

# Tech gadgets Brand Classification

## **ABSTRACT**

This project aims to analyze and discover the common standard features shared by various tech gadgets from various brands that help the brand stand out in the market.

## **PROBLEM STATEMENT**

Each of the technological gadgets is available in a variety of brands. In this modern era, it is difficult to determine which brand a customer should purchase based on their preferences, so clustering analysis and PCA analysis are used to classify the brand based on the characteristics and standard features shared by each brand.

## **INTRODUCTION**

Smartphone brand data is analyzed and clustered based on feature similarities, and the most important features that contribute the most to the brand are identified using dimensionality reduction using PCA.

## **METHODOLOGY USED**

### **Clustering Analysis using K- Means Algorithm**

### **Principle Component Analysis**

Clustering- data classification based on characteristic similarity

PCA –to identify the variables that have the greatest impact on all brands

## **SOFTWARE USED**

Data Collection - **Typeform**

Data Analysis - **R- Studio**

## **TECH GADGETS BRAND ANALYSIS**

### **DATASET :**

**Different brands of tech gadgets with various features are chosen for analysis and collected via typeform.**

**Sample – 129 (Respondents)**

**Gadgets names – Smartphone, Mouse Keyboard, Headphone Camera and Laptop**

**Gadget: Smartphone**

## Source Code

```
library("factoextra") - extracts and visualizes the result of Multivariate data
library("NbClust") - determine the best number of clusters
library("dplyr") - resolves the data manipulation hurdles
library("cluster") - perform the cluster analysis with the k-means algorithm
library(ggbiplot) - to visualize the PCA components in 2D
library("rstatix") - helper package of the univariate and multivariate data
library("FactoMineR") - to perform the principle component analysis
library(parameters) - Utilities for processing the parameters of various statistical models
```

## Data Preprocessing

```
smartphone <-
  read.table(
    file = "D:/Files/CIT/M.Sc.DCS/4th Semester/17MDC46 - PA Lab/PA Sem IV project/smartphone.csv",
    sep = ",",
    dec = ".",
    header = TRUE,
  )

sm = smartphone[2:11]
#preprocessing the data by replacing the NA values with zero
sm[is.na(sm)] = 0
# sm = scale(sm)
sm
```

```
> sm
  Brand Price Performance Quality Design Operate.Platform.system Value Get.used.to..Habit Reputation Services
1      7      8          10      0      0                      0      0              0          0          3
2      0      8           0      0      0                      0      0              0          0          0
3      7      8           0      9      0                      0      0              0          0          3
4      7      0          10      9      0                      0      5              0          0          0
5      0      0          10      0      0                      0      0              0          0          0
6      7      0           0      0      0                      0      0              0          0          0
7      0      8           0      0      0                      4      0              0          0          0
8      0      8           0      0      0                      0      0              0          0          0
9      0      0           0      9      0                      0      0              0          0          0
10     0      0          10      0      0                      0      0              0          0          0
11     0      0           0      9      0                      0      0              0          0          0
12     7      0          10      9      6                      4      0              0          0          0
13     7      8          10      9      0                      0      0              0          1          3
```

The above smartphone data is preprocessed by selecting numeric columns and are scaled to normalize the data to perform the clustering analysis.

## Optimal number of clusters and Quality of a *k*-means partition

```
#Elbow method

fviz_nbclust(sm, kmeans, method = "wss") +
  geom_vline(xintercept = 4, linetype = 2) +
  labs(subtitle = "Elbow method")

#silhouette method

fviz_nbclust(sm, kmeans, method = "silhouette")

#gap statistics method

gap_stat = clusGap(
  sm,
  FUN = kmeans,
  nstart = 25,
  K.max = 10,
  B = 50
```

```

)
fviz_gap_stat(gap_stat)

# Consensus Based Algorithm

n_clust <- n_clusters(sm,
  package = c("easystats", "NbClust", "mclust"),
  standardize = TRUE, fast = TRUE,
  nbclust_method = "kmeans")

n_clust

```

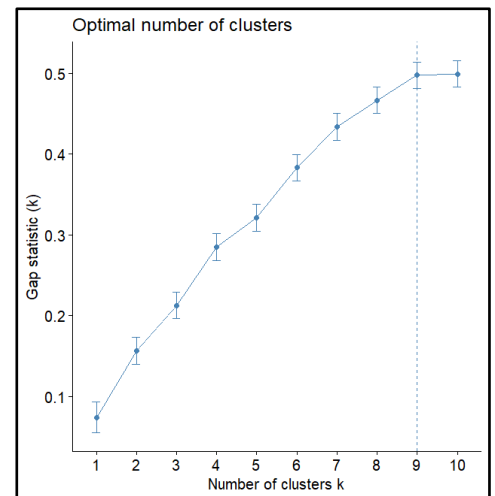
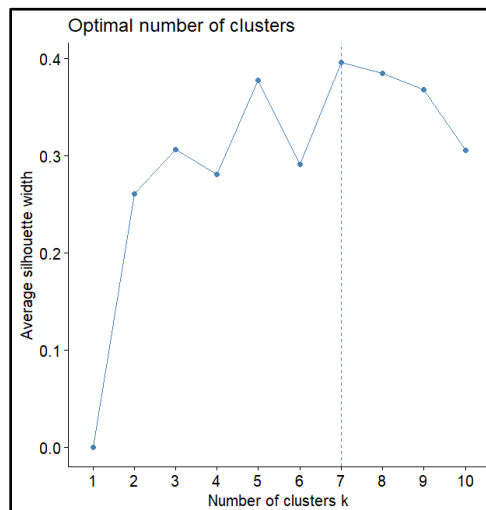
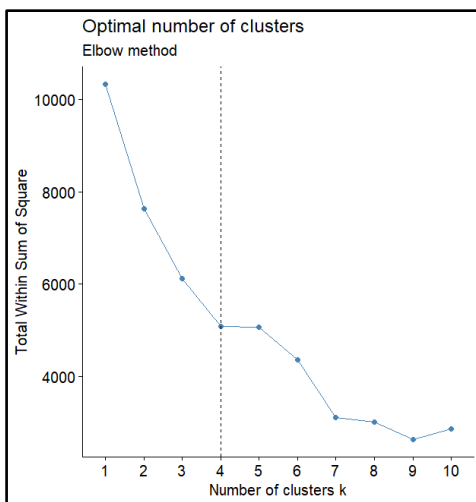
The **Elbow method** looks at the total within-cluster sum of square (WSS) as a function of the number of clusters. Here the optimal number is **4**. The Elbow method is sometimes ambiguous and an alternative is the average silhouette method.

