

# fml assignment 5

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#Importing dataset

```
CEREALS <- read.csv("C:/Users/gaya3/Downloads/CEREALS.csv")
head(CEREALS)
```

```
##              name mfr type calories protein fat sodium fiber carbo
## 1          100%_Bran  N   C        70         4  1   130   10.0   5.0
## 2      100%_Natural_Bran  Q   C       120         3  5    15    2.0   8.0
## 3           All-Bran   K   C        70         4  1   260    9.0   7.0
## 4 All-Bran_with_Extra_Fiber  K   C        50         4  0   140   14.0   8.0
## 5         Almond_Delight  R   C       110         2  2   200    1.0  14.0
## 6  Apple_Cinnamon_Cheerios  G   C       110         2  2   180    1.5  10.5
##  sugars potass vitamins shelf weight cups  rating
## 1      6      280       25    3      1 0.33 68.40297
## 2      8      135        0    3      1 1.00 33.98368
## 3      5      320       25    3      1 0.33 59.42551
## 4      0      330       25    3      1 0.50 93.70491
## 5      8       NA       25    3      1 0.75 34.38484
## 6     10       70       25    1      1 0.75 29.50954
```

```
dim(CEREALS)
```

```
## [1] 77 16
```

#Loading packages

```
library(cluster)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##   filter, lag
```

```
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
library(caret)
```

```
## Loading required package: ggplot2
```

```
## Loading required package: lattice
```

```
library(dendextend)
```

```
## Warning: package 'dendextend' was built under R version 4.2.3
```

```
##
## -----
## Welcome to dendextend version 1.17.1
## Type citation('dendextend') for how to cite the package.
##
## Type browseVignettes(package = 'dendextend') for the package vignette.
## The github page is: https://github.com/talgalili/dendextend/
##
## Suggestions and bug-reports can be submitted at: https://github.com/talgalili/dendextend/i
## ssues
## You may ask questions at stackoverflow, use the r and dendextend tags:
##   https://stackoverflow.com/questions/tagged/dendextend
##
## To suppress this message use: suppressPackageStartupMessages(library(dendextend))
## -----
```

```
##
## Attaching package: 'dendextend'
```

```
## The following object is masked from 'package:stats':
##
##   cutree
```

```
library(knitr)
library(factoextra)
```

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

```
library(readr)
```

#Omitting the NULL values

```
CEREALS <- na.omit(CEREALS)
dim(CEREALS)
```

```
## [1] 74 16
```

```
head(CEREALS)
```

```
##           name mfr type calories protein fat sodium fiber carbo
## 1      100%_Bran   N   C       70      4   1   130  10.0   5.0
## 2    100%_Natural_Bran   Q   C      120      3   5    15   2.0   8.0
## 3           All-Bran   K   C       70      4   1   260   9.0   7.0
## 4 All-Bran_with_Extra_Fiber   K   C       50      4   0   140  14.0   8.0
## 6   Apple_Cinnamon_Cheerios   G   C      110      2   2   180   1.5  10.5
## 7       Apple_Jacks   K   C      110      2   0   125   1.0  11.0
##  sugars potass vitamins shelf weight cups   rating
## 1      6      280      25    3      1 0.33 68.40297
## 2      8      135       0    3      1 1.00 33.98368
## 3      5      320      25    3      1 0.33 59.42551
## 4      0      330      25    3      1 0.50 93.70491
## 6     10       70      25    1      1 0.75 29.50954
## 7     14       30      25    2      1 1.00 33.17409
```

#Creating a dataset with the Numeric Values

```
df1<- data.frame(CEREALS[,4:16])
df2 <- na.omit(df1)
```

#Normalizing the data

```
df1 <- scale(df1)
head(df1)
```

```
##      calories   protein      fat    sodium      fiber      carbo      sugars
## 1 -1.8659155  1.3817478  0.0000000 -0.3910227  3.22866747 -2.5001396 -0.2542051
## 2  0.6537514  0.4522084  3.9728810 -1.7804186 -0.07249167 -1.7292632  0.2046041
## 3 -1.8659155  1.3817478  0.0000000  1.1795987  2.81602258 -1.9862220 -0.4836096
## 4 -2.8737823  1.3817478 -0.9932203 -0.2702057  4.87924705 -1.7292632 -1.6306324
## 6  0.1498180 -0.4773310  0.9932203  0.2130625 -0.27881412 -1.0868662  0.6634132
## 7  0.1498180 -0.4773310 -0.9932203 -0.4514312 -0.48513656 -0.9583868  1.5810314
##      potass  vitamins      shelf    weight      cups      rating
## 1  2.5605229 -0.1818422  0.9419715 -0.2008324 -2.0856582  1.8549038
## 2  0.5147738 -1.3032024  0.9419715 -0.2008324  0.7567534 -0.5977113
## 3  3.1248675 -0.1818422  0.9419715 -0.2008324 -2.0856582  1.2151965
## 4  3.2659536 -0.1818422  0.9419715 -0.2008324 -1.3644493  3.6578436
## 6 -0.4022862 -0.1818422 -1.4616799 -0.2008324 -0.3038480 -0.9165248
## 7 -0.9666308 -0.1818422 -0.2598542 -0.2008324  0.7567534 -0.6553998
```

