

CAT 3 PROJECT:

ANALYSIS OF PERSONALITY AND SOCIAL BEHAVIOUR

Problem statement:

To classify general public of different age groups based on the analysis of 'Personality and Social Behaviour'.

Methods:

1. Elbow method of clustering.
2. Kmeans clustering.
3. Hierarchical clustering
4. PCA

Tools and libraries:

1. R studio.
2. library(dplyr)
3. library(factoextra)
4. library(ggplot2)
5. library(ggpubr)
6. library(cluster)

Code:

```
library(dplyr)
```

```
library(factoextra)
```

```
library(ggplot2)
```

```
library(ggpubr)
```

```
library(cluster)
```

```
dataset1=read.csv("C:\\Users\\Navika M S\\OneDrive\\Documents\\Semester 4\\PA  
lab\\Revision material\\Personality and Social Behavior Dataset For Analysis.csv")
```

```
print(dataset1)
```

```
summary(dataset1)
```

```
datasetm=read.csv("C:\\Users\\Navika M S\\OneDrive\\Documents\\Semester 4\\PA  
lab\\male.csv")
```

```
print(datasetm)
```

```
summary(datasetm)
```

```
datasetf=read.csv("C:\\Users\\Navika M S\\OneDrive\\Documents\\Semester 4\\PA  
lab\\female.csv")
```

```
print(datasetf)
```

```
summary(datasetf)
```

```

fviz_nbclust(dataset, kmeans, method="wss")
labs(subtitle="Elbow Method")
results11<-kmeans(dataset1,5)
results11
results11$size
results11$cluster
dataset1[,1:7] <-scale(dataset1[,1:7])
plot(dataset1[c("AGE","MIND")],col=results11$cluster)
points(results11$centers,pch=2,col="red")
plot(dataset1[c("AGE","ENERGY")],col=results11$cluster)
points(results11$centers,pch=2,col="red")
plot(dataset1[c("AGE","NATURE")],col=results11$cluster)
points(results11$centers,pch=2,col="red")
plot(dataset1[c("AGE","TACTICS")],col=results11$cluster)
points(results11$centers,pch=2,col="red")
plot(dataset1[c("AGE","IDENTITY")],col=results11$cluster)
points(results11$centers,pch=2,col="red")
clusplot(dataset1,results11$cluster)
sil <- silhouette(results11$cluster, dist(dataset1))
fviz_silhouette(sil)
#EUCLIDEAN
data.exc1<-dist(dataset1,method="euclidean")
round(as.matrix(data.exc1)[1:7,1:7],1)
# Use hcut() which compute hclust and cut the tree
hc.cut <- hcut(dataset1, k = 5, hc_method = "complete")
# Visualize dendrogram
fviz_dend(hc.cut, show_labels = TRUE, rect = TRUE)
res.pca <- prcomp(dataset1, scale = TRUE)
fviz_eig(res.pca)

```

```

fviz_pca_ind(res.pca,
              col.ind = "cos2", # Color by the quality of representation
              gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
              repel = TRUE    # Avoid text overlapping
)

# Dimension reduction using PCA
res.pca <- prcomp(dataset1, scale = TRUE)
res.pca

# Coordinates of individuals
ind.coord <- as.data.frame(get_pca_ind(res.pca)$coord)

# Add clusters obtained using the K-means algorithm
ind.coord$cluster <- factor(results11$cluster)

# Add Species groups from the original data sett
ind.coord$Species <- df$Species

# Data inspection
head(ind.coord)

# Percentage of variance explained by dimensions
eigenvalue <- round(get_eigenvalue(res.pca), 1)
variance.percent <- eigenvalue$variance.percent
head(eigenvalue)

ggscatter(
  ind.coord, x = "Dim.1", y = "Dim.2",
  color = "cluster", palette = "npg", ellipse = TRUE, ellipse.type = "convex",
  shape = "circle", size = 1.5, legend = "right", ggtheme = theme_bw(),
  xlab = paste0("Dim 1 (", variance.percent[1], "% )" ),
  ylab = paste0("Dim 2 (", variance.percent[2], "% )" )
) +
  stat_mean(aes(color = cluster), size = 4)

```

Output:

	GENDER	AGE	MIND	ENERGY	NATURE	TACTICS	IDENTITY
1	1	2	3	3	2	3	2
2	2	1	3	3	2	3	3
3	2	2	2	1	1	2	3
4	2	2	3	3	3	3	3
5	2	2	3	2	3	3	2
6	1	5	2	3	2	3	3
7	1	3	2	3	2	3	3
8	1	2	3	3	2	3	2
9	1	2	3	3	2	3	3
10	2	2	2	2	2	2	2
11	1	2	2	2	1	3	2
12	1	2	4	2	3	3	3
13	1	2	2	3	2	2	3
14	2	4	2	3	2	2	3
15	2	2	2	2	1	2	1
16	1	2	3	3	2	2	2
17	2	1	2	2	3	2	1
18	1	2	3	3	2	2	3
19	1	2	3	3	2	3	2
20	2	1	3	2	2	2	2
21	1	2	2	3	2	3	3
22	1	1	2	3	3	3	3
23	2	2	2	3	2	3	3
24	1	2	2	3	2	3	3
25	1	4	2	3	2	3	3
26	2	1	3	3	2	2	3
27	2	2	3	3	2	2	3
28	1	2	3	3	2	2	3
29	2	1	3	3	2	3	3
30	2	1	3	2	2	2	2
31	1	2	3	2	2	2	2
32	1	2	1	1	1	1	2
33	1	2	1	1	1	1	2
34	2	1	2	2	3	2	3
35	1	1	2	2	2	3	2
36	2	2	3	3	2	3	2
37	1	4	3	3	2	3	3
38	1	4	2	1	1	2	3
39	1	4	3	3	3	3	3
40	1	2	3	2	3	3	2
41	1	4	2	3	2	3	3
42	1	4	2	3	2	3	3
43	2	4	2	2	2	2	2

```
> summary(dataset1)
```