The **Silhouette method** measures the Quality of a clustering and determines how well each point lies within its cluster. The Silhouette method suggests **7** clusters.

The optimal number of clusters is the one that maximizes the **gap statistic**. This method suggests only **9** clusters.

Here, the 3 approaches suggest a different number of clusters.

Because no method is clearly better, a fourth alternative is to run many methods and take the number of clusters that is the most agreed upon (i.e., find the consensus).



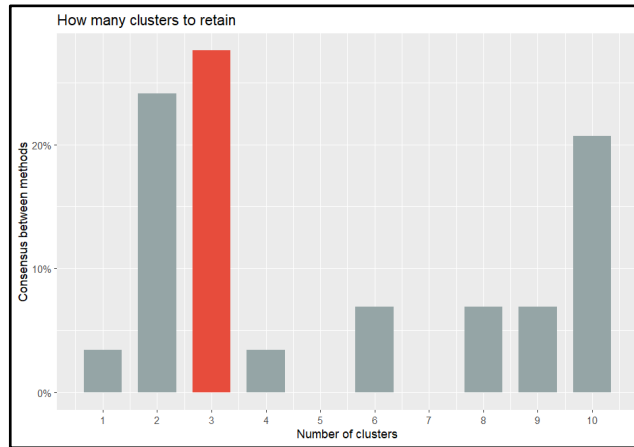
```

> # Consensus Based Algorithm
>
> n_clust <- n_clusters(sm,
+   package = c("easystats", "NbClust", "mclust"),
+   standardize = TRUE, fast = TRUE,
+   nbclust_method = "kmeans")
> n_clust
# Method Agreement Procedure:

The choice of 3 clusters is supported by 8 (27.59%) methods out of 29 (Hartigan, Marriot, trcovw, Tracew, Ratkowsky, Ball, PtBiserial, Mixture (EII)).
>

```

Based on all indices, most methods suggest to retain **3 clusters**, followed by a 4-clusters solution.



## K- Means Clustering

# Elbow Method - 4 , silhouette = 7 , gap statistic - 9 , Consense based algorithm -3

```
smart_result_e = kmeans(sm , 4 , nstart = 25)
smart_result_e
```

```
fviz_cluster(
  smart_result_e,
  data = sm,
  palette = c("#2E9FDF", "#00AFBB", "#E7B800", "violet"),
  geom = "point",
  ellipse.type = "convex",
  ggtheme = theme_bw()
)
```

```
# silhouette = 7
smart_result_s = kmeans(sm , 7)
smart_result_s
```

```
fviz_cluster(
  smart_result_s,
  data = sm,
  palette = c(
    "#2E9FDF",
    "#00AFBB",
    "#E7B800",
    "violet",
    "red",
    "pink",
    "green"
  ),
  geom = "point",
  ellipse.type = "convex",
  ggtheme = theme_bw()
)
```

```
# gap statistic - 9
smart_result_g = kmeans(sm , 9)
smart_result_g
```

```
fviz_cluster(
  smart_result_g,
  data = sm,
  palette = c(
    "#2E9FDF",
    "#00AFBB",
    "#E7B800",
    "violet",
    "red",

```

```

    "pink",
    "green",
    "brown",
    "yellow"
  ),
  geom = "point",
  ellipse.type = "convex",
  ggtheme = theme_bw()
)

# Final Optimal Cluster #Consense based algorithm is taken with the optimal clusters of 3

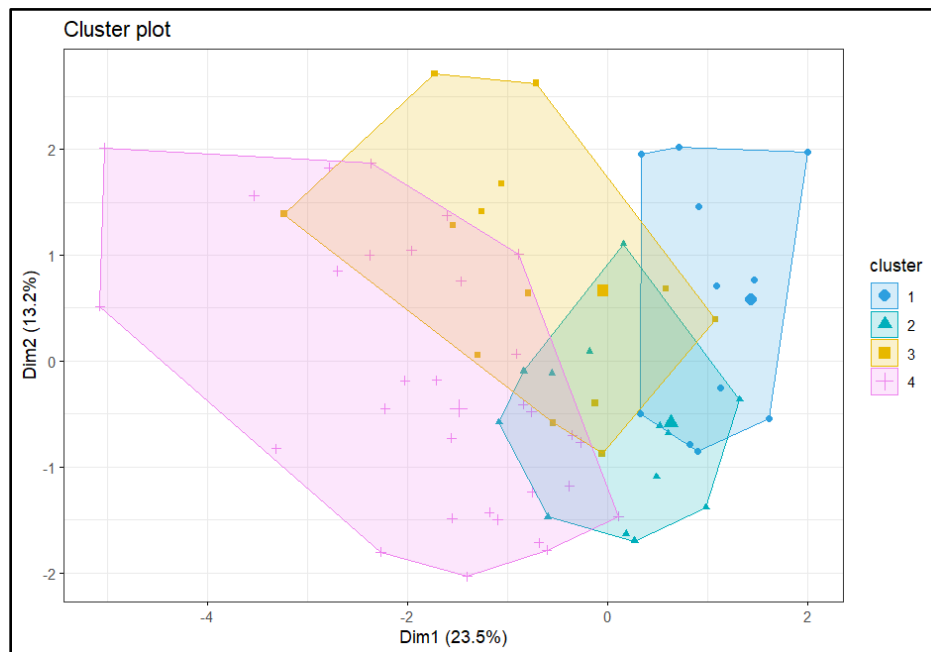
optimal_cluster = kmeans(sm , 3 , nstart = 25)
optimal_cluster

fviz_cluster(
  optimal_cluster,
  data = sm,
  palette = c("#2E9FDF",
              "#00AFBB",
              "#E7B800"),
  geom = "point",
  ellipse.type = "convex",
  ggtheme = theme_bw()
)

#quality of k-means partition
BSS = optimal_cluster$betweenss
TSS = optimal_cluster$totss
# We calculate the quality of the partition
BSS / TSS * 100