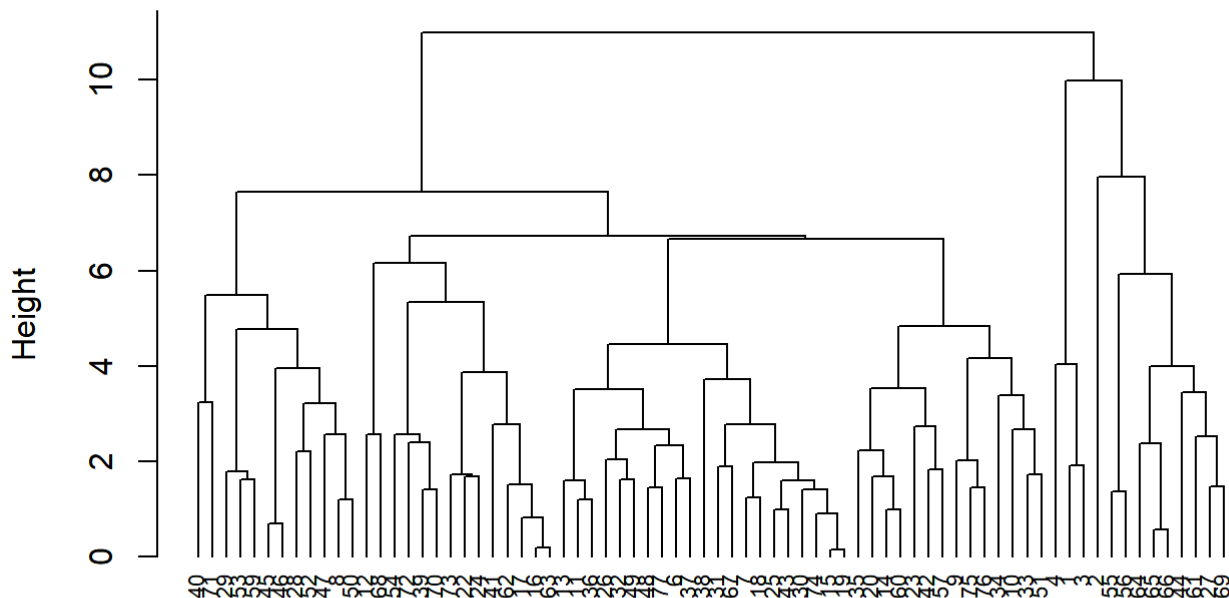
#Applying hierarchical clustering using Euclidean distance method.

```
distance <- dist(df1, method= "euclidean")
Hist_cluster <- hclust(distance, method = "complete")
```

#Plotting of the dendrogram

```
plot(Hist_cluster, cex = 0.7, hang = -1)
```

## Cluster Dendrogram



#Using Agnes function to perform clustering with single linkage, complete linkage average linkage and Ward.

```
hc.single <- agnes(df1, method = "single")
hc.complete <- agnes(df1, method = "complete")
hc.average <- agnes(df1, method = "average")
hc.ward <- agnes(df1, method = "ward")
```

#Determining the best method

```
print(hc.single$ac)
```

```
## [1] 0.6067859
```

```
print(hc.complete$ac)
```

```
## [1] 0.8353712
```

```
print(hc.average$ac)
```

```
## [1] 0.7766075
```