GENDER	AGE	MIND	ENERGY	NATURE	TACTICS	IDENTITY
Min. :1.00	Min. :1.00	Min. :1.00	Min. :1.00	Min. :1.00	Min. :1.00	Min. :1.00
1st Qu.:1.00	1st Qu.:2.00	1st Qu.:2.00	1st Qu.:2.00	1st Qu.:2.00	1st Qu.:2.00	1st Qu.:2.00
Median :1.00	Median :2.00	Median :2.00	Median :3.00	Median :2.00	Median :3.00	Median :3.00
Mean :1.38	Mean :2.48	Mean :2.48	Mean :2.52	Mean :2.02	Mean :2.52	Mean :2.52
3rd Qu.:2.00	3rd Qu.:4.00	3rd Qu.:3.00	3rd Qu.:3.00	3rd Qu.:2.00	3rd Qu.:3.00	3rd Qu.:3.00
Max. :2.00	Max. :5.00	Max. :4.00	Max. :3.00	Max. :3.00	Max. :3.00	Max. :3.00

```

> fviz_nbclust(dataset, kmeans, method="wss")
> labs(subtitle="Elbow Method")
$subtitle
[1] "Elbow Method"

attr(,"class")
[1] "labels"
>
> results11<-kmeans(dataset1,5)
> results11
K-means clustering with 5 clusters of sizes 19, 10, 3, 5, 13

Cluster means:
  GENDER    AGE    MIND  ENERGY  NATURE  TACTICS  IDENTITY
1 1.368421 1.789474 2.789474 2.947368 2.157895 2.684211 2.736842
2 1.600000 1.500000 2.500000 2.000000 2.400000 2.300000 2.000000
3 1.333333 4.333333 2.000000 1.666667 1.000000 2.333333 2.000000
4 1.400000 2.000000 1.600000 1.400000 1.000000 1.800000 2.000000
5 1.230769 4.000000 2.461538 2.923077 2.153846 2.769231 2.923077

Clustering vector:
[1] 1 1 4 1 2 5 5 1 1 2 4 1 1 5 4 1 2 1 1 2 1 1 1 1 5 1 1 1 1 2 2 4 4 2 2 1 5 3 5 2 5 5 5 5 2 3 5 5 5 3

Within cluster sum of squares by cluster:
[1] 24.000000 13.900000 4.666667 8.400000 15.384615
(between_SS / total_SS = 61.7 %)

Available components:
[1] "cluster"      "centers"      "totss"        "withinss"     "tot.withinss" "betweenss"    "size"         "iter"
[9] "ifault"
> results11$size
[1] 19 10 3 5 13
> results11$cluster
[1] 1 1 4 1 2 5 5 1 1 2 4 1 1 5 4 1 2 1 1 2 1 1 1 1 5 1 1 1 1 2 2 4 4 2 2 1 5 3 5 2 5 5 5 5 2 3 5 5 5 3
>
> dataset1[,1:7] <-scale(dataset1[,1:7])
>
> plot(dataset1[c("AGE", "MIND")],col=results11$cluster)
> points(results11$centers,pch=2,col="red")
>
> plot(dataset1[c("AGE", "ENERGY")],col=results11$cluster)
> points(results11$centers,pch=2,col="red")
>
> plot(dataset1[c("AGE", "NATURE")],col=results11$cluster)
> points(results11$centers,pch=2,col="red")
>
> plot(dataset1[c("AGE", "TACTICS")],col=results11$cluster)
> points(results11$centers,pch=2,col="red")
>
> plot(dataset1[c("AGE", "IDENTITY")],col=results11$cluster)
> points(results11$centers,pch=2,col="red")
>
> clusplot(dataset1,results11$cluster)
>
> sil <- silhouette(results11$cluster, dist(dataset1))
> fviz_silhouette(sil)
  cluster size ave.sil.width
1         1   19         0.16
2         2   10         0.17
3         3    3         0.05
4         4    5         0.09
5         5   13         0.25
>
> #EUCLIDEAN
> data.exc1<-dist(dataset1,method="euclidean")
> round(as.matrix(data.exc1)[1:7,1:7],1)
  1    2    3    4    5    6    7
1 0.0  2.7  5.0  3.1  3.1  3.4  2.4
2 2.7  0.0  4.3  1.9  2.9  4.2  3.1
3 5.0  4.3  0.0  5.1  4.7  5.1  4.5
4 3.1  1.9  5.1  0.0  2.2  4.0  3.2
5 3.1  2.9  4.7  2.2  0.0  4.6  3.9
6 3.4  4.2  5.1  4.0  4.6  0.0  1.7
7 2.4  3.1  4.5  3.2  3.9  1.7  0.0

```

```

>
> # Use hcut() which compute hclust and cut the tree
> hc.cut <- hcut(dataset1, k = 5, hc_method = "complete")
> # Visualize dendrogram
> fviz_dend(hc.cut, show_labels = TRUE, rect = TRUE)
/
> res.pca <- prcomp(dataset1, scale = TRUE)
> fviz_eig(res.pca)
>
> fviz_pca_ind(res.pca,
+             col.ind = "cos2", # Color by the quality of representation
+             gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
+             repel = TRUE      # Avoid text overlapping
+ )
> # Dimension reduction using PCA
> res.pca <- prcomp(dataset1, scale = TRUE)
> res.pca
Standard deviations (1, .., p=7):
[1] 1.5682067 1.2390121 0.8990680 0.8837855 0.7506537 0.6708209 0.6345826

Rotation (n x k) = (7 x 7):

```

	PC1	PC2	PC3	PC4	PC5	PC6	PC7
GENDER	0.07723116	-0.53503227	0.6798017	-0.4357959	0.05765008	0.1463238	-0.17604903
AGE	-0.07109846	0.60733950	0.5411088	0.1614479	0.48867591	0.2138506	0.15060994
MIND	-0.42026964	-0.35851515	0.0765270	0.3055440	0.44899894	-0.6234309	0.07321646
ENERGY	-0.49033796	0.12342216	0.2164341	-0.2543599	-0.50662508	-0.1330876	0.59867498
NATURE	-0.44553406	-0.32181719	-0.2968844	0.0360027	0.28128147	0.6946220	0.21651221
TACTICS	-0.48913138	0.07826014	0.2038592	0.3845004	-0.39088833	0.1431537	-0.62605118
IDENTITY	-0.36629546	0.30236089	-0.2503436	-0.6905242	0.26017368	-0.1532518	-0.37912186

```

> # Coordinates of individuals
> ind.coord <- as.data.frame(get_pca_ind(res.pca)$coord)
> # Add clusters obtained using the K-means algorithm
> ind.coord$cluster <- factor(results11$cluster)
> # Add Species groups from the original data sett
> ind.coord$Species <- df$Species
> # Data inspection
> head(ind.coord)

