Week 8 Assignment

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```
10.7~\#3
  • a)
obs <- data.frame(</pre>
 X1 = c(1, 1, 0, 5, 6, 4),
 X2 = c(4, 3, 4, 1, 2, 0)
obs %>%
  ggplot(aes(x = X1, y = X2)) +
geom_point()
  3 -
X 2-
  1 -
  0 -
                                  2
                                              X1
  • b)
# set random number seed for replicable results
set.seed(2017)
# assign the labels
obs$cluster <- sample(rep(c(1, 2), each = 3), 6, replace = FALSE)
```

```
# print the labels
obs
##
     X1 X2 cluster
## 1 1 4
## 2 1 3
                 1
## 3 0 4
## 4 5 1
                 1
## 5 6 2
                 2
## 6 4 0
# plot the labels
obs %>%
  dplyr::select(X1, X2, cluster) %>%
  ggplot(aes(x = X1, y = X2, color = as.factor(cluster))) +
  geom_point()
   4 -
   3 -
                                                                       as.factor(cluster)
X 2-
                                                                        • 1
                                                                        • 2
   1 -
   0 -
                          2
       0
                                   X1
      c)
centroids <-
  obs %>%
  dplyr::group_by(cluster) %>%
  dplyr::summarise(centroid_X1 = round(mean(X1), 2), centroid_X2 = round(mean(X2), 2)) %>%
  as.data.frame()
centroids
## cluster centroid_X1 centroid_X2
```

1

1

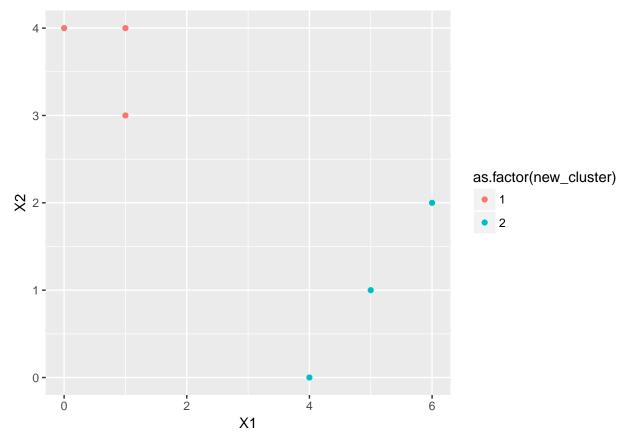
2.00

```
## 2
           2
                    3.67
                                 2.00
     d)
# create distance from cluster1 centroid
obs$dist from cluster1 centroid <- round(</pre>
  sqrt((obs$X1 - centroids$centroid_X1[centroids$cluster == 1])^2 +
       (obs$X2 - centroids$centroid_X2[centroids$cluster == 1])^2), 4)
# create distance from cluster2 centroid
obs$dist_from_cluster2_centroid <- round(</pre>
  sqrt((obs$X1 - centroids$centroid_X1[centroids$cluster == 2])^2 +
       (obs$X2 - centroids$centroid_X2[centroids$cluster == 2])^2), 4)
obs$new_cluster <-
  ifelse(obs$dist_from_cluster1_centroid <= obs$dist_from_cluster2_centroid, 1, 2)</pre>
obs
    X1 X2 cluster dist_from_cluster1_centroid dist_from_cluster2_centroid
## 1 1 4
                 2
                                         1.6640
                                                                      3.3360
## 2 1 3
                 1
                                         1.0530
                                                                      2.8511
## 3 0 4
                                         2.4019
                                                                      4.1796
## 4 5 1
                 1
                                         3.4335
                                                                      1.6640
## 5 6 2
                 2
                                         4.0557
                                                                      2.3300
## 6 4 0
                                         3.3360
                                                                      2.0270
## new cluster
## 1
               1
## 2
## 3
               1
## 4
               2
## 5
               2
## 6
cluster_difference <- FALSE</pre>
while(cluster_difference == FALSE){
# reset obs$cluster to obs$new cluster
obs$cluster <- obs$new_cluster</pre>
# create distance from cluster1 centroid
obs$dist_from_cluster1_centroid <- round(</pre>
  sqrt((obs$X1 - centroids$centroid_X1[centroids$cluster == 1])^2 +
       (obs$X2 - centroids$centroid_X2[centroids$cluster == 1])^2), 4)
# create distance from cluster2 centroid
obs$dist_from_cluster2_centroid <- round(</pre>
  sqrt((obs$X1 - centroids$centroid_X1[centroids$cluster == 2])^2 +
       (obs$X2 - centroids$centroid_X2[centroids$cluster == 2])^2), 4)
obs$new_cluster <-
  ifelse(obs$dist_from_cluster1_centroid <= obs$dist_from_cluster2_centroid, 1, 2)</pre>
```