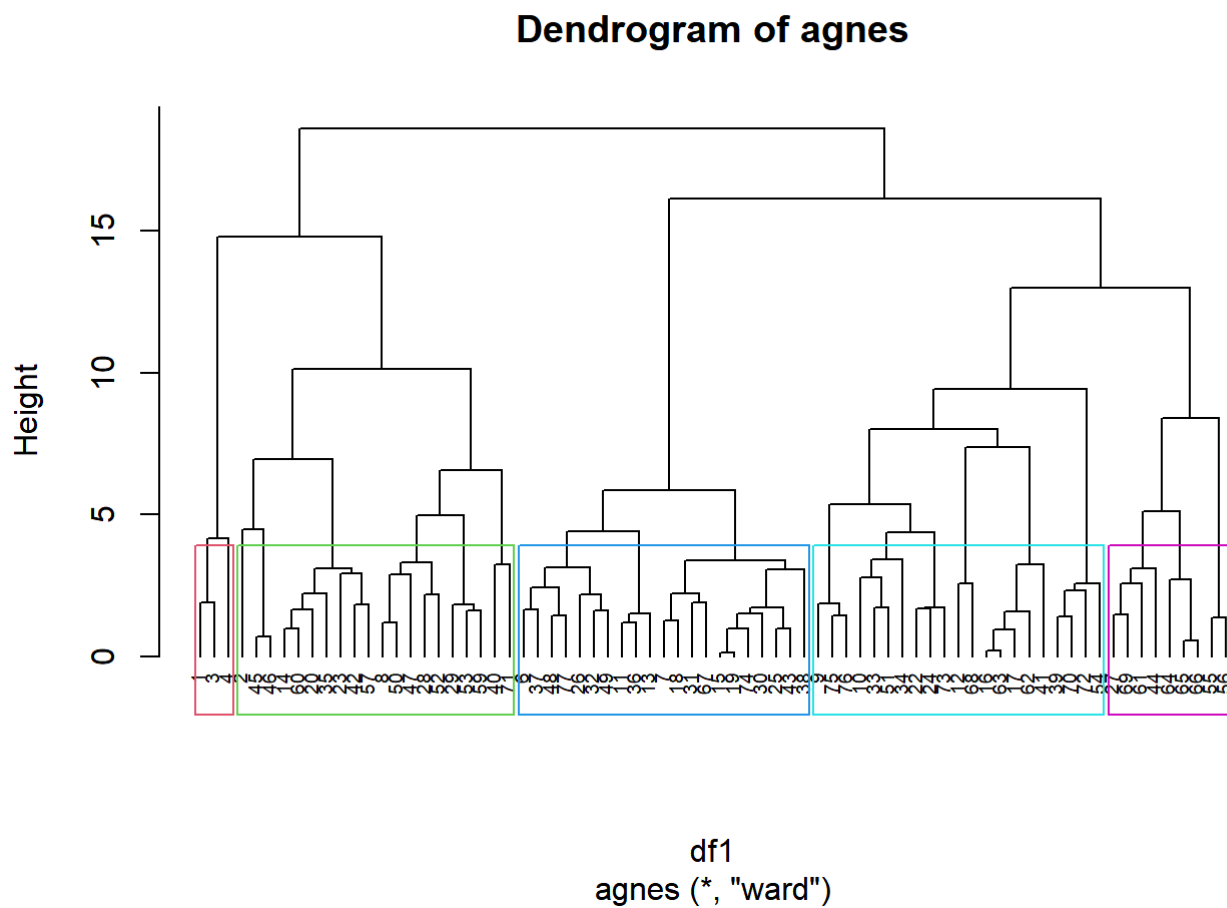
```
print(hc.ward$ac)
```

```
## [1] 0.9046042
```

#With a rating of 0.9046042, the ward technique is superior to the other methods.

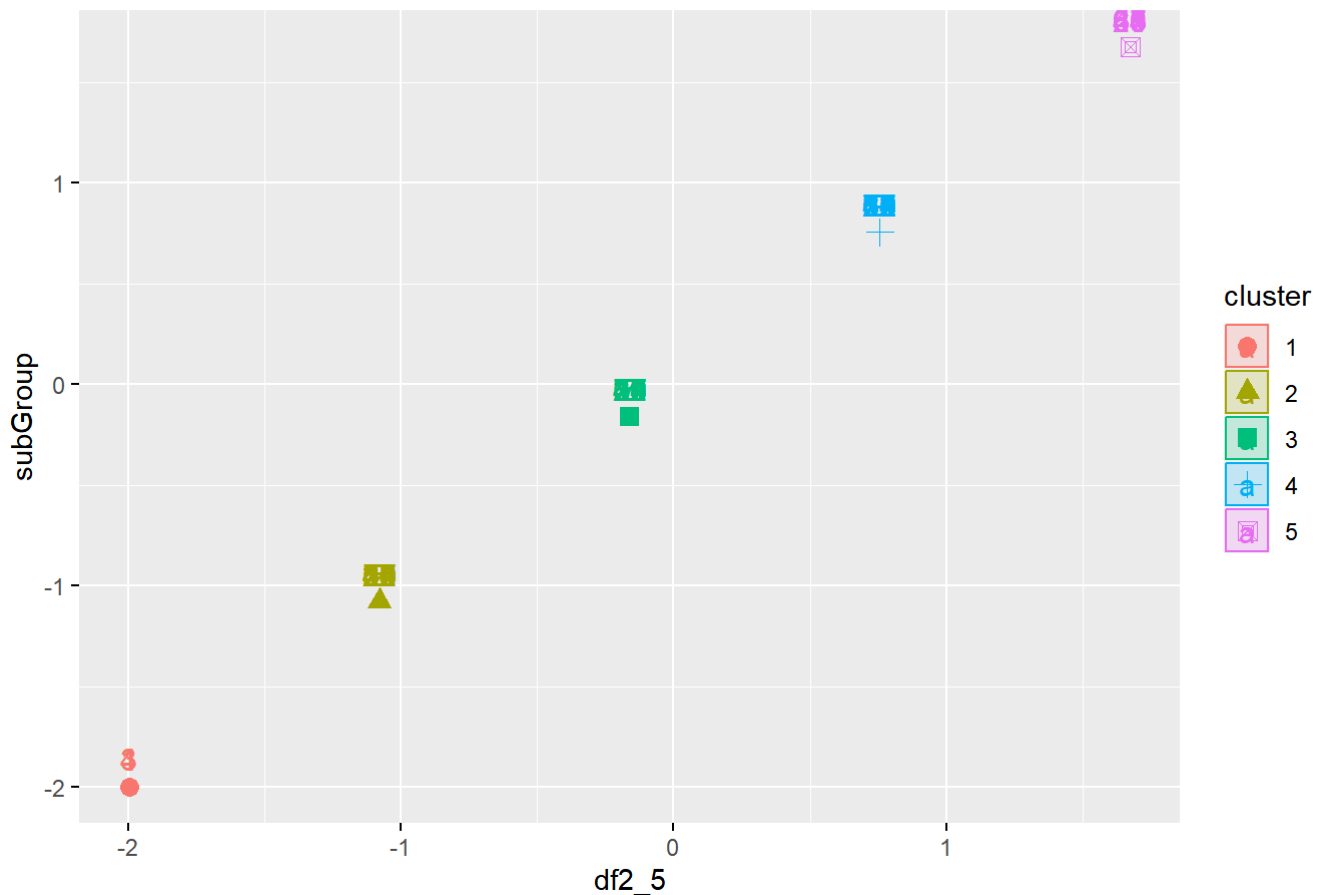
#Choosing the number of clusters

```
pltree(hc.ward, cex = 0.6, hang = -1, main = "Dendrogram of agnes")
df2_5 <- cutree(hc.ward, k = 5)
rect.hclust(hc.ward , k=5, border = 2:7)
```



```
subGroup <- cutree(hc.ward, k=5)
df2_5 <- as.data.frame(cbind(df2_5,subGroup))
fviz_cluster(list(data=df2_5, cluster = subGroup))
```