```

## Elbow Method



```
K-means clustering with 4 clusters of sizes 31, 25, 22, 40

Cluster means:
  Brand      Price Performance Quality      Design Operate.Platform.system
1 2.032258 3.354839         0         0 0.7741935              0.1290323
2 1.680000 3.200000        10         0 0.9600000              0.8000000
3 1.590909 1.818182         0         9 1.6363636              0.3636364
4 3.675000 5.000000        10         9 3.1500000              1.3000000

  Value Get.used.to..Habit Reputation  Services
1 0.483871      0.58064516 0.0000000 0.09677419
2 0.400000      0.16000000 0.0400000 0.12000000
3 1.136364      0.09090909 0.1363636 0.81818182
4 1.750000      0.25000000 0.2000000 0.75000000

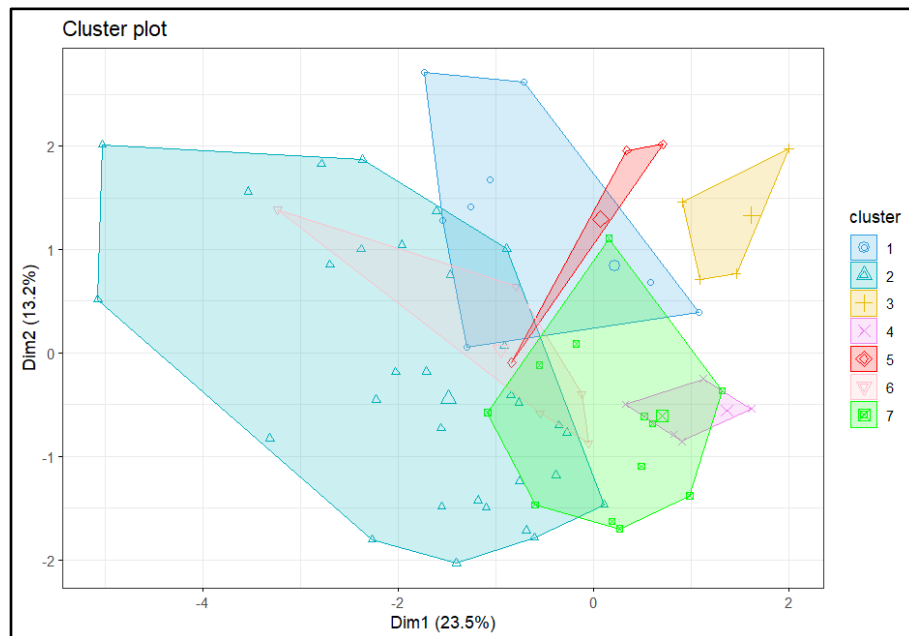
Clustering vector:
[1] 2 1 3 4 2 1 1 1 3 2 3 4 4 2 1 2 2 4 2 3 3 3 1 3 1 2 2 4 4 2 4 2 1 4
[35] 4 1 1 1 1 1 2 3 4 2 1 1 1 1 4 4 4 2 4 4 4 3 4 1 3 3 3 4 2 4 3 4 3 3
[69] 2 4 4 1 4 2 4 4 3 4 2 1 1 1 2 1 4 4 2 3 4 1 3 4 3 4 3 4 1 2 2 2 1 4
[103] 2 2 1 4 1 4 3 4 1 3 1 4 4 4 1 4
```

Within cluster sum of squares by cluster:

```
[1] 1038.9677 855.3600 765.0455 1907.1750
(between_SS / total_SS = 55.8 %)
```

The WSS value of cluster 4 in Elbow method is high 1907.1750 which means that there is dissimilarity in the clusters members

