

Sharon Kasturiarachi

MIS 64060 Fundamentals of Machine Learning

Dr. Rouzbeh Razavi

Assignment 4- Answers

Kasturiarachi-Assignment 4

```
> summary(Pharmaceuticals)
      Symbol      Name      Market_Cap      Beta      PE_Ratio
Length:21      Length:21      Min.   : 0.41      Min.   :0.1800      Min.   : 3.60
Class :character Class :character 1st Qu.: 6.30      1st Qu.:0.3500      1st Qu.:18.90
Mode  :character Mode  :character Median : 48.19      Median :0.4600      Median :21.50
                        Mean  : 57.65      Mean  :0.5257      Mean  :25.46
                        3rd Qu.: 73.84      3rd Qu.:0.6500      3rd Qu.:27.90
                        Max.   :199.47      Max.   :1.1100      Max.   :82.50
      ROE      ROA      Asset_Turnover      Leverage      Rev_Growth      Net_Profit_Margin
Min.   : 3.9      Min.   : 1.40      Min.   :0.3      Min.   :0.0000      Min.   : -3.17      Min.   : 2.6
1st Qu.:14.9      1st Qu.: 5.70      1st Qu.:0.6      1st Qu.:0.1600      1st Qu.: 6.38      1st Qu.:11.2
Median :22.6      Median :11.20      Median :0.6      Median :0.3400      Median : 9.37      Median :16.1
Mean   :25.8      Mean   :10.51      Mean   :0.7      Mean   :0.5857      Mean   :13.37      Mean   :15.7
3rd Qu.:31.0      3rd Qu.:15.00      3rd Qu.:0.9      3rd Qu.:0.6000      3rd Qu.:21.87      3rd Qu.:21.1
Max.   :62.9      Max.   :20.30      Max.   :1.1      Max.   :3.5100      Max.   :34.21      Max.   :25.5
Median_Recommendation Location      Exchange
Length:21      Length:21      Length:21
Class :character Class :character Class :character
Mode  :character Mode  :character Mode  :character

> colnames(Pharmaceuticals)
 [1] "Symbol"      "Name"      "Market_Cap"      "Beta"
 [5] "PE_Ratio"    "ROE"      "ROA"      "Asset_Turnover"
 [9] "Leverage"    "Rev_Growth"      "Net_Profit_Margin"      "Median_Recommendation"
[13] "Location"    "Exchange"

> str(Pharmaceuticals)
spec_tbl_df [21 x 14] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
 $ Symbol      : chr [1:21] "ABT" "AGN" "AHM" "AZN" ...
 $ Name        : chr [1:21] "Abbott Laboratories" "Allergan, Inc." "Amersham plc" "AstraZeneca PL
C" ...
 $ Market_Cap  : num [1:21] 68.44 7.58 6.3 67.63 47.16 ...
 $ Beta        : num [1:21] 0.32 0.41 0.46 0.52 0.32 1.11 0.5 0.85 1.08 0.18 ...
 $ PE_Ratio    : num [1:21] 24.7 82.5 20.7 21.5 20.1 27.9 13.9 26 3.6 27.9 ...
 $ ROE         : num [1:21] 26.4 12.9 14.9 27.4 21.8 3.9 34.8 24.1 15.1 31 ...
 $ ROA         : num [1:21] 11.8 5.5 7.8 15.4 7.5 1.4 15.1 4.3 5.1 13.5 ...
 $ Asset_Turnover : num [1:21] 0.7 0.9 0.9 0.9 0.6 0.6 0.9 0.6 0.3 0.6 ...
 $ Leverage    : num [1:21] 0.42 0.6 0.27 0 0.34 0 0.57 3.51 1.07 0.53 ...
 $ Rev_Growth  : num [1:21] 7.54 9.16 7.05 15 26.81 ...
 $ Net_Profit_Margin : num [1:21] 16.1 5.5 11.2 18 12.9 2.6 20.6 7.5 13.3 23.4 ...
 $ Median_Recommendation: chr [1:21] "Moderate Buy" "Moderate Buy" "Strong Buy" "Moderate Sell" ...
 $ Location    : chr [1:21] "US" "CANADA" "UK" "UK" ...
 $ Exchange    : chr [1:21] "NYSE" "NYSE" "NYSE" "NYSE" ...
- attr(*, "spec")=
```