Cluster plot



#It is concluded that 5 clusters can be selected.

#Creating Partitions

```
set.seed(123)
df_A <- df2 [1:55,]
df_B <- df2 [56:74,]
```

#Performing Hierarchical Clustering, considering k = 5.

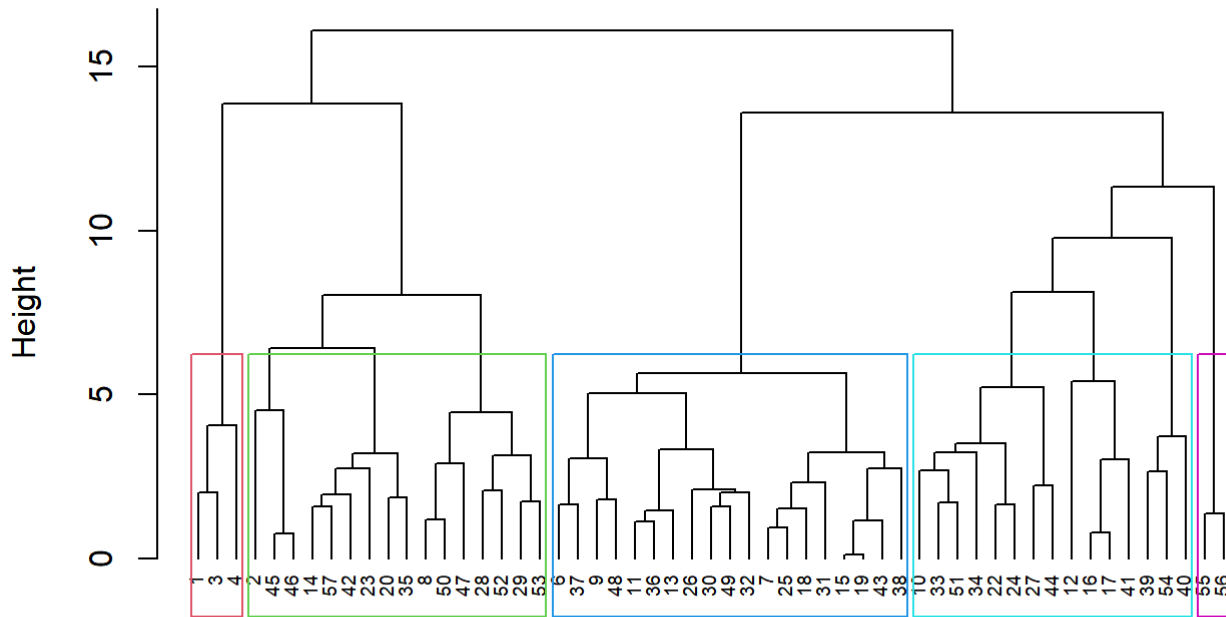
```
Ag.single <- agnes(scale(df_A), method = "single")
Ag.complete <- agnes(scale(df_A), method = "complete")
Ag.average <- agnes(scale(df_A), method = "average")
Ag.ward <- agnes(scale(df_A), method = "ward")

cbind(single= Ag.single$ac , complete=Ag.complete$ac , average= Ag.average$ac , ward= Ag.ward$ac)
```

```
##          single  complete  average    ward
## [1,] 0.6564842 0.8120228 0.7449303 0.8808195
```

```
pltree(Ag.ward, cex = 0.6, hang = -1, main = "Dendrogram of Agnes Using Ward")
rect.hclust(Ag.ward, k = 5, border = 2:7)
```

