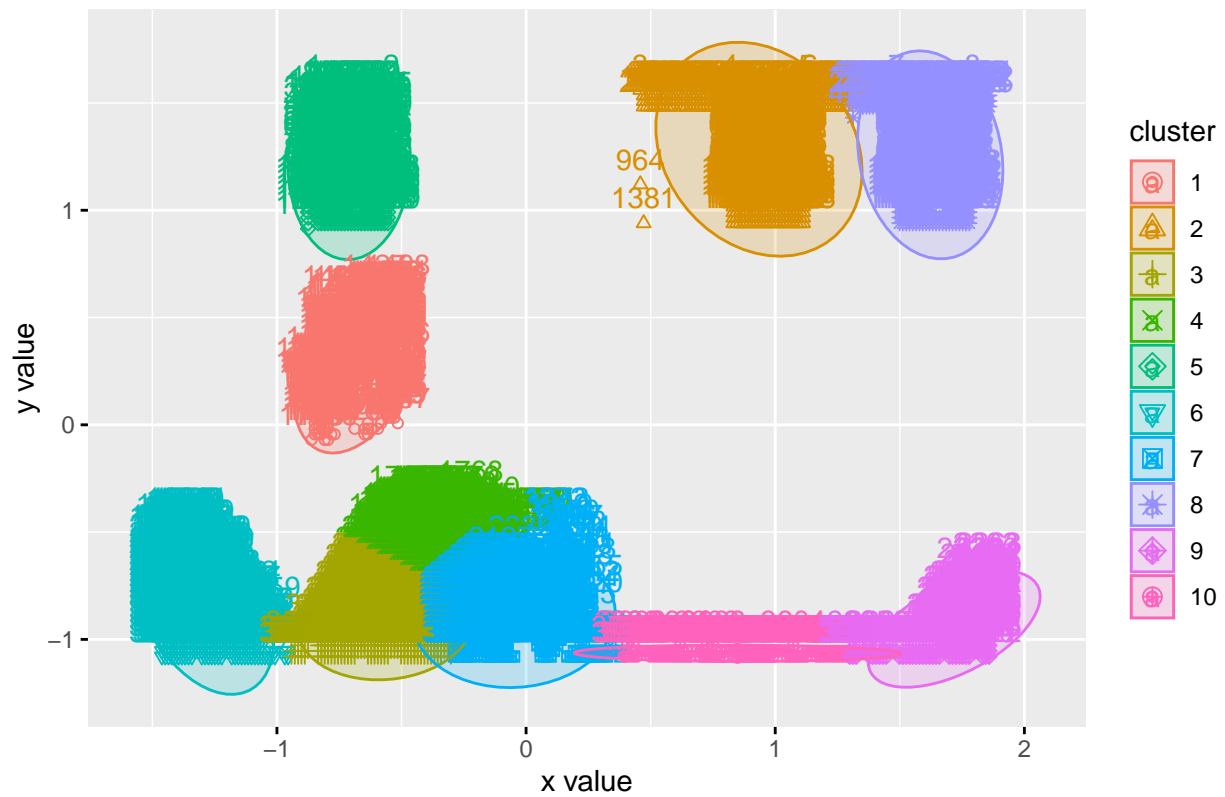
```
# Compute k-means with k = 9
set.seed(123)
km.res <- kmeans(df_scale, 9, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



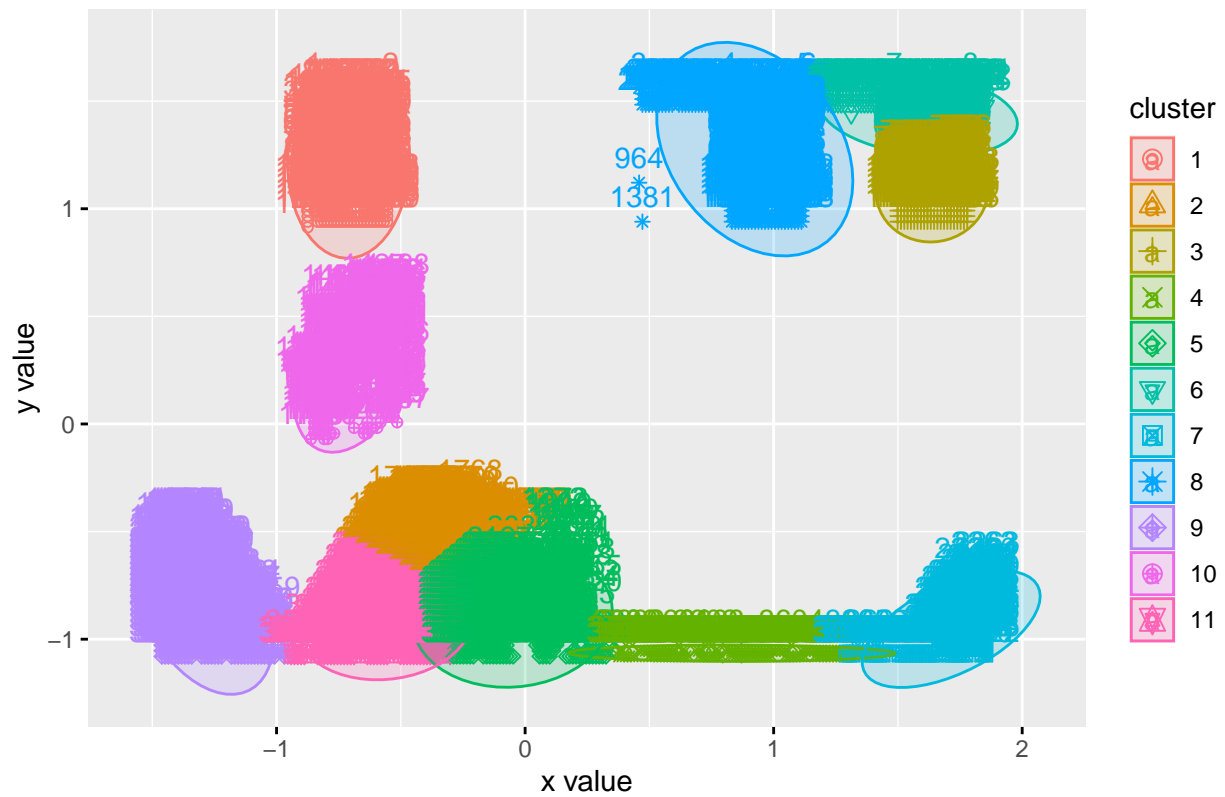
```
# Compute k-means with k = 10
set.seed(123)
km.res <- kmeans(df_scale, 10, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