After removing the variables that we do not want to include in the model

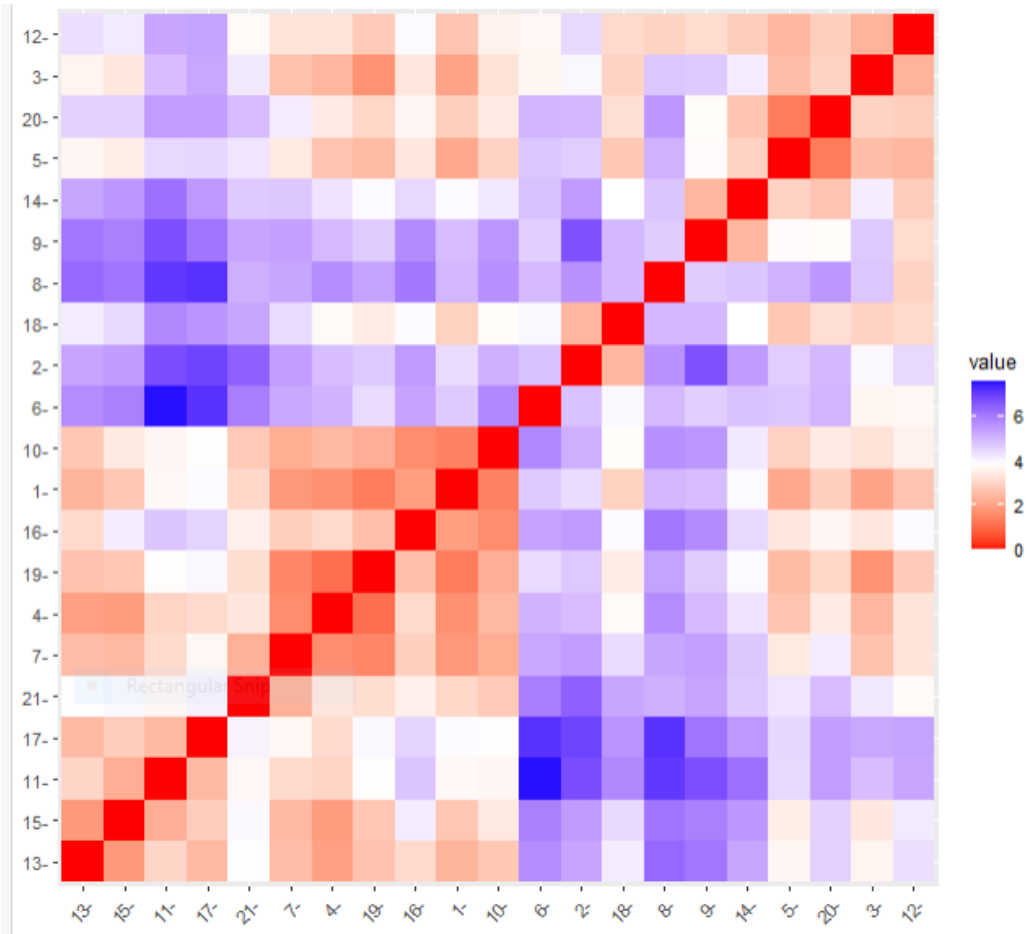
```
#Remove all categorical variables
Pharmaceutical_data$Symbol<- NULL
Pharmaceutical_data$Name <- NULL
Pharmaceutical_data$Median_Recommendation<-NULL
Pharmaceutical_data$Location <-NULL
Pharmaceutical_data$Exchange <-NULL
str(Pharmaceutical_data)
data.frame': 21 obs. of 9 variables:
 $ Market_Cap      : num  68.44 7.58 6.3 67.63 47.16 ...
 $ Beta            : num  0.32 0.41 0.46 0.52 0.32 1.11 0.5 0.85 1.08 0.18 ...
 $ PE_Ratio        : num  24.7 82.5 20.7 21.5 20.1 27.9 13.9 26 3.6 27.9 ...
 $ ROE             : num  26.4 12.9 14.9 27.4 21.8 3.9 34.8 24.1 15.1 31 ...
 $ ROA             : num  11.8 5.5 7.8 15.4 7.5 1.4 15.1 4.3 5.1 13.5 ...
 $ Asset_Turnover   : num  0.7 0.9 0.9 0.9 0.6 0.6 0.9 0.6 0.3 0.6 ...
 $ Leverage         : num  0.42 0.6 0.27 0 0.34 0 0.57 3.51 1.07 0.53 ...
 $ Rev_Growth       : num  7.54 9.16 7.05 15 26.81 ...
 $ Net_Profit_Margin: num  16.1 5.5 11.2 18 12.9 2.6 20.6 7.5 13.3 23.4 ...
 |
```