## Dendrogram of Agnes Using Ward



scale(df\_A)  
agnes (\*, "ward")

```
cut2 <- cutree(Ag.ward, k = 5)
```

#Calculating the centroids.

```
Result <- as.data.frame(cbind(df_A, cut2))
Result[Result$cut2==1,]
```

```
##  calories protein fat sodium fiber carbo sugars potass vitamins shelf weight
## 1      70      4  1   130    10    5      6   280      25      3      1
## 3      70      4  1   260     9    7      5   320      25      3      1
## 4      50      4  0   140    14    8      0   330      25      3      1
##  cups   rating cut2
## 1 0.33 68.40297    1
## 3 0.33 59.42551    1
## 4 0.50 93.70491    1
```

```
Centroid1 <- colMeans(Result[Result$cut2==1,])
Result[Result$cut2==2,]
```

```
##      calories protein fat sodium fiber carbo sugars potass vitamins shelf weight
## 2          120      3  5      15   2.0   8.0      8   135         0     3   1.00
## 8          130      3  2     210   2.0  18.0      8   100        25     3   1.33
## 14         110      3  2     140   2.0  13.0      7   105        25     3   1.00
## 20         110      3  3     140   4.0  10.0      7   160        25     3   1.00
## 23         100      2  1     140   2.0  11.0     10   120        25     3   1.00
## 28         120      3  2     160   5.0  12.0     10   200        25     3   1.25
## 29         120      3  0     240   5.0  14.0     12   190        25     3   1.33
## 35         120      3  3      75   3.0  13.0      4   100        25     3   1.00
## 42         100      4  2     150   2.0  12.0      6    95        25     2   1.00
## 45         150      4  3      95   3.0  16.0     11   170        25     3   1.00
## 46         150      4  3     150   3.0  16.0     11   170        25     3   1.00
## 47         160      3  2     150   3.0  17.0     13   160        25     3   1.50
## 50         140      3  2     220   3.0  21.0      7   130        25     3   1.33
## 52         130      3  2     170   1.5  13.5     10   120        25     3   1.25
## 53         120      3  1     200   6.0  11.0     14   260        25     3   1.33
## 57         100      4  1     135   2.0  14.0      6   110        25     3   1.00
##      cups   rating cut2
## 2  1.00 33.98368     2
## 8  0.75 37.03856     2
## 14 0.50 40.40021     2
## 20 0.50 40.44877     2
## 23 0.75 36.17620     2
## 28 0.67 40.91705     2
## 29 0.67 41.01549     2
## 35 0.33 45.81172     2
## 42 0.67 45.32807     2
## 45 1.00 37.13686     2
## 46 1.00 34.13976     2
## 47 0.67 30.31335     2
## 50 0.67 40.69232     2
## 52 0.50 30.45084     2
## 53 0.67 37.84059     2
## 57 0.50 49.51187     2
```

```
Centroid2 <- colMeans(Result[Result$cut2==2,])
Result[Result$cut2==3,]
```



```
##      calories protein fat sodium fiber carbo sugars potass vitamins shelf weight
## 6         110      2  2   180   1.5  10.5    10    70      25     1     1
## 7         110      2  0   125   1.0  11.0    14    30      25     2     1
## 9          90      2  1   200   4.0  15.0     6   125      25     1     1
## 11        120      1  2   220   0.0  12.0    12    35      25     2     1
## 13        120      1  3   210   0.0  13.0     9    45      25     2     1
## 15        110      1  1   180   0.0  12.0    13    55      25     2     1
## 18        110      1  0    90   1.0  13.0    12    20      25     2     1
## 19        110      1  1   180   0.0  12.0    13    65      25     2     1
## 25        110      2  1   125   1.0  11.0    13    30      25     2     1
## 26        110      1  0   200   1.0  14.0    11    25      25     1     1
## 30        110      1  1   135   0.0  13.0    12    25      25     2     1
## 31        100      2  0    45   0.0  11.0    15    40      25     1     1
## 32        110      1  1   280   0.0  15.0     9    45      25     2     1
## 36        120      1  2   220   1.0  12.0    11    45      25     2     1
## 37        110      3  1   250   1.5  11.5    10    90      25     1     1
## 38        110      1  0   180   0.0  14.0    11    35      25     1     1
## 43        110      2  1   180   0.0  12.0    12    55      25     2     1
## 48        100      2  1   220   2.0  15.0     6    90      25     1     1
## 49        120      2  1   190   0.0  15.0     9    40      25     2     1
##      cups   rating cut2
## 6  0.75 29.50954    3
## 7  1.00 33.17409    3
## 9  0.67 49.12025    3
## 11 0.75 18.04285    3
## 13 0.75 19.82357    3
## 15 1.00 22.73645    3
## 18 1.00 35.78279    3
## 19 1.00 22.39651    3
## 25 1.00 32.20758    3
## 26 0.75 31.43597    3
## 30 0.75 28.02576    3
## 31 0.88 35.25244    3
## 32 0.75 23.80404    3
## 36 1.00 21.87129    3
## 37 0.75 31.07222    3
## 38 1.33 28.74241    3
## 43 1.00 26.73451    3
## 48 1.00 40.10596    3
## 49 0.67 29.92429    3
```

```
Centroid3 <- colMeans(Result[Result$cut2==3,])
Result[Result$cut2==4,]
```

```
##      calories protein fat sodium fiber carbo sugars potass vitamins shelf weight
## 10         90       3  0   210    5   13     5   190       25     3    1.0
## 12        110       6  2   290    2   17     1   105       25     1    1.0
## 16        110       2  0   280    0   22     3    25       25     1    1.0
## 17        100       2  0   290    1   21     2    35       25     1    1.0
## 22        110       2  0   220    1   21     3    30       25     3    1.0
## 24        100       2  0   190    1   18     5    80       25     3    1.0
## 27        100       3  0     0    3   14     7   100       25     2    1.0
## 33        100       3  1   140    3   15     5    85       25     3    1.0
## 34        110       3  0   170    3   17     3    90       25     3    1.0
## 39        110       2  1   170    1   17     6    60      100     3    1.0
## 40        140       3  1   170    2   20     9    95      100     3    1.3
## 41        110       2  1   260    0   21     3    40       25     2    1.0
## 44        100       4  1     0    0   16     3    95       25     2    1.0
## 51         90       3  0   170    3   18     2    90       25     3    1.0
## 54        100       3  0   320    1   20     3    45      100     3    1.0
##      cups   rating cut2
## 10 0.67 53.31381    4
## 12 1.25 50.76500    4
## 16 1.00 41.44502    4
## 17 1.00 45.86332    4
## 22 1.00 46.89564    4
## 24 0.75 44.33086    4
## 27 0.80 58.34514    4
## 33 0.88 52.07690    4
## 34 0.25 53.37101    4
## 39 1.00 36.52368    4
## 40 0.75 36.47151    4
## 41 1.50 39.24111    4
## 44 1.00 54.85092    4
## 51 1.00 59.64284    4
## 54 1.00 41.50354    4
```

```
Centroid4 <- colMeans(Result[Result$cut2==4,])

Centroids <- rbind(Centroid1, Centroid2, Centroid3, Centroid4)
x2 <- as.data.frame(rbind(Centroids[, -14], df_B))
```