## Silhouette Method



```

K-means clustering with 7 clusters of sizes 17, 40, 16, 13, 3, 5, 24

Cluster means:
      Brand  Price Performance  Quality  Design Operate.Platform.system
1 1.235294 0.000000  0.000000    9 1.0588235 0.4705882
2 3.675000 5.000000 10.000000    9 3.1500000 1.3000000
3 2.625000 0.000000  0.000000    0 0.0000000 0.0000000
4 1.076923 8.000000  0.000000    0 0.9230769 0.3076923
5 4.666667 0.000000  3.333333    0 6.0000000 0.0000000
6 2.800000 8.000000  0.000000    9 3.6000000 0.0000000
7 1.458333 3.333333 10.000000    0 0.7500000 0.8333333

      Value Get.used.to..Habit Reputation  Services
1 0.8823529 0.1176471 0.11764706 0.7058824
2 1.7500000 0.2500000 0.20000000 0.7500000
3 0.6250000 0.8750000 0.00000000 0.1875000
4 0.0000000 0.0000000 0.00000000 0.0000000
5 3.3333333 1.3333333 0.00000000 0.0000000
6 2.0000000 0.0000000 0.20000000 1.2000000
7 0.2083333 0.1666667 0.04166667 0.1250000

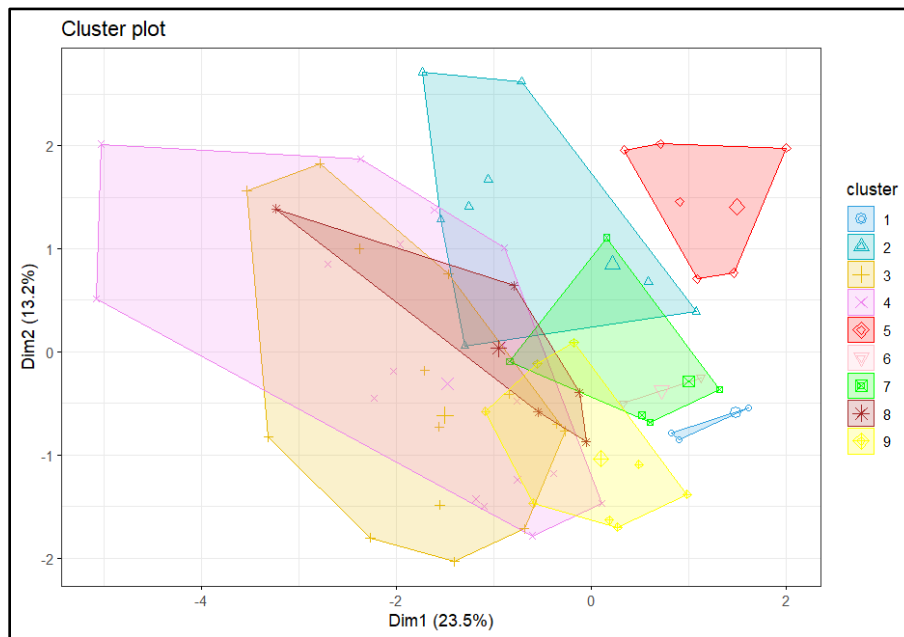
Clustering vector:
[1] 7 4 6 2 7 3 4 4 1 7 1 2 2 7 3 7 7 2 7 1 1 1 3 1 3 7 7 2 2 7 2 7 4 2
[35] 2 5 4 3 4 3 5 1 2 7 3 3 3 2 2 2 7 2 2 2 1 2 3 1 1 6 2 7 2 1 2 1 6
[69] 7 2 2 4 2 7 2 2 1 2 7 3 3 3 7 3 2 2 7 1 2 4 6 2 1 2 1 2 4 7 7 5 2
[103] 7 7 4 2 4 2 6 2 4 1 3 2 2 2 4 2

Within cluster sum of squares by cluster:
[1] 333.0588 1907.1750 251.6875 158.6154 118.6667 143.6000 766.0000
(between_SS / total_SS = 64.4 %)

```

The WSS value of cluster 2 in Silhouette method is high 1907.1705 which means that there is dissimilarity in the cluster's members

## Gap Statistics





```

K-means clustering with 9 clusters of sizes 11, 17, 18, 22, 18, 2, 15, 5, 10

Cluster means:
  Brand      Price Performance Quality    Design
1 0.0000000 8.000000      0      0 0.5454545
2 1.2352941 0.000000      0      9 1.0588235
3 2.3333333 3.555556     10      9 5.3333333
4 4.7727273 6.181818     10      9 1.3636364
5 2.7222222 0.000000      0      0 0.6666667
6 7.0000000 8.000000      0      0 3.0000000
7 0.9333333 0.000000     10      0 0.8000000
8 2.8000000 8.000000      0      9 3.6000000
9 2.8000000 8.000000     10      0 1.2000000

Operate.Platform.system      Value Get.used.to..Habit Reputation
1      0.3636364 0.0000000      0.0000000 0.0000000
2      0.4705882 0.8823529      0.1176471 0.1176471
3      1.5555556 1.9444444      0.2222222 0.1111111
4      1.0909091 1.5909091      0.2727273 0.2727273
5      0.0000000 0.8333333      1.0000000 0.0000000
6      0.0000000 0.0000000      0.0000000 0.0000000
7      0.5333333 0.3333333      0.1333333 0.0000000
8      0.0000000 2.0000000      0.0000000 0.2000000
9      1.2000000 0.5000000      0.2000000 0.1000000

Services
1 0.0000000
2 0.7058824
3 0.5000000
4 0.9545455
5 0.1666667
6 0.0000000
7 0.0000000
8 1.2000000
9 0.3000000

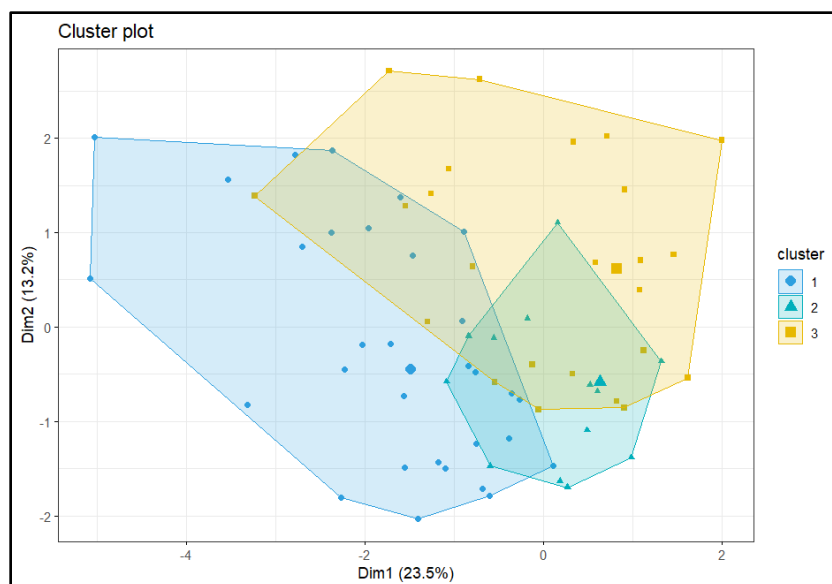
Clustering vector:
[1] 9 1 8 4 7 5 1 1 2 7 2 3 4 7 5 7 9 4 9 2 2 2 5 2 5 9 7 3 3 7 4 9 1 4
[35] 4 5 6 5 1 5 7 2 3 9 5 5 5 5 4 4 3 7 3 4 4 2 3 5 2 2 8 3 9 3 2 4 2 8
[69] 7 4 4 1 4 7 4 3 2 4 9 5 5 5 9 5 3 4 7 2 4 1 8 3 2 4 2 3 6 7 9 7 5 3
[103] 7 7 1 3 1 4 8 4 1 2 5 3 3 3 1 4

Within cluster sum of squares by cluster:
[1] 47.27273 333.05882 751.22222 867.04545 362.61111 18.00000 202.13333
[8] 143.60000 243.90000
(between_SS / total_SS = 71.3 %)