```
cluster_difference <- all(obs$cluster == obs$new_cluster)</pre>
print(obs)
    {\tt X1~X2~cluster~dist\_from\_cluster1\_centroid~dist\_from\_cluster2\_centroid}
## 1 1 4
                                        1.6640
                                                                     3.3360
## 2 1 3
                 1
                                        1.0530
                                                                     2.8511
## 3 0 4
                 1
                                        2.4019
                                                                     4.1796
                 2
## 4 5 1
                                        3.4335
                                                                     1.6640
## 5 6 2
                 2
                                        4.0557
                                                                     2.3300
## 6 4 0
                 2
                                        3.3360
                                                                     2.0270
## new_cluster
## 1
## 2
               1
## 3
## 4
               2
## 5
               2
## 6
               2
```

The clustering only took one iteration until the algorithm converged.

```
• f)
obs %>%
  dplyr::select(X1, X2, new_cluster) %>%
  ggplot(aes(x = X1, y = X2, color = as.factor(new_cluster))) +
  geom_point()
```



```
# load the data
data("USArrests")

# scale the data so that each variable has a mean of 0 and a variance of 1
usa_scaled <- as.data.frame(scale(USArrests))

# verify that the scaled data's variables all have mean 0 and variance 1
apply(usa_scaled, 2, function(x) round(c(mean(x), var(x)), 2))

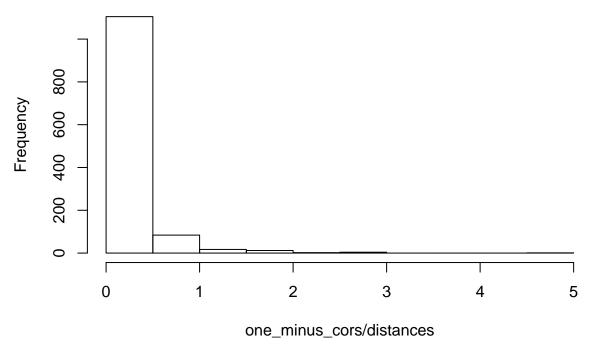
## Murder Assault UrbanPop Rape
## [1,] 0 0 0 0 0
```

```
## [2,] 1 1 1 1 1
# create data frame of combinations of rows of usa_scaled data set
# there are choose(50, 2) = 1225 ways to choose 2 observations out of 50
combs <- t(combn(nrow(usa_scaled), 2))
head(combs)</pre>
```

```
##
       [,1] [,2]
## [1,]
          1
## [2,]
          1
## [3,]
          1
## [4,]
          1
              5
## [5,]
          1
## [6,]
          1
###----- Determine Correlations -----
```

```
# cycle through combs to get each correlation between rows
one_minus_cors <- rep(0, nrow(combs))</pre>
# create function for getting correlations between each observation
one_minus_cor_usa <- function(x){</pre>
  res <- cor(as.numeric(usa_scaled[combs[x, 1], ]),
            as.numeric(usa_scaled[combs[x, 2], ]))
 res <- 1 - res
  res
}
# determine 1 - r_{ij} for each pair of observations
one_minus_cors <- sapply(1:nrow(combs), function(x) one_minus_cor_usa(x))</pre>
###----- Determine Squared Euclidean Distances ------
# cycle through combs to get distance between rows
distances <- dist(usa_scaled)^2
###----- Determine Proportionality ----
summary(one_minus_cors/distances)
##
      Min. 1st Qu.
                      Median
                                 Mean 3rd Qu.
                                                   Max.
## 0.000086 0.069140 0.133900 0.234200 0.262600 4.888000
hist(one_minus_cors/distances)
```

Histogram of one_minus_cors/distances



As can be seen from the summary above, there is no evidence of proportionality. If there were proportionality, there would only be one value. I believe the wording of this question is either wrong or simply hard to understand. The question refers to distances and correlations between the i^{th} and j^{th} observations, which

would lead me to believe it is referring to the rows of the data set (i.e., 50 rows; one per state). If the authors meant to refer to the distances and correlations between i^{th} and j^{th} variables (i.e., the 4 variables of Murder, Assault, UrbanPop, and Rape), then there is constant proportionality. This can be shown below.

```
# squared distance matrix
distance_matrix <- dist(t(usa_scaled))^2

# 1 - correlation matrix
one_minus_cor_mat <- as.dist(1-cor(usa_scaled))

# demonstration of proportionality
distance_matrix/one_minus_cor_mat</pre>
```