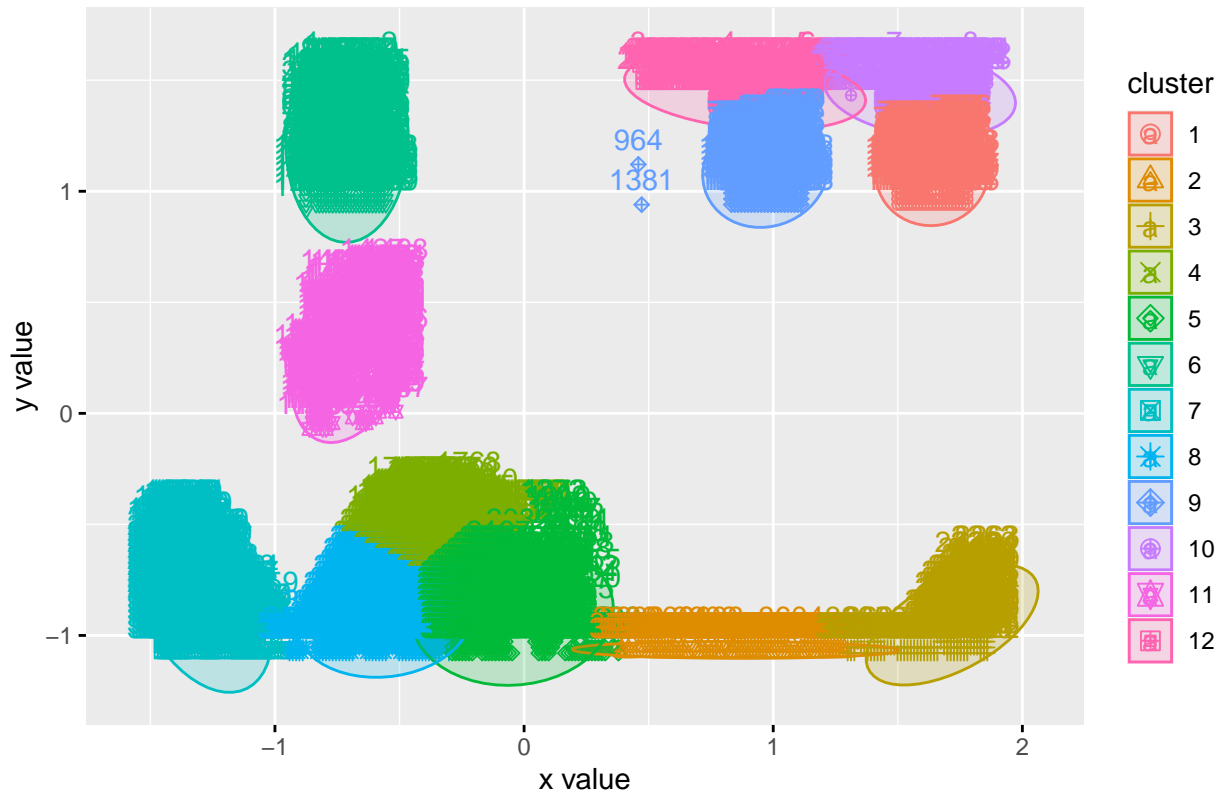
```
# Compute k-means with k = 11
set.seed(123)
km.res <- kmeans(df_scale, 11, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

Cluster plot



