DSC520 Week10 Exercise 11.2.1

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```
# Load the packages
library(tidyverse)
## -- Attaching packages ----- tidyverse 1.3.1 --
## v ggplot2 3.3.5 v purrr 0.3.4

## v tibble 3.1.6 v dplyr 1.0.7

## v tidyr 1.1.4 v stringr 1.4.0

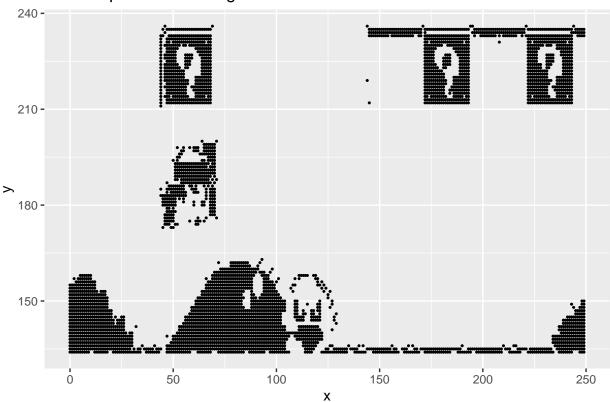
## v readr 2.1.1 v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
library(cluster)
library(factoextra)
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
library(caTools)
setwd("/Users/Jagadeesh/Documents/GitHub/dsc520")
# Load Clustering Data Dataset
cluster_df <- read.csv("data/clustering-data.csv")</pre>
\# Check structure of clustering_data_df
str(cluster_df)
## 'data.frame': 4022 obs. of 2 variables:
## $ x: int 46 69 144 171 194 195 221 244 45 47 ...
## $ y: int 236 236 236 236 236 236 236 235 235 ...
{\it \# Check sample rows of clustering\_data\_df}
head(cluster_df)
##
```

1 46 236

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

```
# i.Plot the data set using a scatter plot.
library(ggplot2)
ggplot(data=cluster_df,aes(x=x,y=y)) + geom_point(size=0.4) + ggtitle("Scatter-plot of clustering data"
```

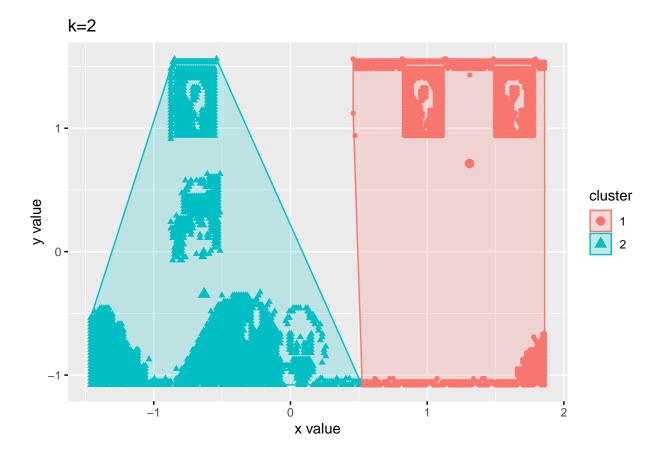
Scatter-plot of clustering data



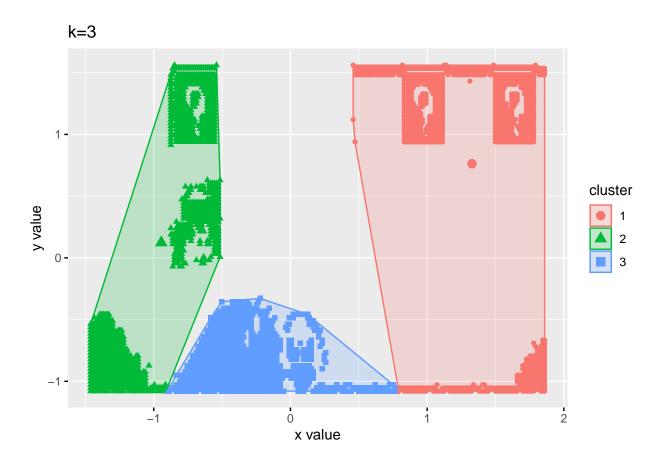
```
# ii.Fit the dataset using the k-means algorithm from k=2 to k=12. Create a scatter plot of the resulta
set.seed(123)
kmeans_2 <- kmeans(cluster_df, 2, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_3 <- kmeans(cluster_df, 3, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_4 <- kmeans(cluster_df, 4, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_5 <- kmeans(cluster_df, 5, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_6 <- kmeans(cluster_df, 6, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_7 <- kmeans(cluster_df, 7, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_8 <- kmeans(cluster_df, 8, iter.max = 300, nstart = 10)</pre>
```