#Calculating the distanceance.

```

#df_A_distance
#Clustered_df_A <-cutree ()
#Clusters_A <-as.data.frame(cbind(df_A, Clustered_df_A))
#Identify clusters in each partition.
#Clust.1 <-colMeans (Clusters_A [Clusters_A$ Clustered_df_A == "1" ,])
#Centroid <-rbind(Clust.1, Clust.2, ..... )

distance1 <- get_dist(x2)
Matrix <- as.matrix(distance1)
data.frame <- data.frame(data=seq(1,nrow(df_B),1), Clusters = rep(0,nrow(df_B)))

for(i in 1:nrow(df_B))
{data.frame[i,2] <- which.min(Matrix[i+4, 1:4])}
data.frame

```

```

##      data Clusters
## 1      1         1
## 2      2         2
## 3      3         2
## 4      4         3
## 5      5         4
## 6      6         2
## 7      7         2
## 8      8         2
## 9      9         3
## 10     10        4
## 11     11        2
## 12     12        4
## 13     13        2
## 14     14        4
## 15     15        4
## 16     16        3
## 17     17        4
## 18     18        4
## 19     19        3

```

```

cbind(df2$SubGroup[51:74], data.frame$Clusters)

```

```
##           [,1]
## [1,]      1
## [2,]      2
## [3,]      2
## [4,]      3
## [5,]      4
## [6,]      2
## [7,]      2
## [8,]      2
## [9,]      3
## [10,]     4
## [11,]     2
## [12,]     4
## [13,]     2
## [14,]     4
## [15,]     4
## [16,]     3
## [17,]     4
## [18,]     4
## [19,]     3
```

```
table(df2$SubGroup[51:74] == data.frame$Clusters)
```

```
## < table of extent 0 >
```

```
#We can conclude that it is partially stable.
```

#Clustering Healthy CEREALS.

```
Healthy_CEREALS <- CEREALS
Healthy_CEREALS_na <- na.omit(Healthy_CEREALS)
Clust_healthy <- cbind(Healthy_CEREALS_na, subGroup)

Clust_healthy[Clust_healthy$subGroup==1,]
```

```
##           name mfr type calories protein fat sodium fiber carbo
## 1          100%_Bran  N  C      70      4  1   130   10    5
## 3           All-Bran  K  C      70      4  1   260    9    7
## 4 All-Bran_with_Extra_Fiber  K  C      50      4  0   140   14    8
##  sugars potass vitamins shelf weight cups  rating subGroup
## 1      6    280      25    3      1 0.33 68.40297      1
## 3      5    320      25    3      1 0.33 59.42551      1
## 4      0    330      25    3      1 0.50 93.70491      1
```

```
Clust_healthy[Clust_healthy$subGroup==2,]
```