```
# Compute k-means with k = 12
set.seed(123)
km.res <- kmeans(df_scale, 12, nstart = 25)
fviz_cluster(km.res, df_scale, ellipse.type = "norm")
```

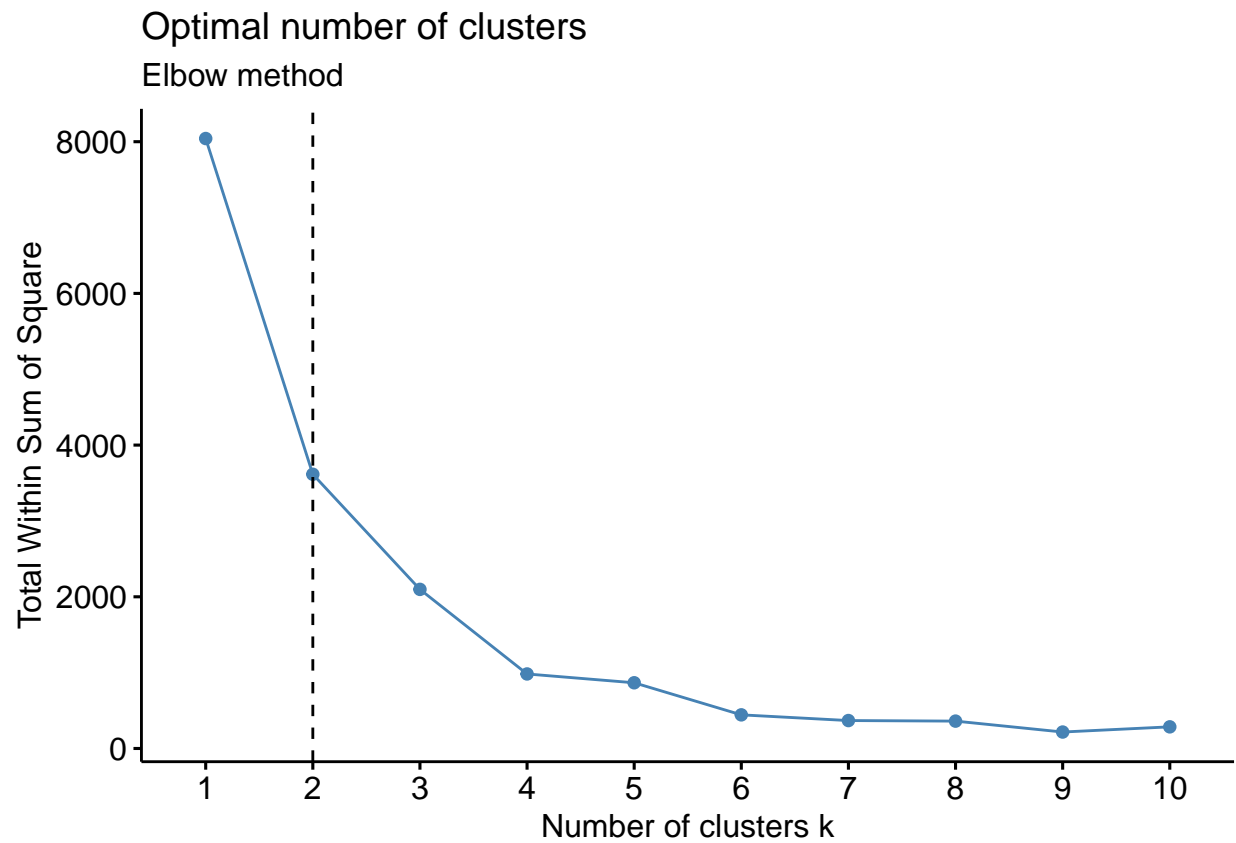
## Cluster plot



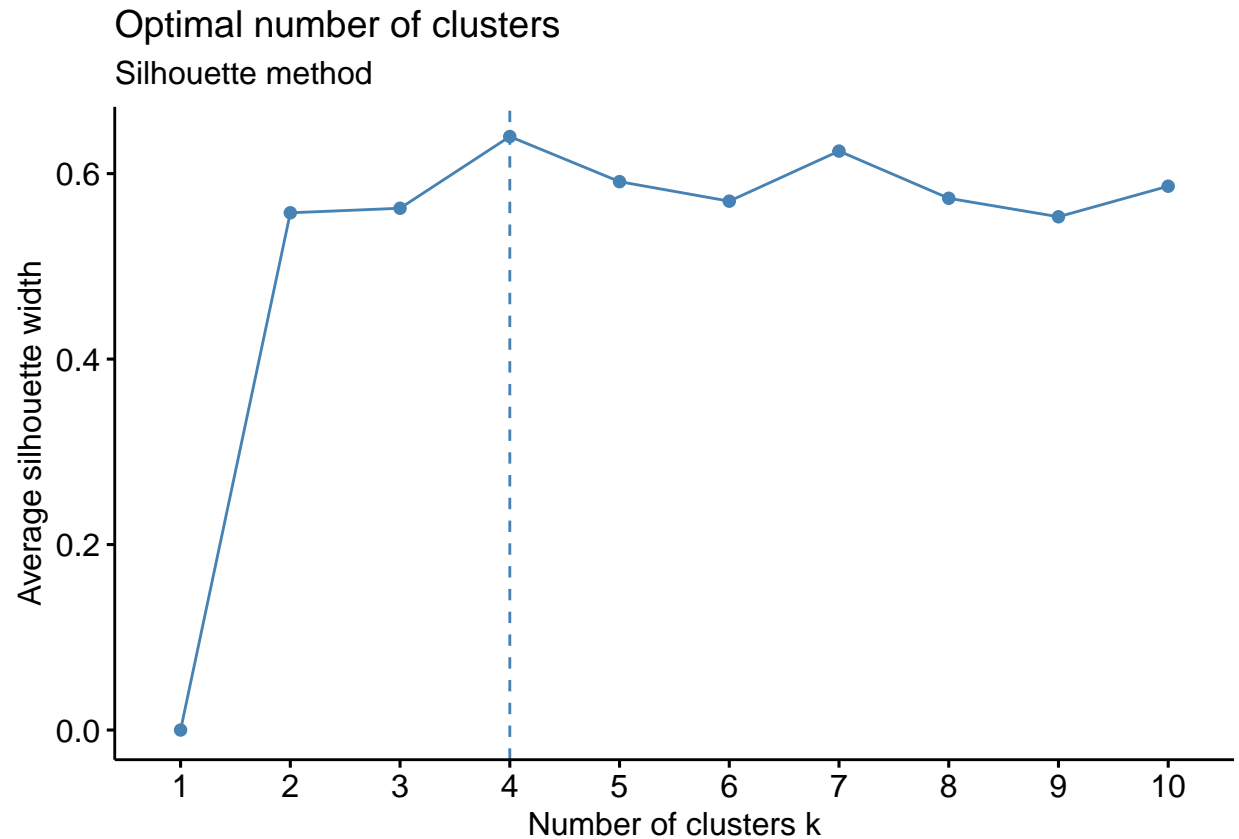
Find the optimal cluster using Elbow, Silhouette, Gap statistic method