```
set.seed(123)
kmeans_9 <- kmeans(cluster_df, 9, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_10 <- kmeans(cluster_df, 10, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_11 <- kmeans(cluster_df, 11, iter.max = 300, nstart = 10)
set.seed(123)
kmeans_12 <- kmeans(cluster_df, 12, iter.max = 300, nstart = 10)

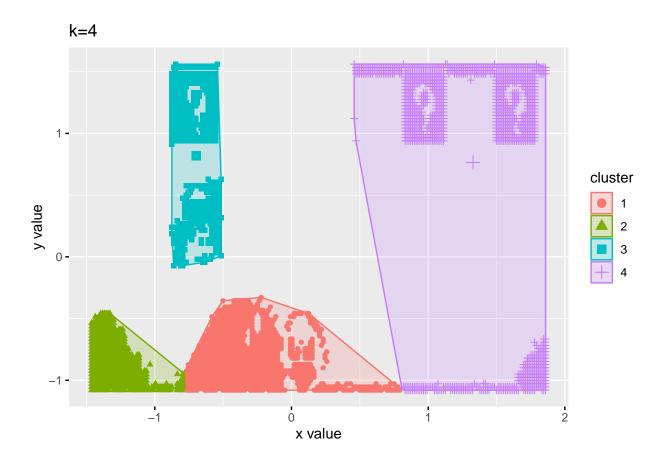
# Plots to compare
fviz_cluster(kmeans_2, geom="point",data=cluster_df) + ggtitle("k=2")</pre>
```



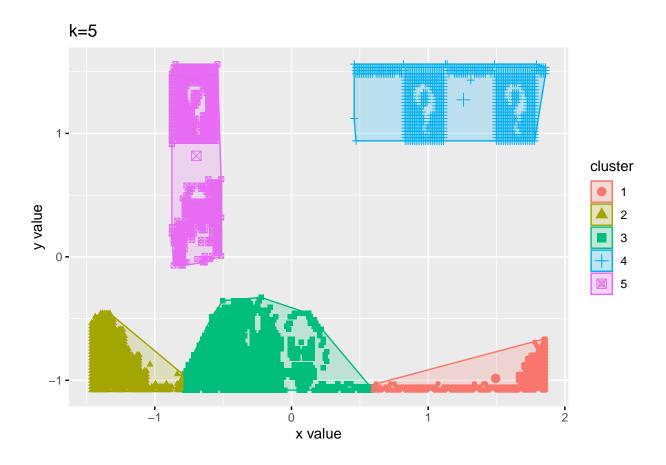
fviz_cluster(kmeans_3, geom="point",data=cluster_df) + ggtitle("k=3")