```

The WSS value of cluster 4 in Gap statistic method is high 867.04545 which means that there is dissimilarity in the clusters members

## Optimal Clusters



K-means clustering with 3 clusters of sizes 40, 25, 53

Cluster means:

	Brand	Price	Performance	Quality	Design	Operate	Platform	system
1	3.675000	5.000000	10	9.000000	3.150000			1.3000000
2	1.680000	3.200000	10	0.000000	0.960000			0.8000000
3	1.849057	2.716981	0	3.735849	1.132075			0.2264151

	Value	Get.used.to..Habit	Reputation	Services
1	1.750000	0.2500000	0.20000000	0.7500000
2	0.400000	0.1600000	0.04000000	0.1200000
3	0.754717	0.3773585	0.05660377	0.3962264

Clustering vector:

```
[1] 2 3 3 1 2 3 3 3 3 2 3 1 1 2 3 2 2 1 2 3 3 3 3 3 2 2 1 1 2 1 2 3 1
[35] 1 3 3 3 3 3 2 3 1 2 3 3 3 3 1 1 1 2 1 1 1 3 1 3 3 3 3 1 2 1 3 1 3 3
[69] 2 1 1 3 1 2 1 1 3 1 2 3 3 3 2 3 1 1 2 3 1 3 3 1 3 1 3 1 3 2 2 2 3 1
[103] 2 2 3 1 3 1 3 1 3 3 3 1 1 1 3 1
```

Within cluster sum of squares by cluster:

```
[1] 1907.175 855.360 2904.981
(between_SS / total_SS = 45.1 %)
```

The WSS value of cluster 3 in Consensus based Algorithm method is high 867.04545 which means that there is dissimilarity in the clusters members

## QUALITY TESTING

Centers= 3 (consensus based algorithm)

With scaling

```
> #quality of k-means partition
> BSS = smart_result_e$betweenss
> TSS = smart_result_e$totss
> # We calculate the quality of the partition
> BSS / TSS * 100
[1] 27.13615
>
```

Without scaling

```
>
> #quality of k-means partition
> BSS = smart_result_e$betweenss
> TSS = smart_result_e$totss
> # We calculate the quality of the partition
> BSS / TSS * 100
[1] 45.12983
>
```

Centers= 4

With scaling

```
> #quality of k-means partition
> BSS = smart_result_e$betweenss
> TSS = smart_result_e$totss
> # We calculate the quality of the partition
> BSS / TSS * 100
[1] 34.4876
>
```

Without scaling

```
>
> #quality of k-means partition
> BSS = smart_result_e$betweenss
> TSS = smart_result_e$totss
> # We calculate the quality of the partition
> BSS / TSS * 100
[1] 55.78888
>
```

The classification into **four or more** groups allows for a higher explained percentage and a higher quality.

This will always be the case: with more classes, the partition will be finer, and the *BSS* contribution will be higher. On the other hand, the “model” will be more complex, requiring more classes. In the extreme case where  $k = n$  (each observation is a singleton class), we have  $BSS = TSS$ , but the partition has lost all interest.

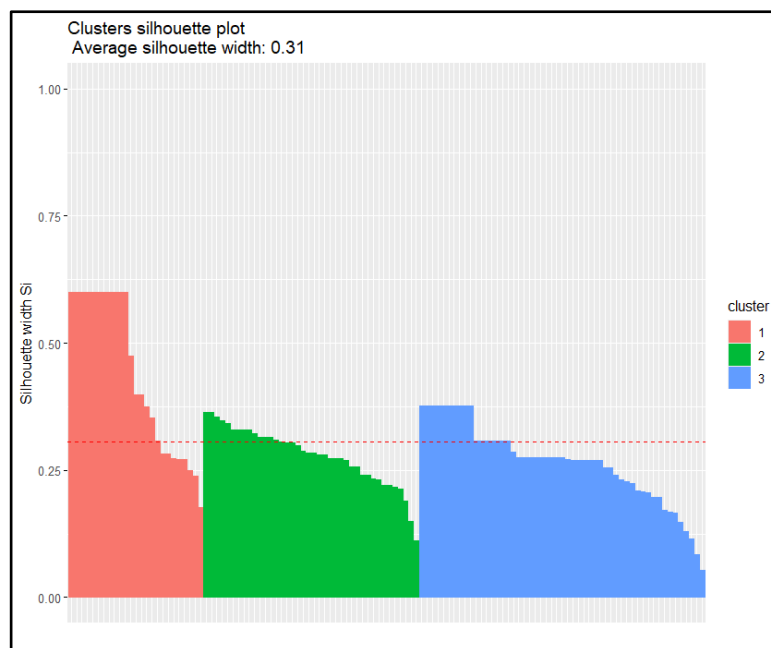
This is the reason we compare partitions via their Quality only for partitions that have the same number of clusters.

So we choose the **3 clusters (without scaling)** based on the optimality using the **consensus based algorithm**. The `nstart()` argument in the function also allows to run the algorithm several times with different initial centers, in order to obtain a potentially better partition

## Visualizations

To confirm that your number of classes is indeed optimal, there is a way to evaluate the Quality of your clustering via the silhouette plot (which shows the silhouette coefficient on the y axis).