```

	Dim.1	Dim.2	Dim.3	Dim.4	Dim.5	Dim.6	Dim.7	cluster
1	-0.8126299	-0.2089837	-0.1335216	1.23081540	-0.8114864	-0.5758419	0.37416620	1
2	-1.1914308	-1.3215828	0.3875720	-0.91897494	-0.6836237	-0.7078555	-0.72962939	1
3	2.5156771	-0.2233651	0.2101749	-1.19229918	0.7987185	-0.5777534	-1.85581985	4
4	-2.0084231	-1.3545243	0.3409609	-0.72124750	0.2075711	0.6530279	-0.23442940	1
5	-0.6535140	-2.0377818	0.4138279	0.79660870	0.5675637	1.1084344	-0.54311543	2
6	-0.9394401	2.3791499	0.7136231	0.04349154	0.1577616	0.6816165	0.02577351	5

```

>
> # Percentage of variance explained by dimensions
> eigenvalue <- round(get_eigenvalue(res.pca), 1)
> variance.percent <- eigenvalue$variance.percent
> head(eigenvalue)

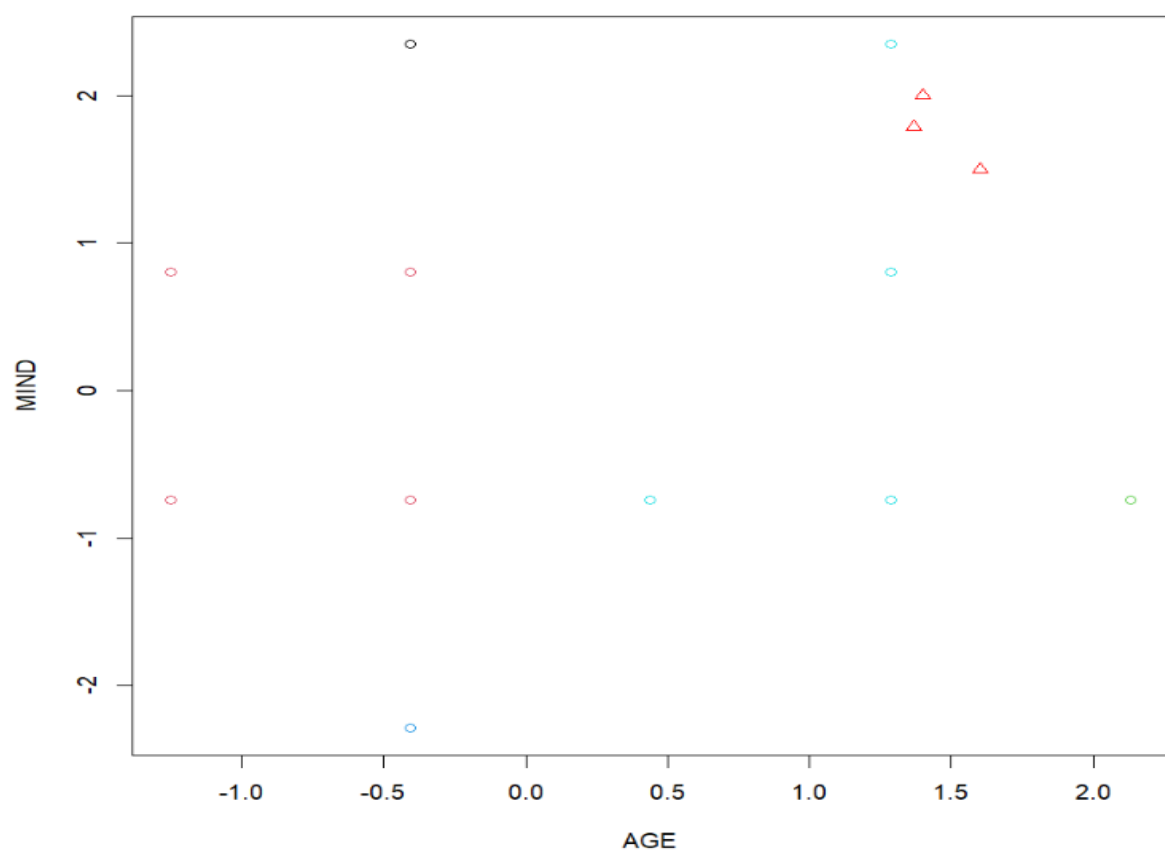
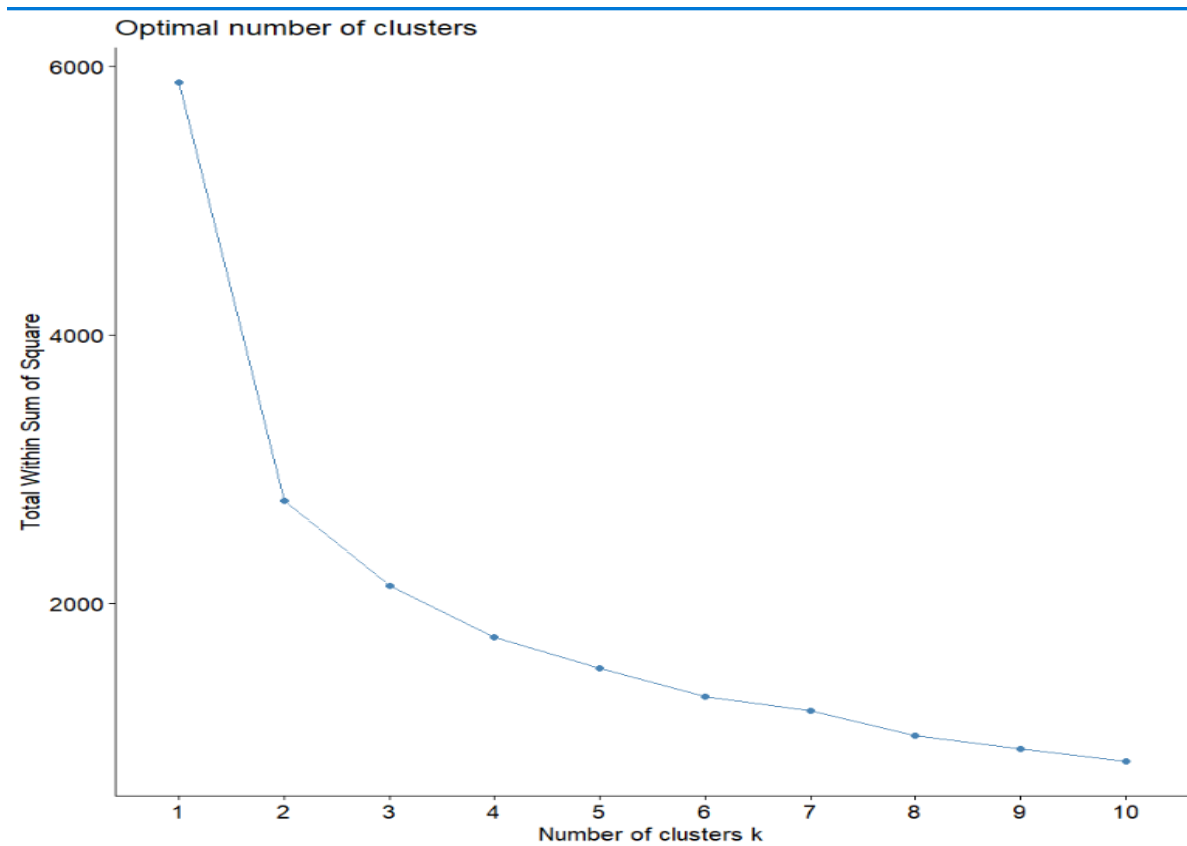
```