Normalizing the data for each variable to be treated equally by the distance measure

```
> Pharmaceutical_data.norm
      Market_Cap      Beta      PE_Ratio      ROE      ROA      Asset_Turnover      Leverage      Rev_Growth
[1,]  0.1840960 -0.80125356 -0.04671323  0.04009035  0.2416121      0.0000000 -0.21209793 -0.52776752
[2,] -0.8544181 -0.45070513  3.49706911 -0.85483986 -0.9422871      0.9225312  0.01828430 -0.38113909
[3,] -0.8762600 -0.25595600 -0.29195768 -0.72225761 -0.5100700      0.9225312 -0.40408312 -0.57211809
[4,]  0.1702742 -0.02225704 -0.24290879  0.10638147  0.9181259      0.9225312 -0.74965647  0.14744734
[5,] -0.1790256 -0.80125356 -0.32874435 -0.26484883 -0.5664461     -0.4612656 -0.31449003  1.21638667
[6,] -0.6953818  2.27578267  0.14948233 -1.45146000 -1.7127612     -0.4612656 -0.74965647 -1.49714434
[7,] -0.1078688 -0.10015669 -0.70887325  0.59693581  0.8617498      0.9225312 -0.02011273 -0.96584257
[8,] -0.9767669  1.26308721  0.03299122 -0.11237924 -1.1677918     -0.4612656  3.74279705 -0.63276071
[9,] -0.9704532  2.15893320 -1.34037772 -0.70899938 -1.0174553     -1.8450624  0.61983791  1.88617085
[10,] 0.2762415 -1.34655112  0.14948233  0.34502953  0.5610770     -0.4612656 -0.07130879 -0.64814764
[11,] 1.0999201 -0.68440408 -0.45749769  2.45971647  1.8389364      1.3837968 -0.31449003  0.76926048
[12,] -0.9393967  0.48409069 -0.34100657 -0.29136529 -0.6979905     -0.4612656  1.10620040  0.05603085
[13,] 1.9841758 -0.25595600  0.18013789  0.18593083  1.0872544      0.9225312 -0.62166634 -0.36213170
[14,] -0.9632863  0.87358895  0.19240011 -0.96753478 -0.9610792     -1.8450624  0.44065173  1.53860717
[15,] 1.2782387 -0.25595600 -0.40231769  0.98142435  0.8429577      1.8450624 -0.39128411  0.36014907
[16,] 0.6654710 -1.30760129 -0.23677768 -0.52338423  0.1288598     -0.9225312 -0.67286239 -1.45369888
[17,] 2.4199899  0.48409069 -0.11415545  1.31287998  1.6322239      0.4612656 -0.54487226  1.10143723
[18,] -0.0240846 -0.48965495  1.90298017 -0.81506519 -0.9047030     -0.4612656 -0.30169102  0.14744734
[19,] -0.4018812 -0.06120687 -0.40231769 -0.21181593  0.5234929      0.4612656 -0.74965647 -0.43544591
[20,] -0.9281345 -1.11285216 -0.43297324 -1.03382590 -0.6979905     -0.9225312 -0.49367621  1.43089863
[21,] -0.1614497  0.40619104 -0.75792214  1.92938746  0.5422849     -0.4612656  0.68383297 -1.17763919

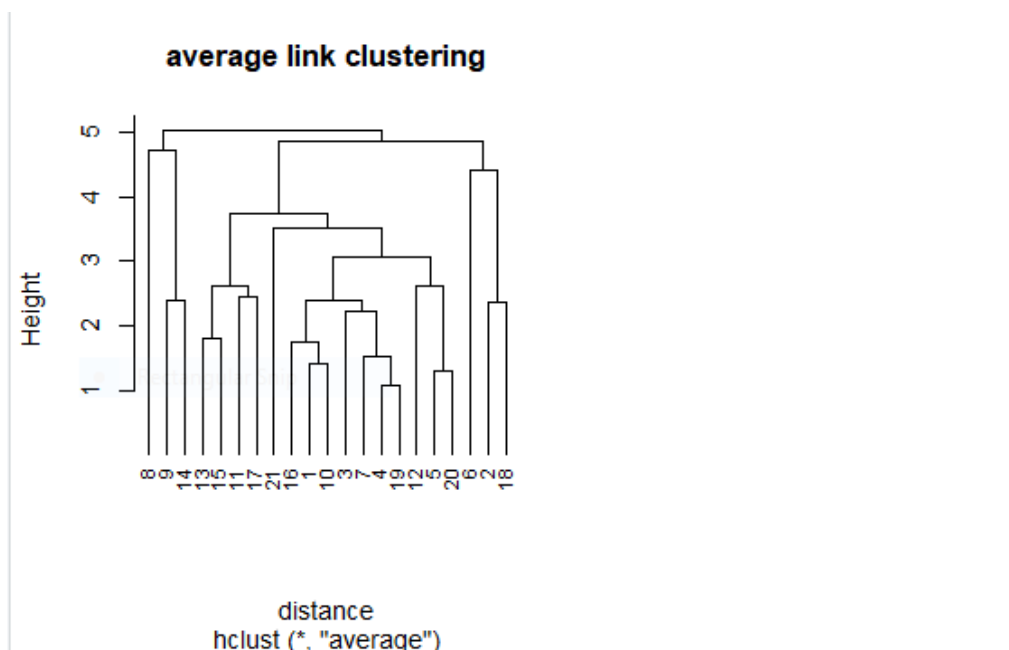
      Net_Profit_Margin
[1,]  0.06168225
[2,] -1.55366706
[3,] -0.68503583
[4,]  0.35122600
[5,] -0.42597037
[6,] -1.99560225
[7,]  0.74744375
[8,] -1.24888417
[9,] -0.36501379
[10,] 1.17413980
[11,] 0.82363947
[12,] -0.71551412
[13,] 0.33598685
[14,] 0.85411776
[15,] -0.24310064
[16,] 1.02174835
[17,] 1.44844440
[18,] -1.27936246
[19,] 0.29026942
[20,] -0.09070919
[21,] 1.49416183
> |
```