We draw the silhouette plot for 3 clusters, as suggested by the average silhouette method:

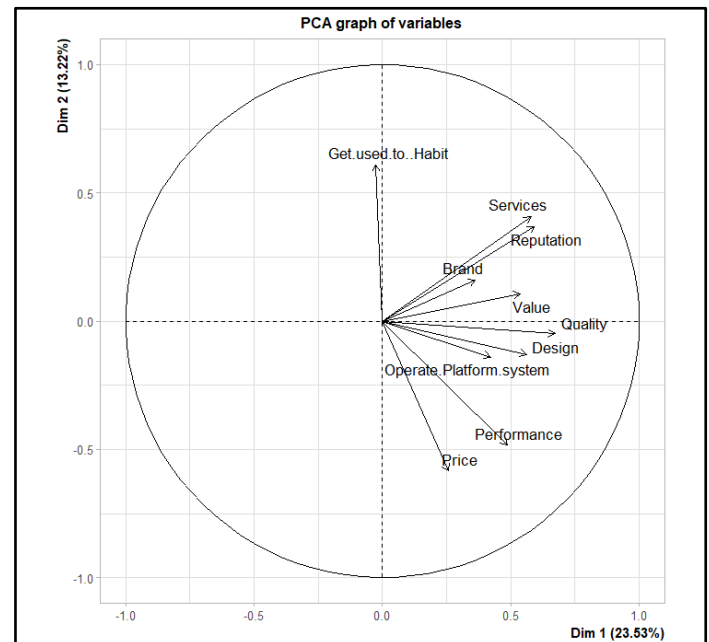
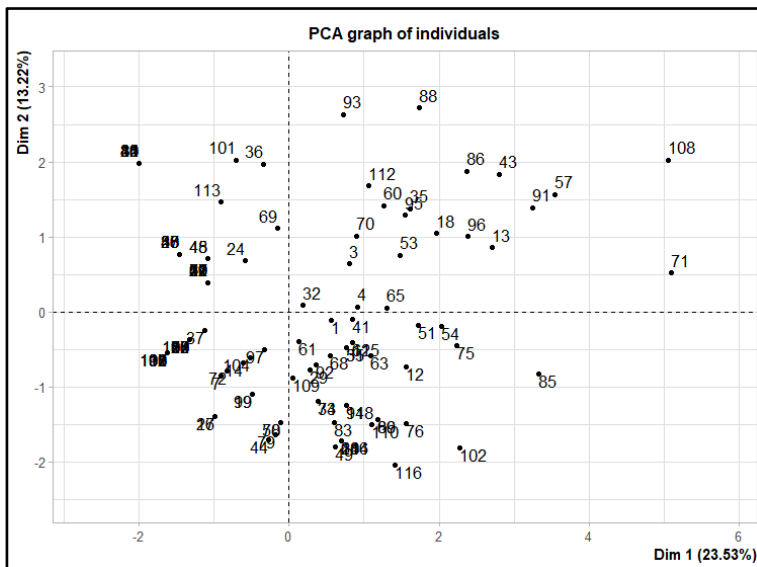


The silhouette plot above and the average silhouette coefficient say that the clustering is good and the clusters are optimal as the value is greater than zero means that the observation is well grouped. The closer the coefficient is to 1, the better the observation is grouped.

## Principle Component Analysis

```
res.pca = PCA(sm)  
res.pca
```

```
> res.pca = PCA(sm)  
> res.pca  
**Results for the Principal Component Analysis (PCA)**  
The analysis was performed on 118 individuals, described by 10 variables  
*The results are available in the following objects:  
  
  name      description  
1  "$eig"    "eigenvalues"  
2  "$var"    "results for the variables"  
3  "$var$coord" "coord. for the variables"  
4  "$var$cor" "correlations variables - dimensions"  
5  "$var$cos2" "cos2 for the variables"  
6  "$var$contrib" "contributions of the variables"  
7  "$ind"    "results for the individuals"  
8  "$ind$coord" "coord. for the individuals"  
9  "$ind$cos2" "cos2 for the individuals"  
10 "$ind$contrib" "contributions of the individuals"  
11 "$call"    "summary statistics"  
12 "$call$centre" "mean of the variables"  
13 "$call$ecart.type" "standard error of the variables"  
14 "$call$row.w" "weights for the individuals"  
15 "$call$col.w" "weights for the variables"  
>
```



The PCA() function performs PCA analysis for the smartphone data and plots the necessary variables in that contributes the most in the dataset

## Eigen Values /Variances

```
#Extract and visualize the eigen values  
get_eig(res.pca)  
#Visualize the eigen values/variances  
fviz_screplot(res.pca , addlabels = TRUE, ylim = c(0, 50))
```

```

> #Extract and visualize the eigen values
> get_eig(res.pca)

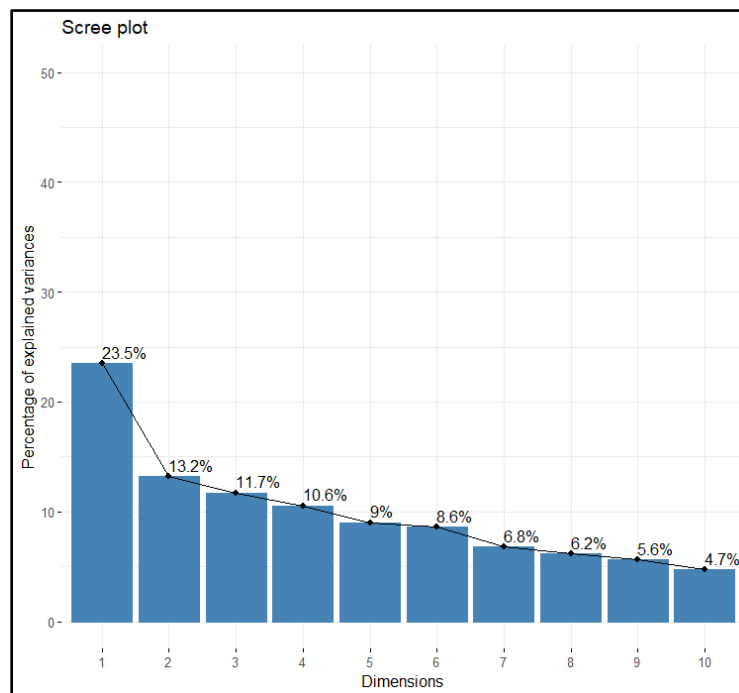
```