##		name	mfr	type	calories	protein	fat	sodium		
## 2		100%_Natural_Bran	Q	C	120	3	5	15		
## 8		Basic_4	G	C	130	3	2	210		
## 14		Clusters	G	C	110	3	2	140		
## 20		Cracklin'_Oat_Bran	K	C	110	3	3	140		
## 23		Crispy_Wheat_&_Raisins	G	C	100	2	1	140		
## 28	Fruit_&_Fibre_Dates,_Walnuts,_and_Oats		P	C	120	3	2	160		
## 29		Fruitful_Bran	K	C	120	3	0	240		
## 35		Great_Grains_Pecan	P	C	120	3	3	75		
## 40		Just_Right_Fruit_&_Nut	K	C	140	3	1	170		
## 42		Life	Q	C	100	4	2	150		
## 45	Muesli_Raisins,_Dates,_&_Almonds		R	C	150	4	3	95		
## 46	Muesli_Raisins,_Peaches,_&_Pecans		R	C	150	4	3	150		
## 47		Mueslix_Crispy_Blend	K	C	160	3	2	150		
## 50		Nutri-Grain_Almond-Raisin	K	C	140	3	2	220		
## 52		Oatmeal_Raisin_Crisp	G	C	130	3	2	170		
## 53		Post_Nat._Raisin_Bran	P	C	120	3	1	200		
## 57		Quaker_Oat_Squares	Q	C	100	4	1	135		
## 59		Raisin_Bran	K	C	120	3	1	210		
## 60		Raisin_Nut_Bran	G	C	100	3	2	140		
## 71		Total_Raisin_Bran	G	C	140	3	1	190		
##	fiber	carbo	sugars	potass	vitamins	shelf	weight	cups	rating	subGroup
## 2	2.0	8.0	8	135	0	3	1.00	1.00	33.98368	2
## 8	2.0	18.0	8	100	25	3	1.33	0.75	37.03856	2
## 14	2.0	13.0	7	105	25	3	1.00	0.50	40.40021	2
## 20	4.0	10.0	7	160	25	3	1.00	0.50	40.44877	2
## 23	2.0	11.0	10	120	25	3	1.00	0.75	36.17620	2
## 28	5.0	12.0	10	200	25	3	1.25	0.67	40.91705	2
## 29	5.0	14.0	12	190	25	3	1.33	0.67	41.01549	2
## 35	3.0	13.0	4	100	25	3	1.00	0.33	45.81172	2
## 40	2.0	20.0	9	95	100	3	1.30	0.75	36.47151	2
## 42	2.0	12.0	6	95	25	2	1.00	0.67	45.32807	2
## 45	3.0	16.0	11	170	25	3	1.00	1.00	37.13686	2
## 46	3.0	16.0	11	170	25	3	1.00	1.00	34.13976	2
## 47	3.0	17.0	13	160	25	3	1.50	0.67	30.31335	2
## 50	3.0	21.0	7	130	25	3	1.33	0.67	40.69232	2
## 52	1.5	13.5	10	120	25	3	1.25	0.50	30.45084	2
## 53	6.0	11.0	14	260	25	3	1.33	0.67	37.84059	2
## 57	2.0	14.0	6	110	25	3	1.00	0.50	49.51187	2
## 59	5.0	14.0	12	240	25	2	1.33	0.75	39.25920	2
## 60	2.5	10.5	8	140	25	3	1.00	0.50	39.70340	2
## 71	4.0	15.0	14	230	100	3	1.50	1.00	28.59278	2

```
Clust_healthy[Clust_healthy$subGroup==3,]
```

##	name	mfr	type	calories	protein	fat	sodium	fiber	carbo
## 6	Apple_Cinnamon_Cheerios	G	C	110	2	2	180	1.5	10.5
## 7	Apple_Jacks	K	C	110	2	0	125	1.0	11.0
## 11	Cap'n'Crunch	Q	C	120	1	2	220	0.0	12.0
## 13	Cinnamon_Toast_Crunch	G	C	120	1	3	210	0.0	13.0
## 15	Cocoa_Puffs	G	C	110	1	1	180	0.0	12.0
## 18	Corn_Pops	K	C	110	1	0	90	1.0	13.0
## 19	Count_Chocula	G	C	110	1	1	180	0.0	12.0
## 25	Froot_Loops	K	C	110	2	1	125	1.0	11.0
## 26	Frosted_Flakes	K	C	110	1	0	200	1.0	14.0
## 30	Fruity_Pebbles	P	C	110	1	1	135	0.0	13.0
## 31	Golden_Crisp	P	C	100	2	0	45	0.0	11.0
## 32	Golden_Grahams	G	C	110	1	1	280	0.0	15.0
## 36	Honey_Graham_Ohs	Q	C	120	1	2	220	1.0	12.0
## 37	Honey_Nut_Cheerios	G	C	110	3	1	250	1.5	11.5
## 38	Honey-comb	P	C	110	1	0	180	0.0	14.0
## 43	Lucky_Charm	G	C	110	2	1	180	0.0	12.0
## 48	Multi-Grain_Cheerios	G	C	100	2	1	220	2.0	15.0
## 49	Nut&Honey_Crunch	K	C	120	2	1	190	0.0	15.0
## 67	Smacks	K	C	110	2	1	70	1.0	9.0
## 74	Trix	G	C	110	1	1	140	0.0	13.0
## 77	Wheaties_Honey_Gold	G	C	110	2	1	200	1.0	16.0
##	sugars	potass	vitamins	shelf	weight	cups	rating	subGroup	
## 6	10	70	25	1	1	0.75	29.50954	3	
## 7	14	30	25	2	1	1.00	33.17409	3	
## 11	12	35	25	2	1	0.75	18.04285	3	
## 13	9	45	25	2	1	0.75	19.82357	3	
## 15	13	55	25	2	1	1.00	22.73645	3	
## 18	12	20	25	2	1	1.00	35.78279	3	
## 19	13	65	25	2	1	1.00	22.39651	3	
## 25	13	30	25	2	1	1.00	32.20758	3	
## 26	11	25	25	1	1	0.75	31.43597	3	
## 30	12	25	25	2	1	0.75	28.02576	3	
## 31	15	40	25	1	1	0.88	35.25244	3	
## 32	9	45	25	2	1	0.75	23.80404	3	
## 36	11	45	25	2	1	1.00	21.87129	3	
## 37	10	90	25	1	1	0.75	31.07222	3	
## 38	11	35	25	1	1	1.33	28.74241	3	
## 43	12	55	25	2	1	1.00	26.73451	3	
## 48	6	90	25	1	1	1.00	40.10596	3	
## 49	9	40	25	2	1	0.67	29.92429	3	
## 67	15	40	25	2	1	0.75	31.23005	3	
## 74	12	25	25	2	1	1.00	27.75330	3	
## 77	8	60	25	1	1	0.75	36.18756	3	

```
Clust_healthy[Clust_healthy$subGroup==4,]
```