Let's look at the distance between observations and using the Euclidean distance to find the similarity between the observations

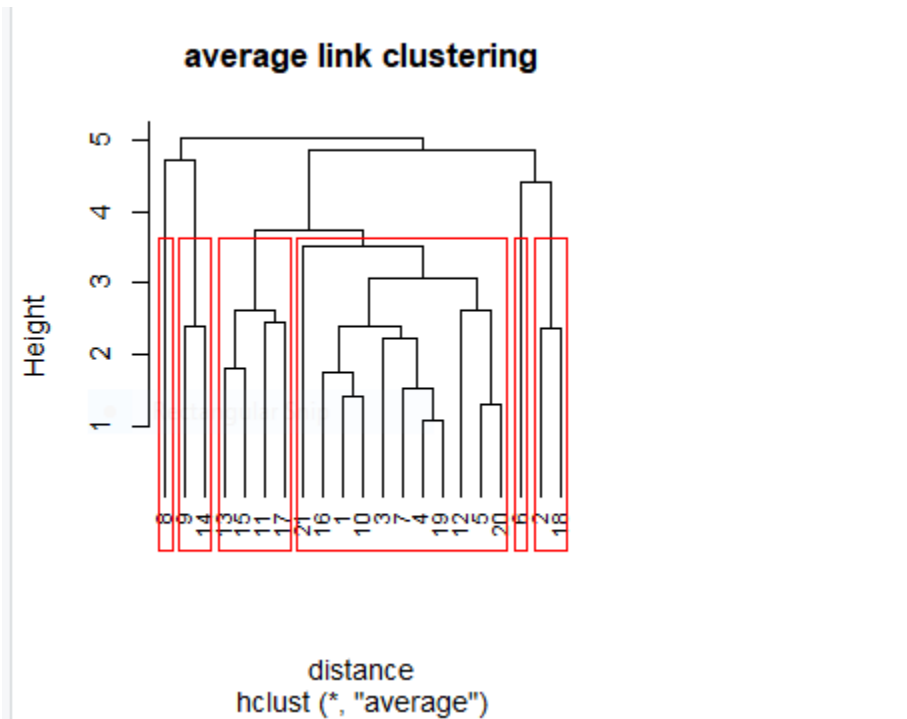


Using the Hierarchical clustering to identify the relationship between individual data points and clusters.