```
library(factoextra)
library(NbClust)

# Elbow method
fviz_nbclust(df_scale, kmeans, method = "wss") +
  geom_vline(xintercept = 2, linetype = 2) +
  labs(subtitle = "Elbow method")
```



```
# Silhouette method  
fviz_nbclust(df_scale, kmeans, method = "silhouette")+  
  labs(subtitle = "Silhouette method")
```

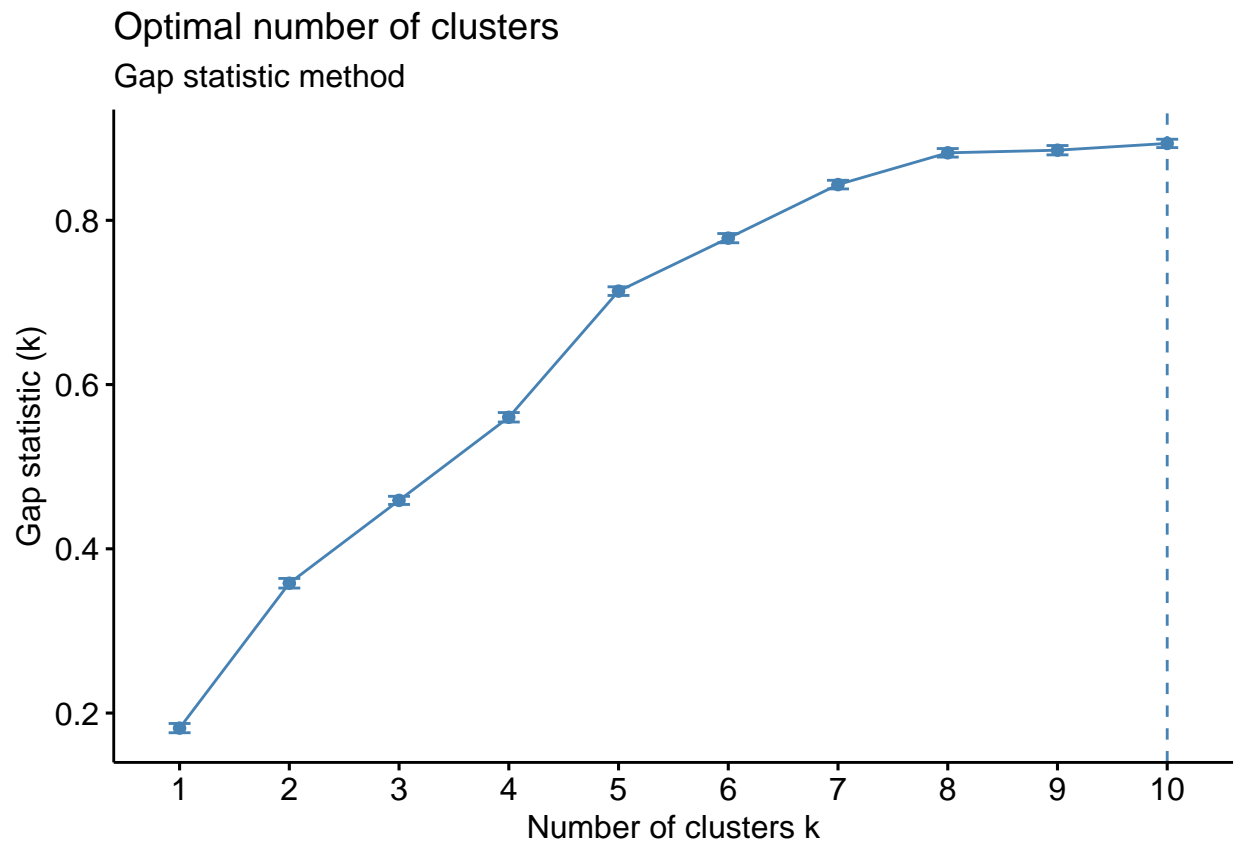


```
# Gap statistic
set.seed(123)
fviz_nbclust(df_scale, kmeans, nstart = 25, method = "gap_stat", nboot = 50)+
  labs(subtitle = "Gap statistic method")
```

```
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
```



```
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
## Warning: did not converge in 10 iterations
```



*##CONCLUSION: Using different method to identify the optimal cluster. Elbow and Shilhouette method both*

## Clustering mean and vector