	eigenvalue	variance.percent	cumulative.variance.percent
Dim.1	2.3529526	23.529526	23.52953
Dim.2	1.3220895	13.220895	36.75042
Dim.3	1.1736112	11.736112	48.48653
Dim.4	1.0556305	10.556305	59.04284
Dim.5	0.9022899	9.022899	68.06574
Dim.6	0.8591302	8.591302	76.65704
Dim.7	0.6796218	6.796218	83.45326
Dim.8	0.6207263	6.207263	89.66052
Dim.9	0.5617762	5.617762	95.27828
Dim.10	0.4721718	4.721718	100.00000

```

>

```



The sum of all the eigenvalues gives a total variance of 10.

The proportion of variation explained by each eigenvalue is given in the second column .The PC1 and PC2 has 36% variability over the data

## Contribution of the Variables

```

# Extract the results from the variables
var = get_pca_var(res.pca)
var

#Coordinates of variables
head(var$coord)

#Contribution of variables
head(var$contrib)

```

```
sm.pca <- prcomp(sm, center = TRUE, scale. = TRUE)
summary(sm.pca)
```

```
> var
Principal Component Analysis Results for variables
=====
Name      Description
1 "$coord" "Coordinates for the variables"
2 "$cor"   "Correlations between variables and dimensions"
3 "$cos2"  "Cos2 for the variables"
4 "$contrib" "contributions of the variables"
>
> #Coordinates of variables
> head(var$coord)
      Dim.1      Dim.2      Dim.3      Dim.4      Dim.5
Brand      0.3618670  0.15861147  0.3643670  0.683767802 -0.25349311
Price      0.2572620 -0.58457364 -0.1490811  0.166685693  0.64438628
Performance 0.4866718 -0.48430004  0.3206401  0.065754001 -0.02844441
Quality     0.6726847 -0.04976176 -0.1972057 -0.002759058 -0.36173487
Design      0.5606938 -0.13072623  0.1230075 -0.063400643  0.05810829
Operate.Platform.system 0.4210595 -0.14003461  0.6041283 -0.335211960 -0.05576266
>
> #Contribution of variables
> head(var$contrib)
      Dim.1      Dim.2      Dim.3      Dim.4      Dim.5
Brand      5.565252  1.9028665 11.312374 4.428997e+01 7.12174168
Price      2.812794 25.8474436 1.893742 2.631993e+00 46.01998253
Performance 10.066051 17.7405933 8.760145 4.095741e-01 0.08967013
Quality     19.231354 0.1872969 3.313711 7.211237e-04 14.50222474
Design     13.360980 1.2926013 1.289256 3.807811e-01 0.37422267
Operate.Platform.system 7.534834 1.4832349 31.098119 1.064454e+01 0.34462025
>
> sm.pca <- prcomp(sm, center = TRUE, scale. = TRUE)
> summary(sm.pca)
Importance of components:
      PC1      PC2      PC3      PC4      PC5      PC6      PC7      PC8      PC9      PC10
Standard deviation 1.5339 1.1498 1.0833 1.0274 0.94989 0.92689 0.82439 0.78786 0.74952 0.68715
Proportion of Variance 0.2353 0.1322 0.1174 0.1056 0.09023 0.08591 0.06796 0.06207 0.05618 0.04722
Cumulative Proportion 0.2353 0.3675 0.4849 0.5904 0.68066 0.76657 0.83453 0.89661 0.95278 1.00000
```

The variables **Performance, Quality, and Design** contribute the most to the PC1 component, while **Price and Performance** contribute the most to the PC2 component. This means that the greater the contribution value, the more the variable contributes to the component.

## Validating the variables with clustering

```
# Contribution of the variables with the
res.km <- kmeans(var$coord, centers = 3, nstart = 25)
grp <- as.factor(res.km$cluster)
# Color variables by groups
fviz_pca_var(res.pca, col.var = grp,
             palette = c("#0073C2FF", "#EFC000FF", "#868686FF"),
             legend.title = "Cluster", addEllipses = TRUE, repel = TRUE)

ggbiplot(
  sm.pca,
  ellipse = TRUE,
  obs.scale = 1,
  var.scale = 1,
  var.axes = T,
  groups = smartphone$Smartphone
) + theme_minimal()
```



In the above graph, the variables **Get. Used to. Habit, Brand** contributes more to the cluster 1

**Services, Reputation, Value, and Quality** contribute more to the cluster 2

**Operating Platform/System, Design, Performance, and Price** contribute more to the cluster 3

Because the above-mentioned variables contributed more to a single cluster, this does not imply that it does not contribute to other groups, but their contribution is less than that of its cluster.

Clusters are formed based on the above criteria. When comparing the clusters with the brands, the above characteristic best fits the top three brands, namely **Samsung, Xiaomi, and Apple** and we can say that customers buys smartphones from these brands based on the characteristic depicted by each of the three clusters regarding the variables.

