

# Assignment\_5

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4/17/2022

```
#Importing required libraries
```

```
library(cluster)
library(caret)
```

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

```
## Warning in register(): Can't find generic 'scale_type' in package ggplot2 to
## register S3 method.
```

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

```
library(dendextend)
```

```
## Warning: package 'dendextend' was built under R version 4.1.3
```

```
##
## -----
## Welcome to dendextend version 1.15.2
## 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/issues
## 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)
```

```
## Warning: package 'factoextra' was built under R version 4.1.3
```

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

```
#Importing dataset
```

```
Cereals<- read.csv("C:/Machine Learning - 1/64060_hbhogadi/Assignment_5/Cereals.csv")
```

```
Data_cereals <- data.frame(Cereals[,4:16])
```

```
#Preprocessing the data
```

```
Data_cereals <- na.omit(Data_cereals)
```

```
#Data Normalization
```

```
Data_cereals_normalized <- scale(Data_cereals)
```

```
#Applying hierarchical clustering to the data using Euclidean distance to the normalize measurements.
```

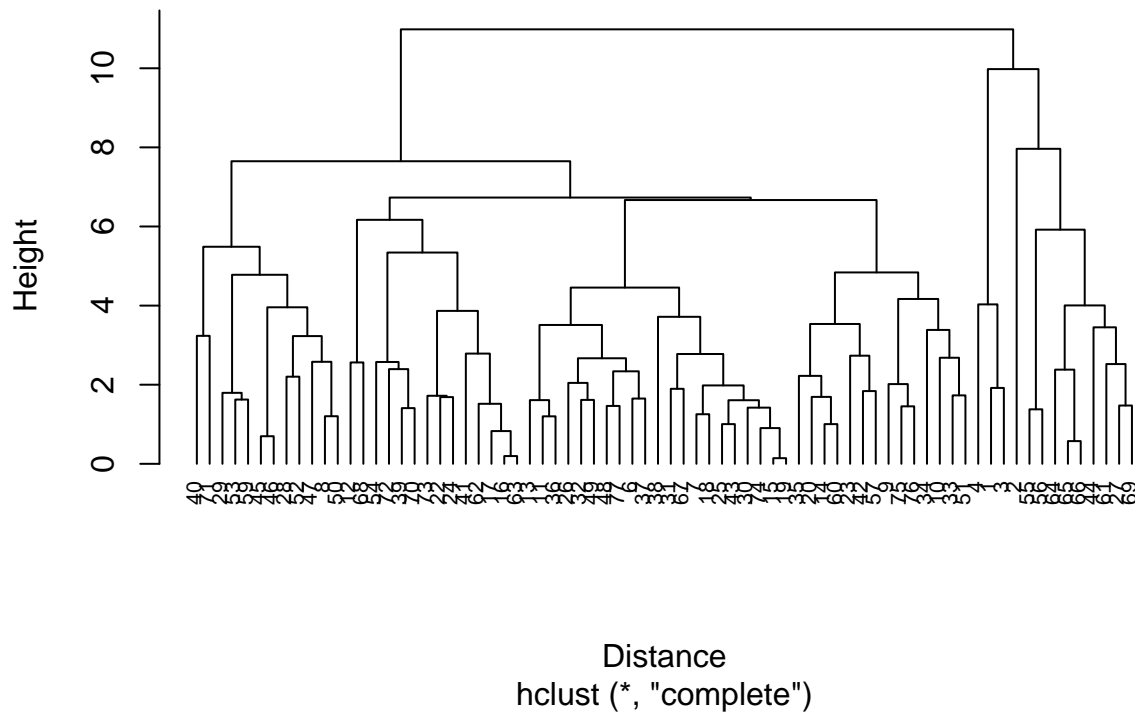
```
Distance <- dist(Data_cereals_normalized, method = "euclidean")
```

```
hierarchial.clust_complete <- hclust(Distance, method = "complete")
```

```
#Plotting the dendogram
```

```
plot(hierarchial.clust_complete, cex = 0.7, hang = -1)
```