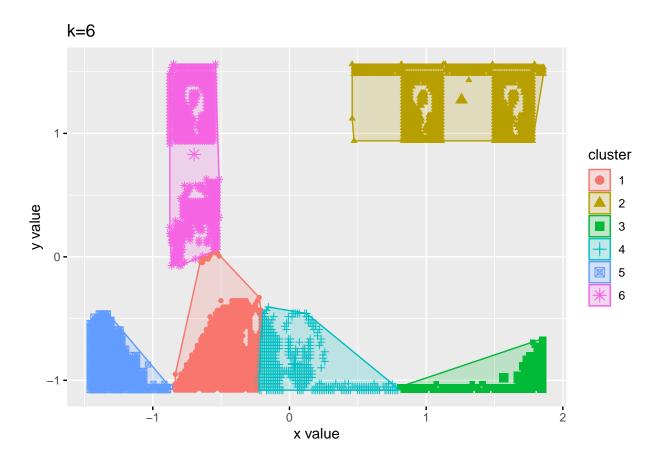
fviz_cluster(kmeans_4, geom="point",data=cluster_df) + ggtitle("k=4")



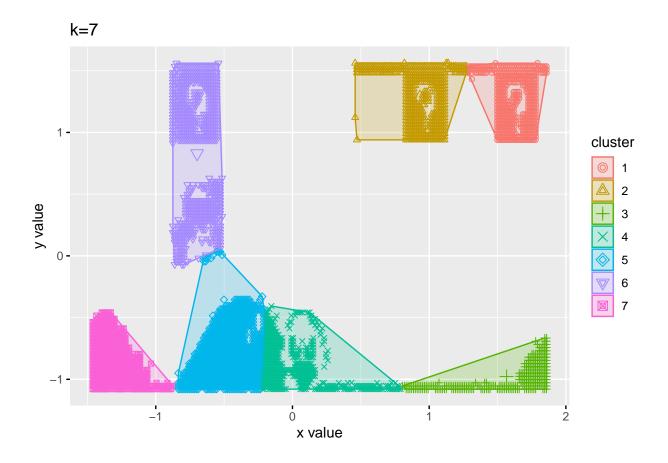
fviz_cluster(kmeans_5, geom="point",data=cluster_df) + ggtitle("k=5")



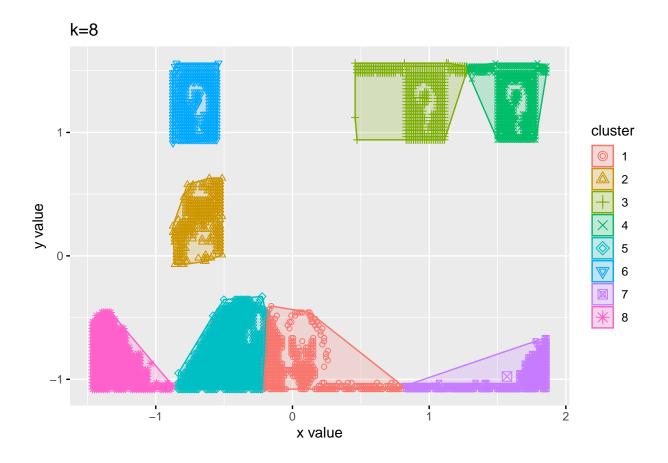
fviz_cluster(kmeans_6, geom="point",data=cluster_df) + ggtitle("k=6")



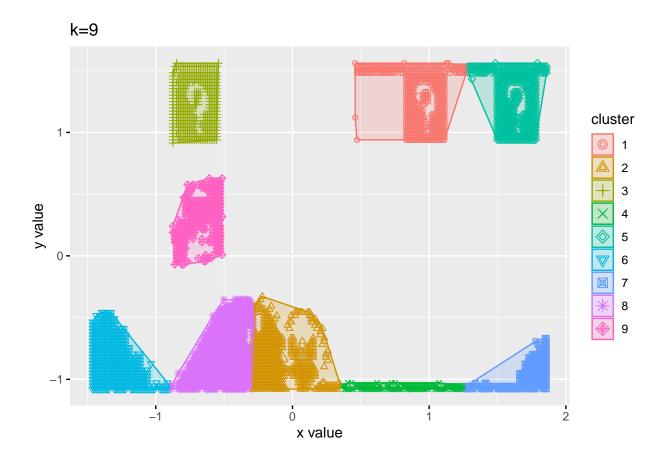
fviz_cluster(kmeans_7, geom="point",data=cluster_df) + ggtitle("k=7")



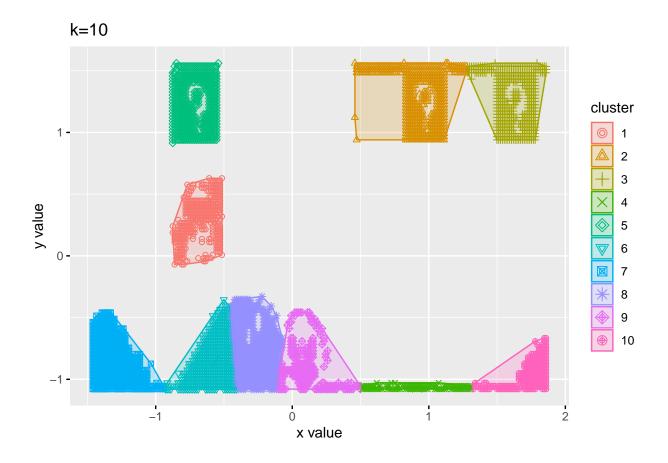
fviz_cluster(kmeans_8, geom="point",data=cluster_df) + ggtitle("k=8")