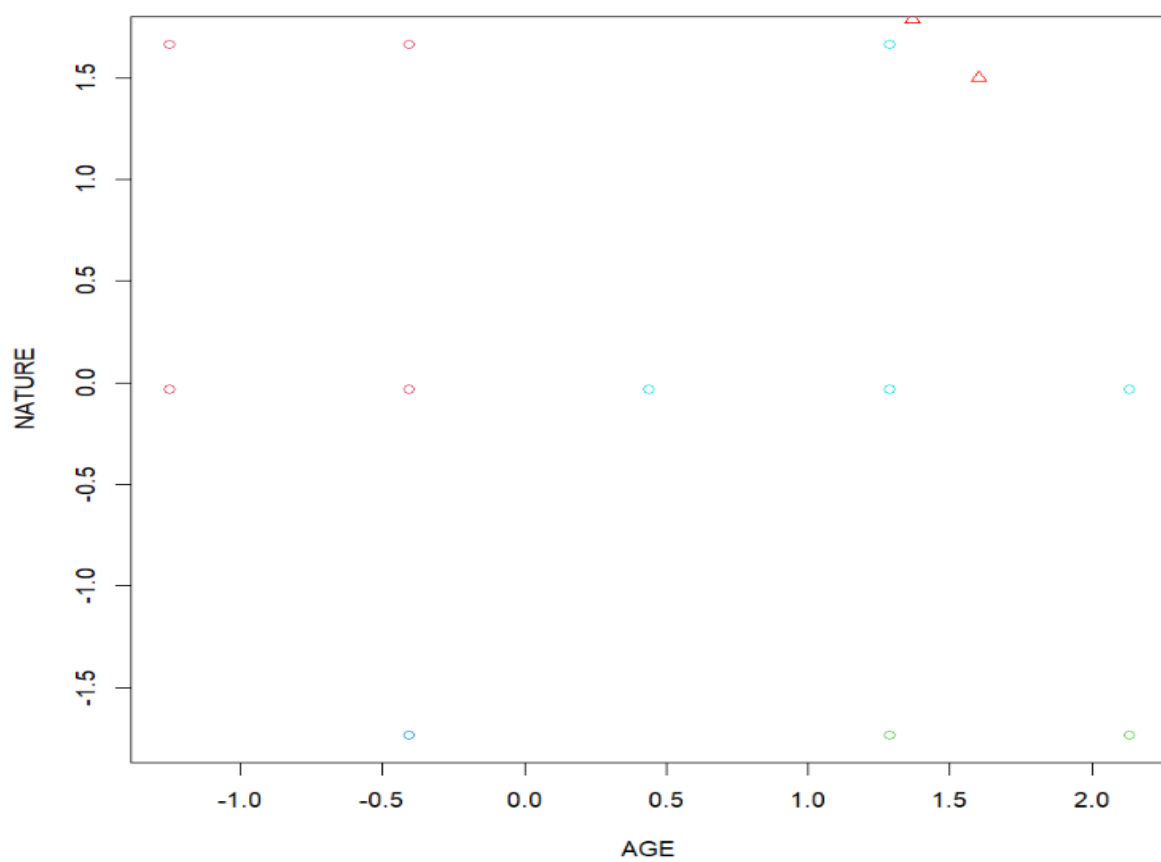
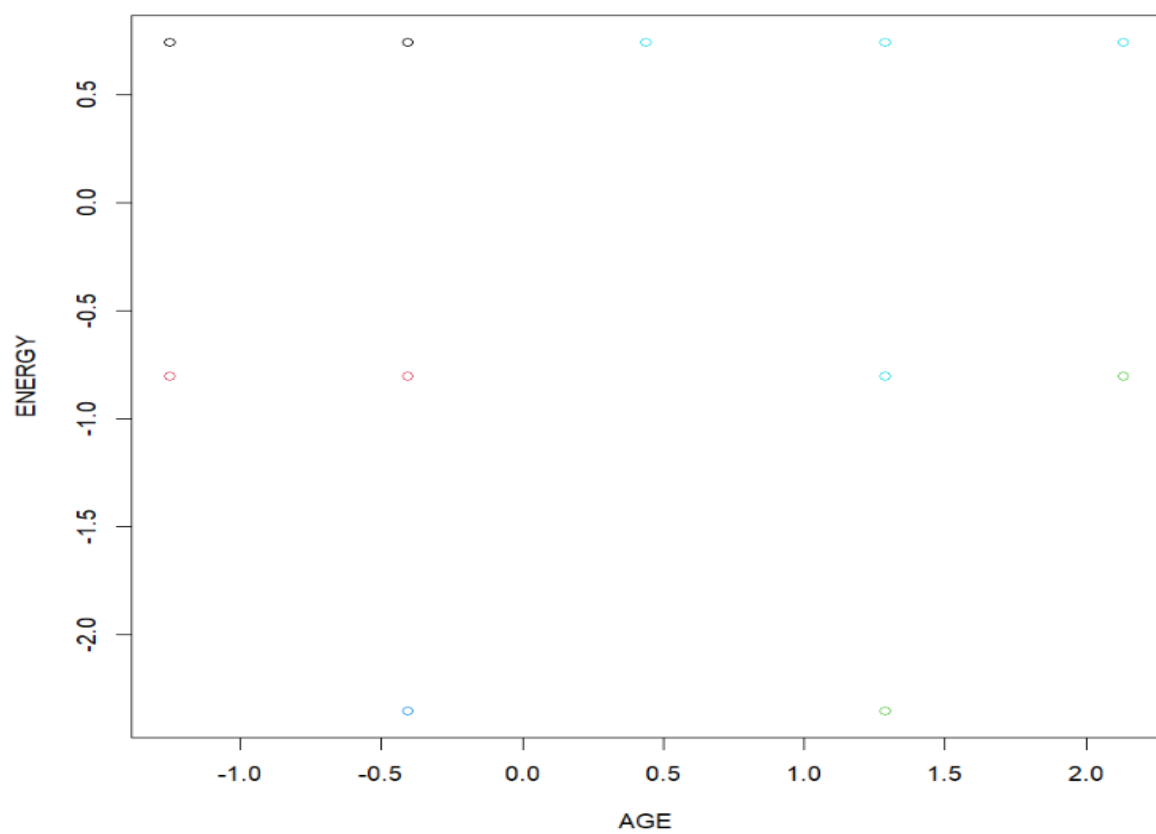
	eigenvalue	variance.percent	cumulative.variance.percent
Dim.1	2.5	35.1	35.1
Dim.2	1.5	21.9	57.1
Dim.3	0.8	11.5	68.6
Dim.4	0.8	11.2	79.8
Dim.5	0.6	8.0	87.8
Dim.6	0.5	6.4	94.2