## Cluster Dendrogram



*#Using agnes function to perform clustering with single linkage, complete linkage, average linkage and*

```
hierarchial.clust_single <- agnes(Data_cereals_normalized, method = "single")
hierarchial.clust_complete <- agnes(Data_cereals_normalized, method = "complete")
hierarchial.clust_average <- agnes(Data_cereals_normalized, method = "average")
hierarchial.clust_ward <- agnes(Data_cereals_normalized, method = "ward")
```

*#Single Linkage vs Complete Linkage vs Average Linkage vs Ward*

```
print(hierarchial.clust_single$ac)
```

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

```
print(hierarchial.clust_complete$ac)
```

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

```
print(hierarchial.clust_average$ac)
```

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

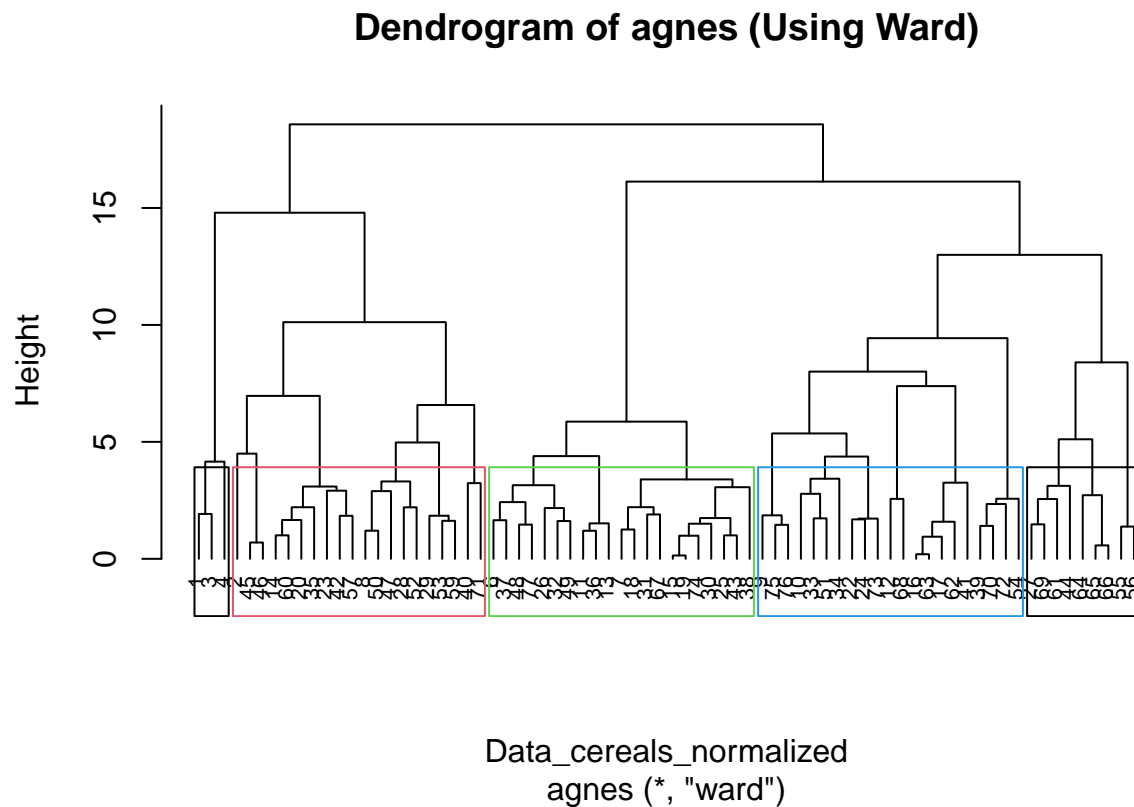
```
print(hierarchial.clust_ward$ac)
```