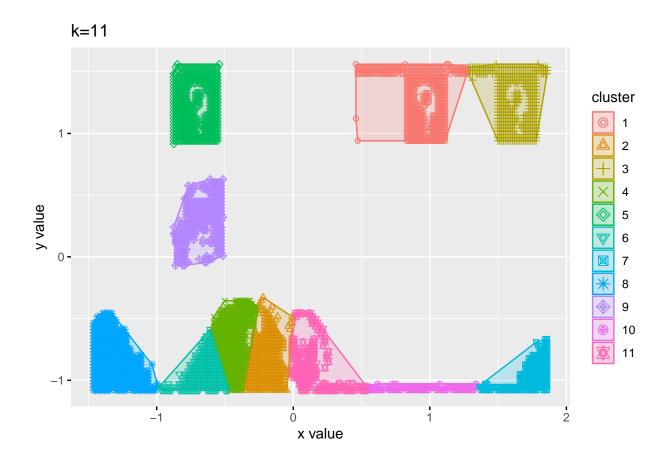
fviz_cluster(kmeans_9, geom="point",data=cluster_df) + ggtitle("k=9")



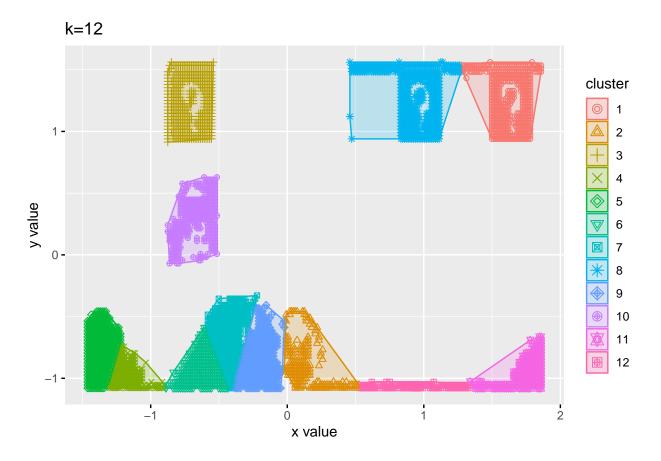
fviz_cluster(kmeans_10, geom="point",data=cluster_df) + ggtitle("k=10")

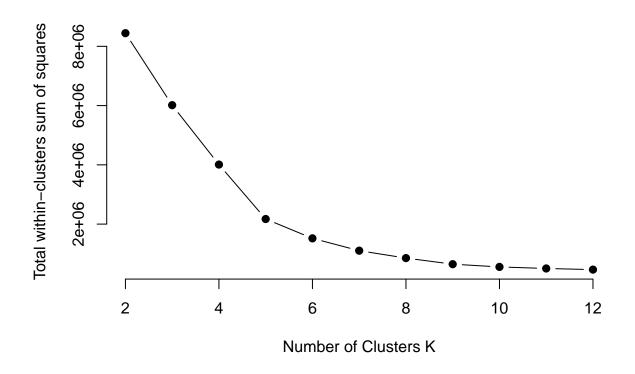


fviz_cluster(kmeans_11, geom="point",data=cluster_df) + ggtitle("k=11")



fviz_cluster(kmeans_12, geom="point",data=cluster_df) + ggtitle("k=12")





The results suggest that 6 is the optimal number of clusters as it appears to ## be the bend of the elbow curve.