The average linkage helps to find the average distance between two clusters



Using the cutree function to cut the dendrogram into 6 clusters. Here the dendrogram shows the relationship between individual data points and clusters, where the height is the distance between clusters.



```
> rect.hclust(fit.average, k=6, border="red" )
> table(clusters)
clusters
 1  2  3  4  5  6
11  2  1  1  2  4
```

Now to determine the optimal number of clusters using the k-means algorithm.

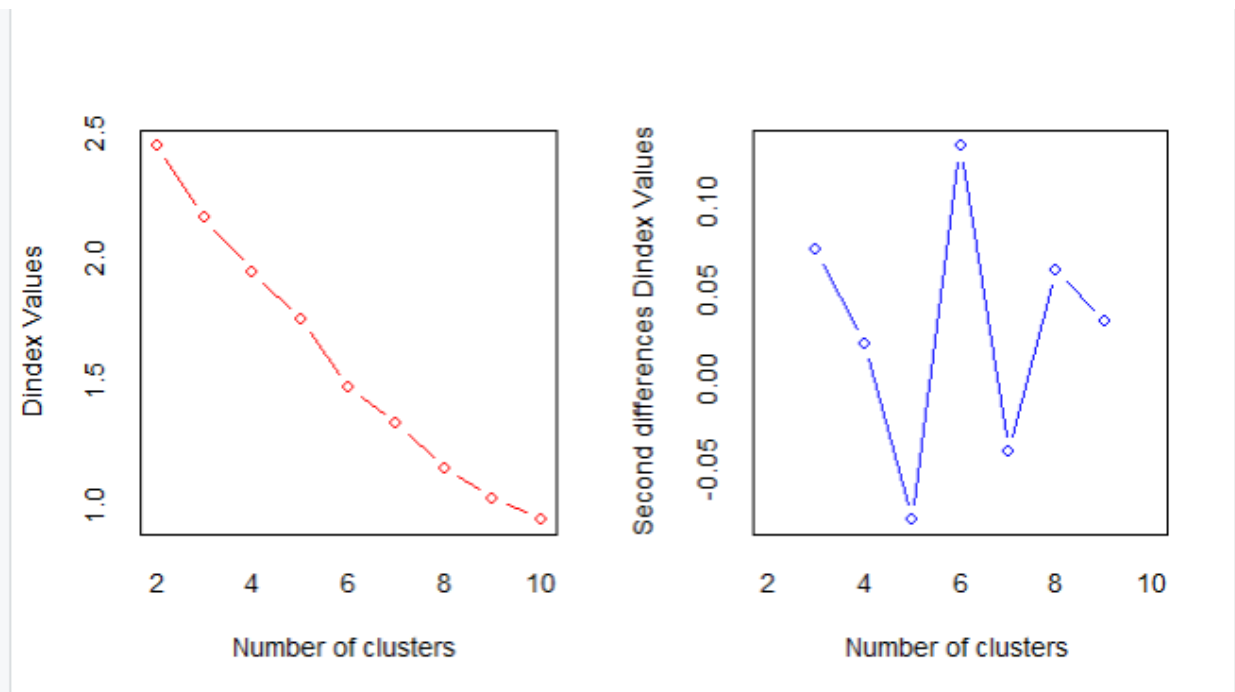
```
> nc <- NbClust(Pharmaceutical_data.norm, distance = "euclidean", min.nc = 2, max.nc = 10, method = "average")
*** : The Hubert index is a graphical method of determining the number of clusters.
      In the plot of Hubert index, we seek a significant knee that corresponds to a
      significant increase of the value of the measure i.e the significant peak in Hubert
      index second differences plot.

*** : The D index is a graphical method of determining the number of clusters.
      In the plot of D index, we seek a significant knee (the significant peak in Dindex
      second differences plot) that corresponds to a significant increase of the value of