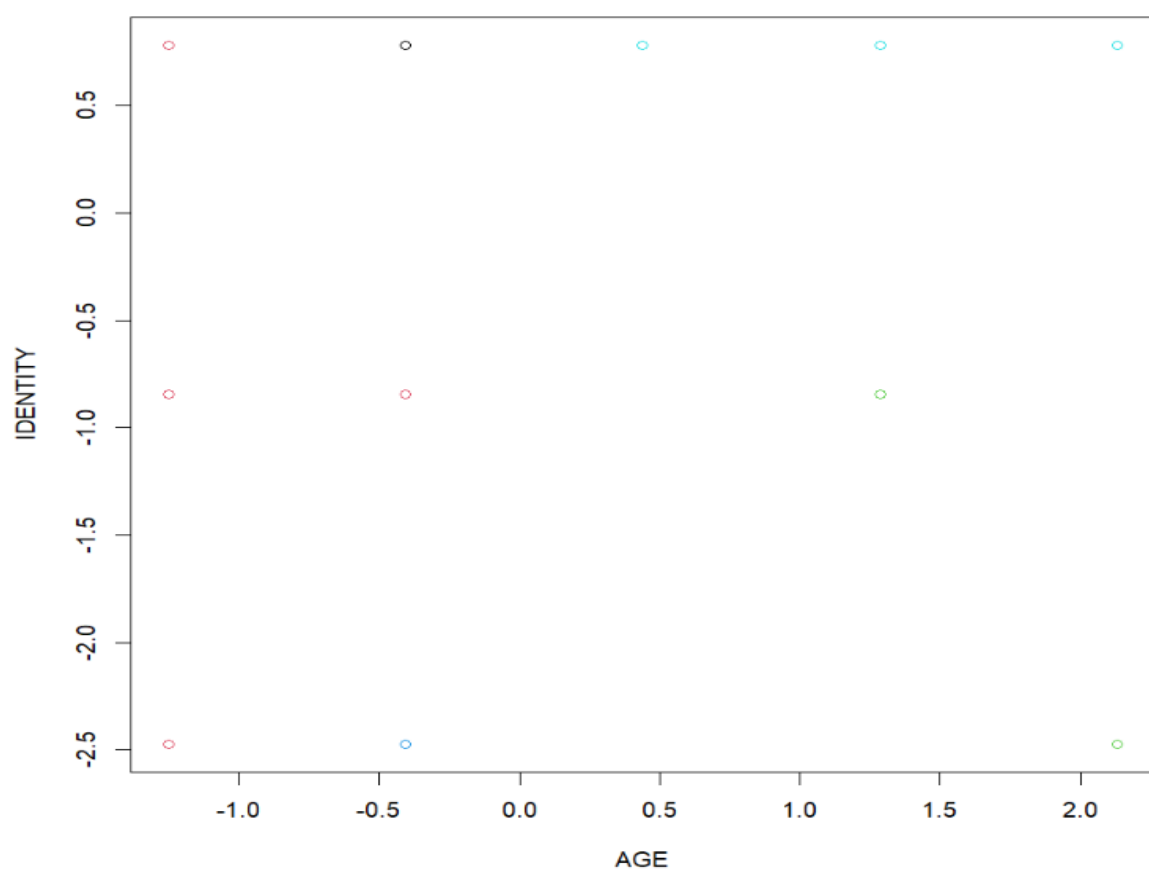
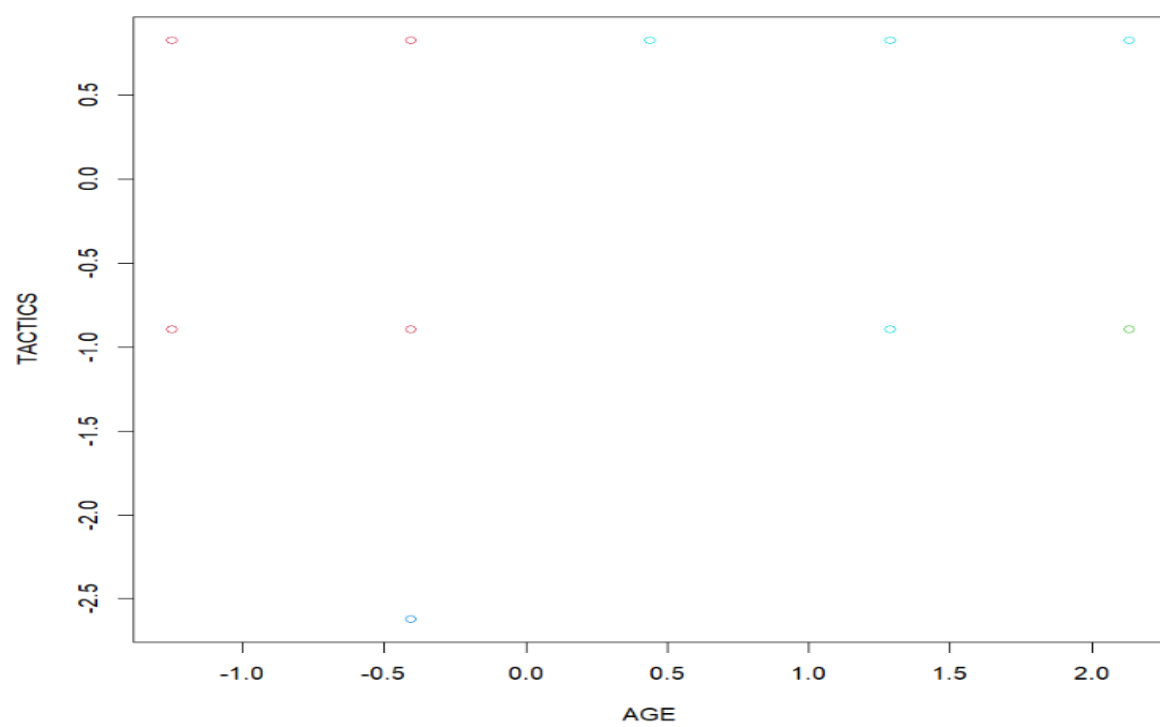
```

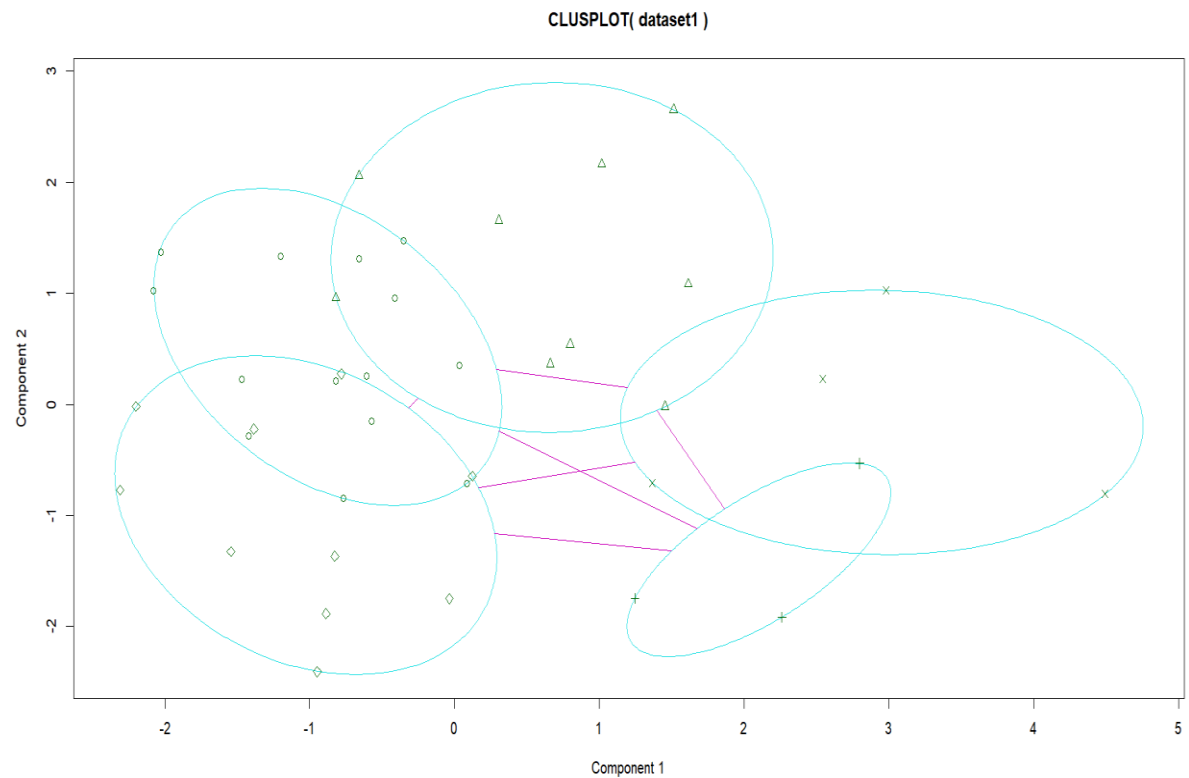
>
> ggscatter(
+   ind.coord, x = "Dim.1", y = "Dim.2",
+   color = "cluster", palette = "npg", ellipse = TRUE, ellipse.type = "convex",
+   shape = "circle", size = 1.5, legend = "right", ggtheme = theme_bw(),
+   xlab = paste0("Dim 1 (", variance.percent[1], "% )"),
+   ylab = paste0("Dim 2 (", variance.percent[2], "% )")
+ ) +
+   stat_mean(aes(color = cluster), size = 4)
> |