```
set.seed(1)

#perform k-means clustering with k = 4 clusters
km <- kmeans(df_scale, centers = 4, nstart = 25)

#view results
km
```

[illegible]

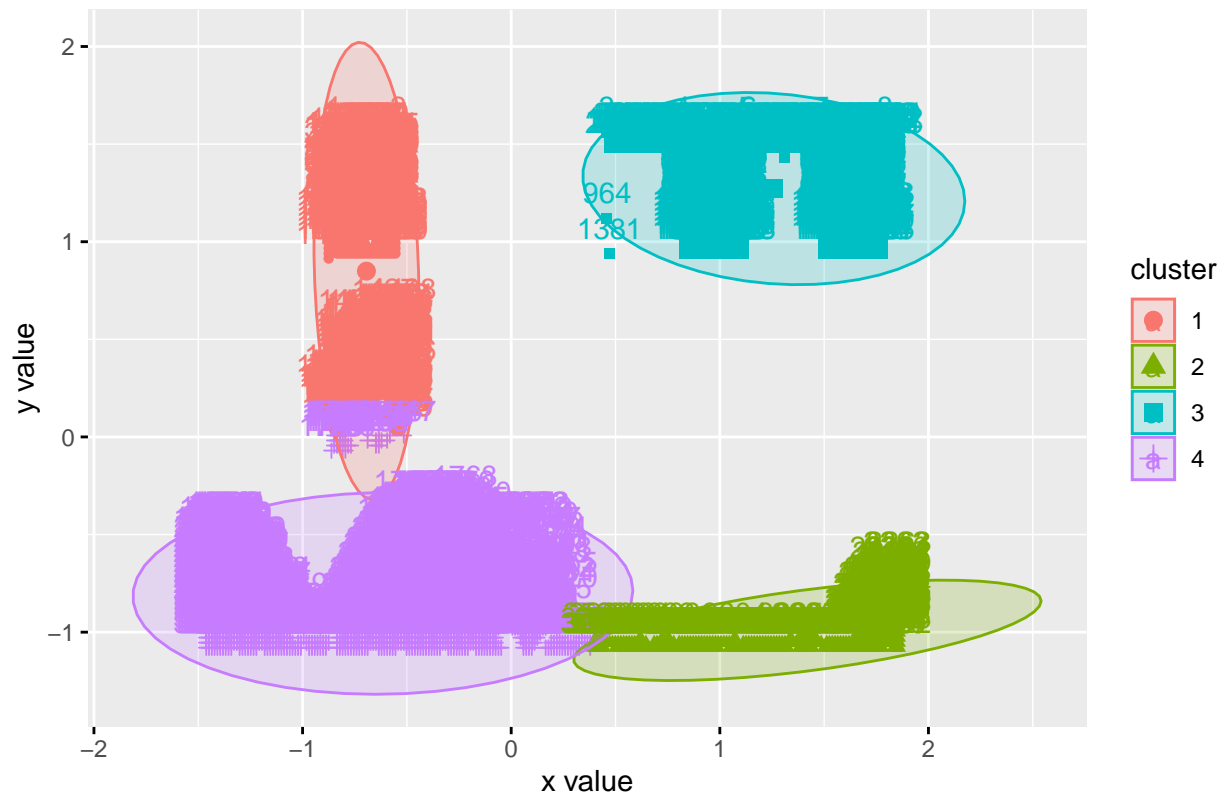
[illegible]

[illegible]

### Plot clusters by using the fviz\_cluster() function

```
#plot results of final k-means model
fviz_cluster(km, df_scale, ellipse.type = "norm")
```

## Cluster plot



## Mean of each cluster

```
#find means of each cluster
aggregate(df, by=list(cluster=km$cluster), mean)
```

```
##   cluster      x      y
## 1      1  57.57162 208.5106
## 2      2 216.05917 137.4408
## 3      3 203.98579 224.8406
## 4      4  63.48175 144.7512
```

## Add clusters to the original data

```
#add cluster assignment to original data
final_data <- cbind(df, cluster = km$cluster)

#view final data
head(final_data)
```

```
##      x  y cluster
## 1  46 236      1
```

##	2	69	236	1
##	3	144	236	3
##	4	171	236	3
##	5	194	236	3
##	6	195	236	3