      the measure.

*****
* Among all indices:
* 4 proposed 2 as the best number of clusters
* 7 proposed 3 as the best number of clusters
* 3 proposed 5 as the best number of clusters
* 1 proposed 6 as the best number of clusters
* 4 proposed 8 as the best number of clusters
* 2 proposed 9 as the best number of clusters
* 2 proposed 10 as the best number of clusters

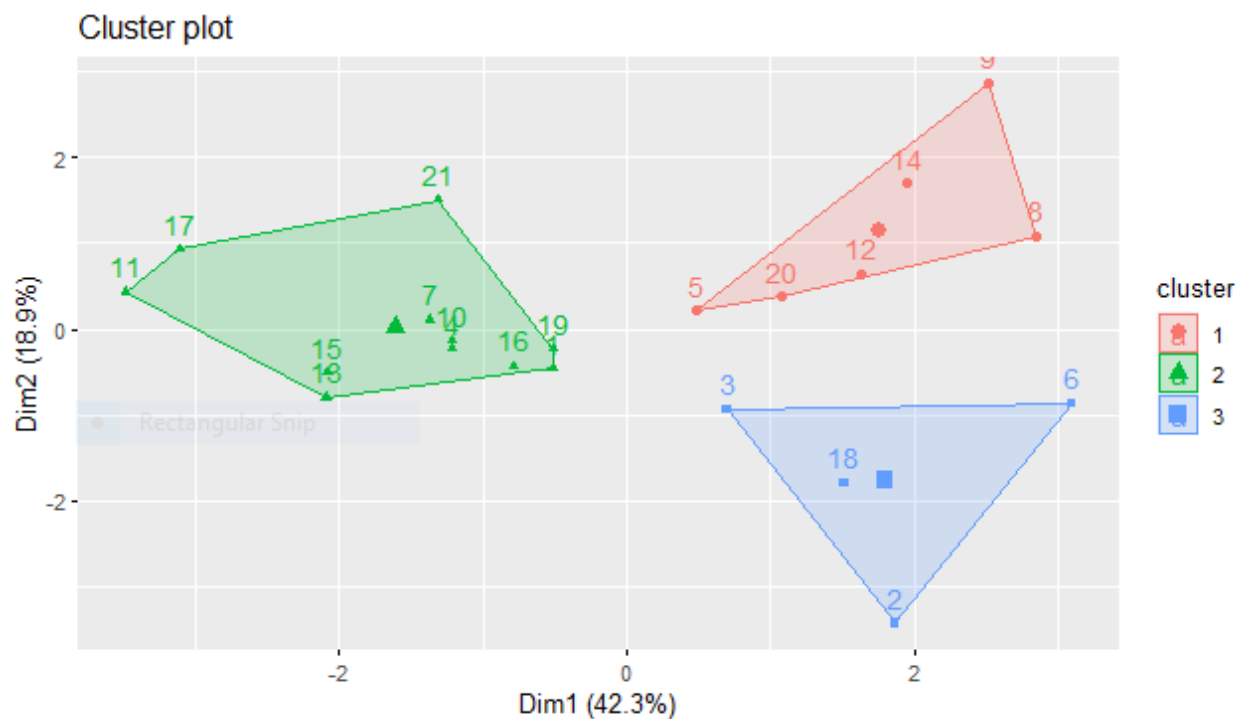
***** Conclusion *****

* According to the majority rule, the best number of clusters is 3
```



So, I will use 3 clusters to perform the k-means where $k=3$ and 25 restarts to perform the cluster analysis

```
> k3$centers
  Market_Cap      Beta  PE_Ratio      ROE      ROA Asset_Turnover  Leverage
1 -0.8261772  0.4775991 -0.3696184 -0.5631589 -0.8514589   -0.9994088  0.8502201
2  0.6733825 -0.3586419 -0.2763512  0.6565978  0.8344159    0.4612656 -0.3331068
3 -0.6125361  0.2698666  1.3143935 -0.9609057 -1.0174553    0.2306328 -0.3592866
  Rev_Growth Net_Profit_Margin
1  0.9158889   -0.3319956
2 -0.2902163    0.6823310
3 -0.5757385   -1.3784169
> k3$size
[1] 6 11 4
> k3$cluster
[1] 2 3 3 2 1 3 2 1 1 2 2 1 2 1 2 2 2 3 2 1 2
>
```