```

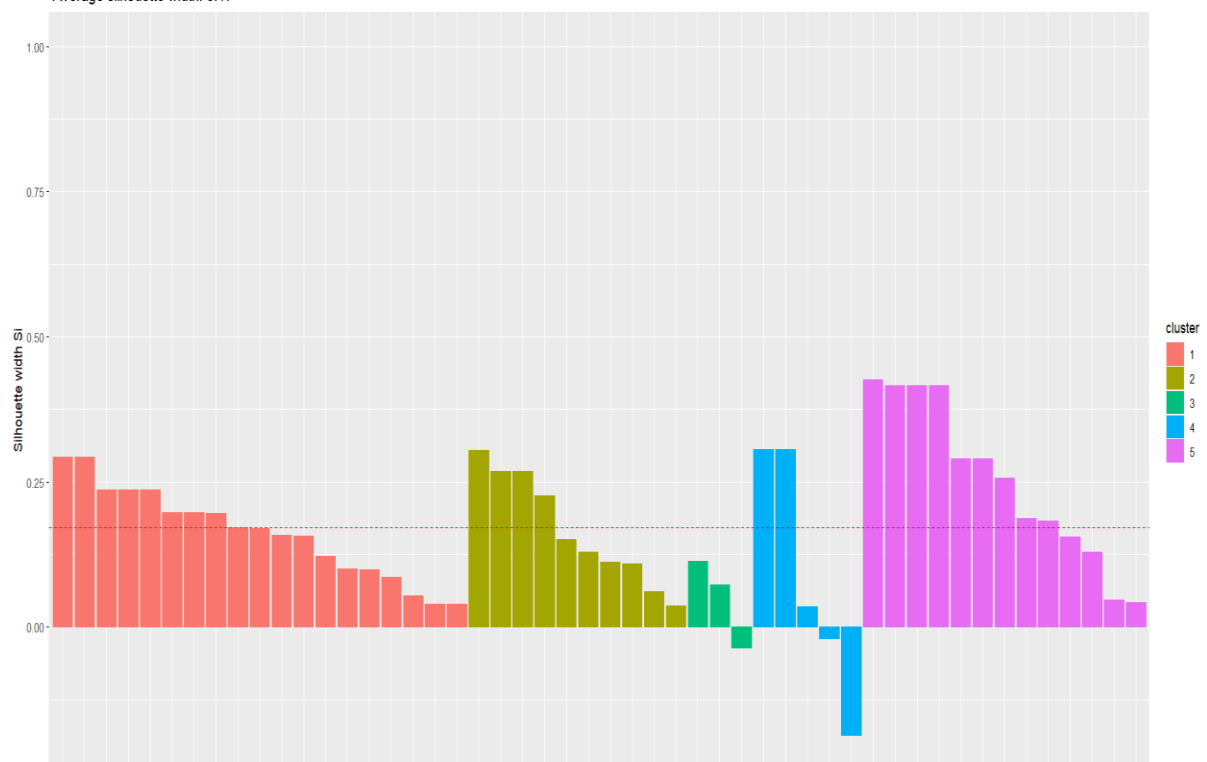


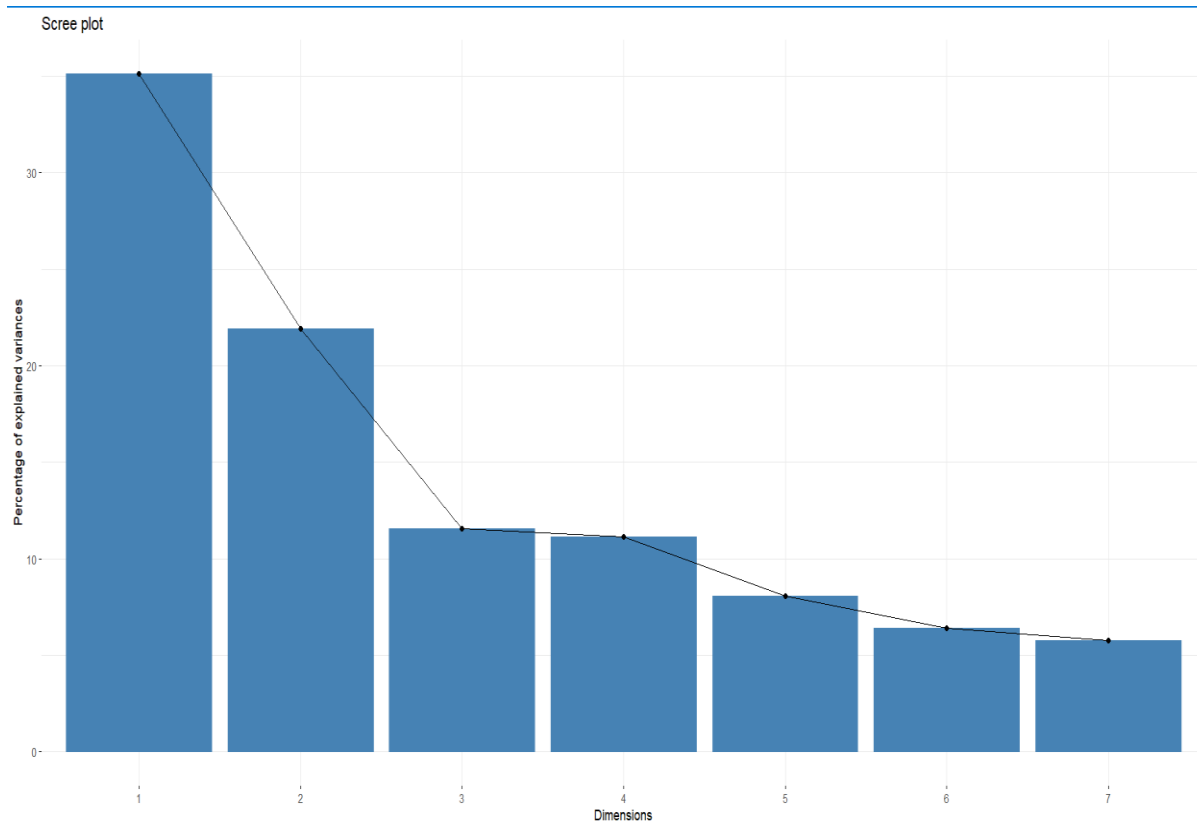
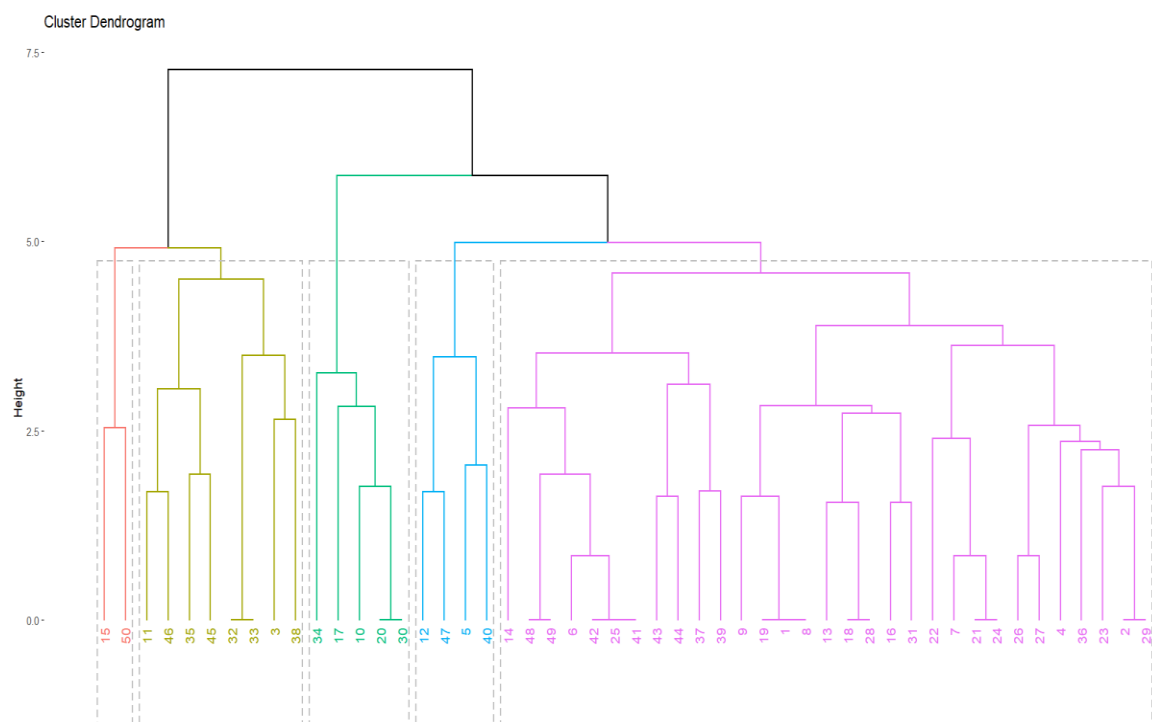