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

```
#Since WARD method has the highest value of 0.9046042, we will consider it.
```

```
#(2) Choosing the clusters:
```

```
pltree(hierarchial.clust_ward, cex = 0.7, hang = -1, main = "Dendrogram of agnes (Using Ward)")
rect.hclust(hierarchial.clust_ward, k = 5, border = 1:4)
```



```
Cluster1 <- cutree(hierarchial.clust_ward, k=5)
```

```
dataframe2 <- as.data.frame(cbind(Data_cereals_normalized,Cluster1))
```

```
#We will choose 5 clusters after observing the distance.
```

```
#Commenting on the structure of the clusters and on their stability
```

```
#Creating Partitions
```

```

set.seed(123)
Partition1 <- Data_cereals[1:50,]
Partition2 <- Data_cereals[51:74,]

#Performing Hierarchical Clustering, considering k = 5.

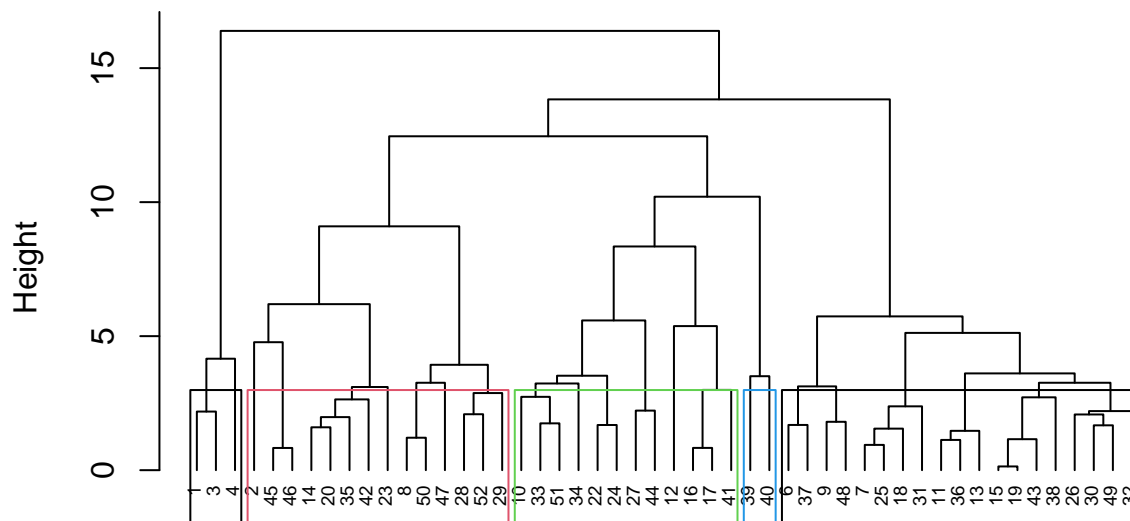
AG_single <- agnes(scale(Partition1), method = "single")
AG_complete <- agnes(scale(Partition1), method = "complete")
AG_average <- agnes(scale(Partition1), method = "average")
AG_ward <- agnes(scale(Partition1), method = "ward")
cbind(single=AG_single$ac , complete=AG_complete$ac , average= AG_average$ac , ward= AG_ward$ac)

##          single complete average      ward
## [1,] 0.6393338 0.8138238 0.7408904 0.8764323

pltree(AG_ward, cex = 0.6, hang = -1, main = "Dendrogram of Agnes with Partitioned Data (Using Ward)")
rect.hclust(AG_ward, k = 5, border = 1:4)

```

## Dendrogram of Agnes with Partitioned Data (Using Ward)



scale(Partition1)  
agnes (\*, "ward")

```
cut_2 <- cutree(AG_ward, k = 5)
```

```
#Calculating the centroids.
```

```
result <- as.data.frame(cbind(Partition1, cut_2))  
result[result$cut_2==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 cut_2  
## 1 0.33 68.40297     1  
## 3 0.33 59.42551     1  
## 4 0.50 93.70491     1
```

```
centroid_1 <- colMeans(result[result$cut_2==1,])  
result[result$cut_2==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  
##   cups   rating cut_2  
## 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
```

```
centroid_2 <- colMeans(result[result$cut_2==2,])  
result[result$cut_2==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	cut_2
## 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