##		name	mfr	type	calories	protein	fat	sodium	fiber	carbo
## 9		Bran_Chex	R	C	90	2	1	200	4	15
## 10		Bran_Flakes	P	C	90	3	0	210	5	13
## 12		Cheerios	G	C	110	6	2	290	2	17
## 16		Corn_Chex	R	C	110	2	0	280	0	22
## 17		Corn_Flakes	K	C	100	2	0	290	1	21
## 22		Crispix	K	C	110	2	0	220	1	21
## 24		Double_Chex	R	C	100	2	0	190	1	18
## 33		Grape_Nuts_Flakes	P	C	100	3	1	140	3	15
## 34		Grape-Nuts	P	C	110	3	0	170	3	17
## 39	Just_Right_Crunchy__	Nuggets	K	C	110	2	1	170	1	17
## 41		Kix	G	C	110	2	1	260	0	21
## 51		Nutri-grain_Wheat	K	C	90	3	0	170	3	18
## 54		Product_19	K	C	100	3	0	320	1	20
## 62		Rice_Chex	R	C	110	1	0	240	0	23
## 63		Rice_Krispies	K	C	110	2	0	290	0	22
## 68		Special_K	K	C	110	6	0	230	1	16
## 70		Total_Corn_Flakes	G	C	110	2	1	200	0	21
## 72		Total_Whole_Grain	G	C	100	3	1	200	3	16
## 73		Triples	G	C	110	2	1	250	0	21
## 75		Wheat_Chex	R	C	100	3	1	230	3	17
## 76		Wheaties	G	C	100	3	1	200	3	17
##	sugars	potass	vitamins	shelf	weight	cups	rating	subGroup		
## 9	6	125	25	1	1	0.67	49.12025	4		
## 10	5	190	25	3	1	0.67	53.31381	4		
## 12	1	105	25	1	1	1.25	50.76500	4		
## 16	3	25	25	1	1	1.00	41.44502	4		
## 17	2	35	25	1	1	1.00	45.86332	4		
## 22	3	30	25	3	1	1.00	46.89564	4		
## 24	5	80	25	3	1	0.75	44.33086	4		
## 33	5	85	25	3	1	0.88	52.07690	4		
## 34	3	90	25	3	1	0.25	53.37101	4		
## 39	6	60	100	3	1	1.00	36.52368	4		
## 41	3	40	25	2	1	1.50	39.24111	4		
## 51	2	90	25	3	1	1.00	59.64284	4		
## 54	3	45	100	3	1	1.00	41.50354	4		
## 62	2	30	25	1	1	1.13	41.99893	4		
## 63	3	35	25	1	1	1.00	40.56016	4		
## 68	3	55	25	1	1	1.00	53.13132	4		
## 70	3	35	100	3	1	1.00	38.83975	4		
## 72	3	110	100	3	1	1.00	46.65884	4		
## 73	3	60	25	3	1	0.75	39.10617	4		
## 75	3	115	25	1	1	0.67	49.78744	4		
## 76	3	110	25	1	1	1.00	51.59219	4		

#Mean ratings to determine the best cluster.

```
mean(Clust_healthy[Clust_healthy$subGroup==1,"rating"])
```

```
## [1] 73.84446
```

```
mean(Clust_healthy[Clust_healthy$subGroup==2,"rating"])
```

```
## [1] 38.26161
```

```
mean(Clust_healthy[Clust_healthy$subGroup==3,"rating"])
```

```
## [1] 28.84825
```

```
mean(Clust_healthy[Clust_healthy$subGroup==4,"rating"])
```

```
## [1] 46.46513
```

*#We can conclude that cluster 1 should be picked since it has the highest value. Cluster 1 can therefore be regarded as a Healthy Cluster.*