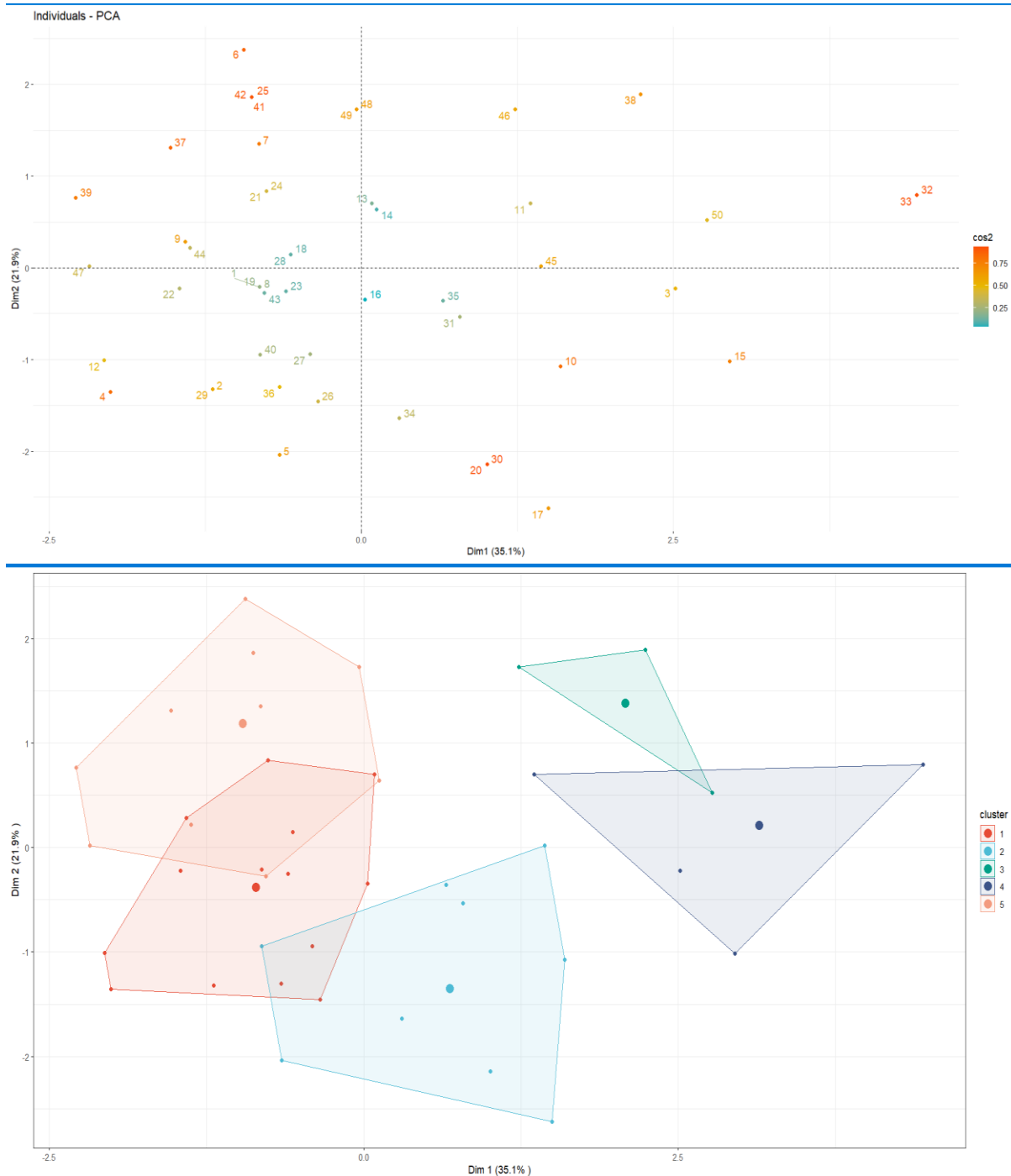




Clusters silhouette plot
Average silhouette width: 0.17







Interpretation:

- From the plot of the elbow method, we could observe that the line is bent at the range 5 making it the optimal number of clusters for the dataset.
- By using k means clustering the clusters have been formed.
- By using PCA we found that the variables whose value is greater than 0.5 has a great influence on once personality and social behaviour.
- The within-cluster sum of squares is a measure of the variability of the observations within each cluster. In general, a cluster that has a small sum of squares is more

compact than a cluster that has a large sum of squares. Clusters that have higher values exhibit greater variability of the observations within the cluster

- In Silhouette method
 - - Observations with a large silhouette S_i (almost 1) are very well clustered.
 - - A small S_i (around 0) means that the observation lies between two clusters.
 - - Observations with a negative S_i are probably placed in the wrong cluster
- For the age category 13-18 it observed that people under this category are **extroverts and intuitive**. They can cope with emotions and are well planned. They are assertive.
- The people under age category 19-24 **have similar personality and behavioural pattern** of people below 19.
- People of age between 25-30 are **extroverts and strongly intuitive** people. Their capacity to cope with emotions are high. They are very well planned and assertive people.
- 31-59 highly **extroverted and highly intuitive**, can easily manage their emotions and are planned.
- Within sum of squares **less**, the cluster is **more compact**.
- Within sum of squares of MIND is the greatest, hence making it the most dissimilar cluster.