```
centroid_3 <- colMeans(result[result$cut_2==3,])
result[result$cut_2==4,]
```

##	calories	protein	fat	sodium	fiber	carbo	sugars	potass	vitamins	shelf	weight
## 10	90	3	0	210	5	13	5	190	25	3	1
## 12	110	6	2	290	2	17	1	105	25	1	1
## 16	110	2	0	280	0	22	3	25	25	1	1
## 17	100	2	0	290	1	21	2	35	25	1	1
## 22	110	2	0	220	1	21	3	30	25	3	1
## 24	100	2	0	190	1	18	5	80	25	3	1
## 27	100	3	0	0	3	14	7	100	25	2	1
## 33	100	3	1	140	3	15	5	85	25	3	1
## 34	110	3	0	170	3	17	3	90	25	3	1

```
## 41      110      2  1    260      0    21      3      40      25      2      1
## 44      100      4  1      0      0    16      3      95      25      2      1
## 51       90      3  0    170      3    18      2      90      25      3      1
##      cups   rating cut_2
## 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
## 41 1.50 39.24111      4
## 44 1.00 54.85092      4
## 51 1.00 59.64284      4
```

```
centroid_4 <- colMeans(result[result$cut_2==4,])
centroids <- rbind(centroid_1, centroid_2, centroid_3, centroid_4)
x2 <- as.data.frame(rbind(centroids[, -14], Partition2))

#Calculating the Distance

Distance_1 <- get_dist(x2)
Matrix_1 <- as.matrix(Distance_1)
dataframe1 <- data.frame(data=seq(1,nrow(Partition2),1), Clusters = rep(0,nrow(Partition2)))
for(i in 1:nrow(Partition2))
{dataframe1[i,2] <- which.min(Matrix_1[i+4, 1:4])}
dataframe1
```

```
##      data Clusters
## 1      1          1
## 2      2          4
## 3      3          3
## 4      4          2
## 5      5          2
## 6      6          1
## 7      7          2
## 8      8          2
## 9      9          3
## 10     10         3
## 11     11         2
## 12     12         2
## 13     13         2
## 14     14         3
## 15     15         4
## 16     16         2
## 17     17         3
## 18     18         2
## 19     19         4
## 20     20         4
## 21     21         3
```



```
## 22 22 4
## 23 23 4
## 24 24 3
```

```
cbind(dataframe2$Cluster1[51:74], dataframe1$Clusters)
```

```
##      [,1] [,2]
## [1,]    2    1
## [2,]    4    4
## [3,]    5    3
## [4,]    5    2
## [5,]    2    2
## [6,]    2    1
## [7,]    2    2
## [8,]    5    2
## [9,]    4    3
## [10,]   4    3
## [11,]   5    2
## [12,]   5    2
## [13,]   5    2
## [14,]   3    3
## [15,]   4    4
## [16,]   5    2
## [17,]   4    3
## [18,]   2    2
## [19,]   4    4
## [20,]   4    4
## [21,]   3    3
## [22,]   4    4
## [23,]   4    4
## [24,]   3    3
```

```
table(dataframe2$Cluster1[51:74] == dataframe1$Clusters)
```

```
##
## FALSE  TRUE
##     12    12
```

*#We can say that the model is partially stable as we are getting 12 FALSE and 12 TRUE*

*#3) The elementary public schools would like to choose a set of cereals to include in their daily cafet*

*#Clustering Healthy Cereals.*

```
Healthy_Cereals <- Cereals
Healthy_Cereals_new <- na.omit(Healthy_Cereals)
HealthyClust <- cbind(Healthy_Cereals_new, Cluster1)
HealthyClust[HealthyClust$Cluster1==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 Cluster1
## 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
```

```
HealthyClust[HealthyClust$Cluster1==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 Cluster1
## 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
```

```
HealthyClust[HealthyClust$Cluster1==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	Cluster1		
## 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		

```
HealthyClust[HealthyClust$Cluster1==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	Cluster1	
## 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(HealthyClust[HealthyClust$Cluster1==1,"rating"])
```

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

```
mean(HealthyClust[HealthyClust$Cluster1==2,"rating"])
```

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

```
mean(HealthyClust[HealthyClust$Cluster1==3,"rating"])
```

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

```
mean(HealthyClust[HealthyClust$Cluster1==4,"rating"])
```

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

*#We can consider cluster 1 since mean ratings of the cluster1 is the highest(i.e. 73.84446).*