Checking for any outliers

```
> dist(k3$centers)
      1      2
2 3.647470
3 3.066970 3.873875
> |
```

(b)

Now that we have the pharmaceutical companies belonging to one of the 3 clusters. We need to dive into each cluster to analyze the characteristics and variables.

Cluster 1-rows 5, 8, 9, 12, 14, 20

Cluster 2-rows 4,7,10,11, 15, 16,17,18, 19,21

Cluster 3-rows 2, 3, 6, 18

```
> k3$size
[1] 6 11 4
```

	Group.1	Market_Cap	Beta	PE_Ratio	ROE	ROA	Asset_Turnover	Leverage
1	1	-0.8261772	0.4775991	-0.3696184	-0.5631589	-0.8514589	-0.9994088	0.8502201
2	2	0.6733825	-0.3586419	-0.2763512	0.6565978	0.8344159	0.4612656	-0.3331068
3	3	-0.6125361	0.2698666	1.3143935	-0.9609057	-1.0174553	0.2306328	-0.3592866
		Rev_Growth	Net_Profit_Margin					
1		0.9158889	-0.3319956					
2		-0.2902163	0.6823310					
3		-0.5757385	-1.3784169					

This indicates that there are 6 pharmaceuticals in cluster 1, 11 pharmaceuticals in cluster 2 and 4 pharmaceuticals in cluster 3

Cluster 1- is characterized by companies that high revenue growth, low ROA and ROE, lowest Market cap and Asset Turnover, highly leveraged, high beta more volatile

Cluster 2- companies that have high net profit margin, high ROE and ROA and highest market Cap

Cluster 3- companies with lowest revenue growth and profit margin, high PE ratio, low ROE, and ROA

The Hierarchical cluster algorithm shows Cluster 1 and Cluster 2 have similar pattern, consisting of companies that are highly profitable and low risk investment. On the other hand, Cluster 3 is comprised of not-for-profit companies with low leverage, and less debt. Moreover, Cluster 1 includes not-for-profit companies with high volatility due to high beta and considering revenue growth and high level of leverage like Cluster 3. The stock price is undervalued.

C) Is there a pattern in the clusters with respect to the numerical variables (10 to 12)? (those not used in forming the clusters)

Yes, there is a pattern in the clusters with respect to variable Median Recommendation categorized into Moderate Buy, Moderate Sell, Strong Buy, and hold.

When comparing the clusters, Cluster 2 has the highest Market Cap, highest ROE and ROA, and highest Asset Turnover, yet does not indicate a moderate sell. On the other hand, Cluster 3 with high PE ratio, lowest revenue growth and profit margin, lowest ROE, ROA, and lowest asset turnover has a strong buy recommendation as the stock price is undervalued. Cluster 1 high beta, leverage, revenue growth and low market cap on hold recommendation.

(d) Naming the clusters according to variables

Cluster 1- highest revenue growth, highest leverage, highest beta, low ROA and ROE, lowest Market cap and Asset Turnover-Risky yet high revenue

Cluster 2 -High ROA, high ROE, high net profit and least risk- Moderate Buy

Cluster 3 - lowest revenue growth and profit margin, high PE ratio, low ROE, and ROA- Strong Buy

Environment	History	Connections	Tutorial
R Global Environment			
Data			
fit.average	List of 7		
k3	List of 9		
nc	List of 4		
Pharmaceutical_data	21 obs. of 9 variables		
Pharmaceutical_data.norm	num [1:21, 1:9] 0.184 -0.854 -0.876 0.17 -0.179 ...		
Pharmaceuticals	21 obs. of 14 variables		
Pharmaceuticals_data	21 obs. of 14 variables		
Values			
clusters	int [1:21] 1 2 1 1 1 3 1 4 5 1 ...		
distance	'dist' num [1:210] 4.42 2.02 1.67 2.11 4.69 ...		