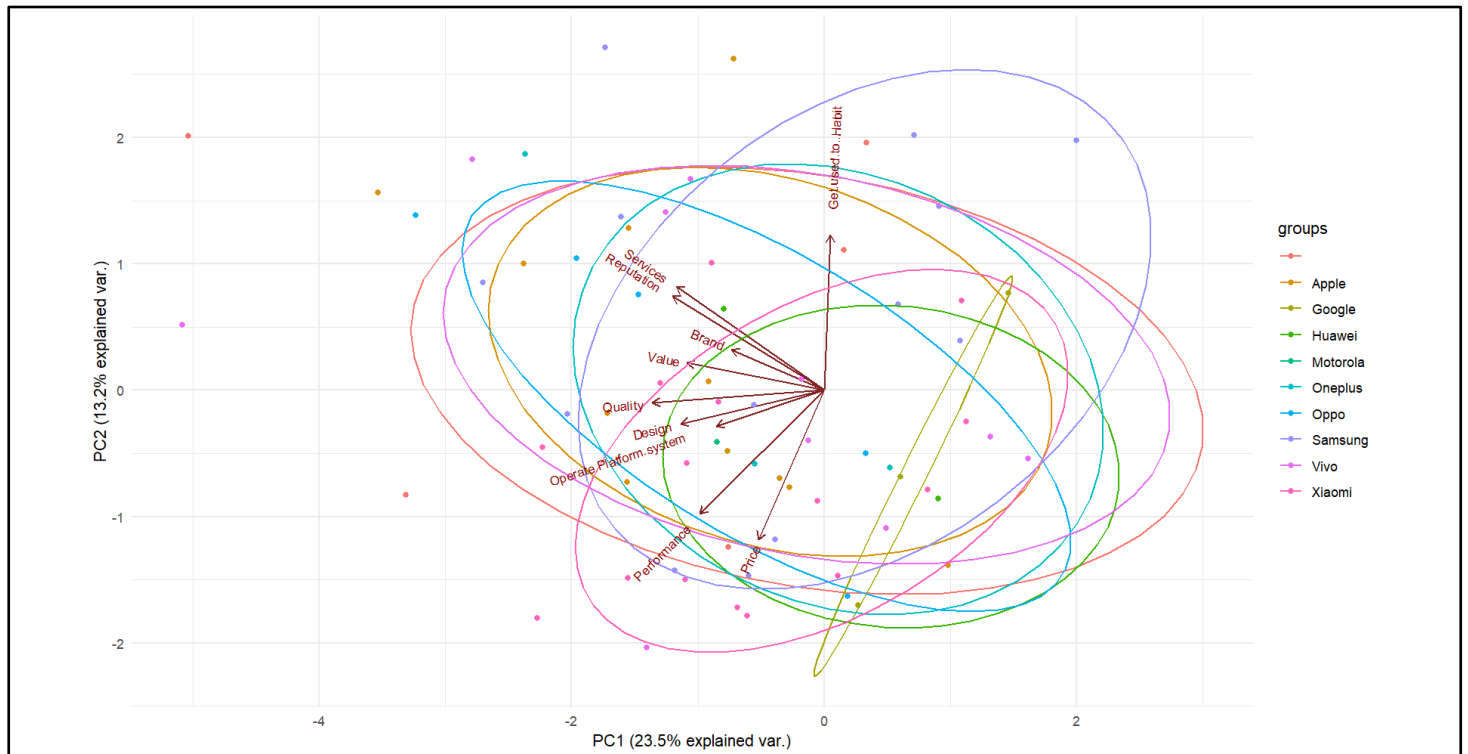
### For example :

The person who seeks for best design, price, and performance can buy Apple Smartphones

The person who is Brand-specific will buy Samsung smartphones

The person who strives for the best Quality goes for Xiaomi smartphones

The above graph represents the overall brands and the variables that contributes for each of the brand on a 2 D plane.

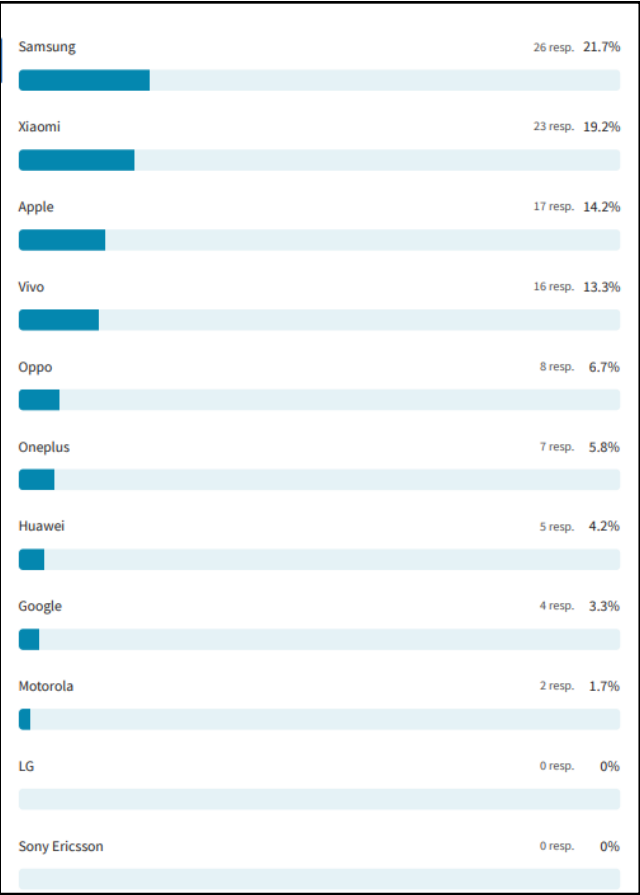


## CONCLUSION

The above clustering analysis creates clusters based on customers' preferences for brands, and through PCA analysis, the features that contribute the most to each brand of smartphone are identified and validated with the clusters, making it easier for consumers to select brands based on their preferences. For each brand, the output is verified and compared with survey respondents. The techniques described above can also be applied to other types of technology to determine brand categorization based on features.

- **Smartphone - Samsung**
- **Mouse - Logitech**
- **Headphone - Oneplus**
- **Camera – Canon**
- **Keyboard - Apple**
- **Laptop - HP**





## REFERENCE

**Clustering -** <https://statsandr.com/blog/clustering-analysis-k-means-and-hierarchical-clustering-by-hand-and-in-r/>

<https://www.r-bloggers.com/2021/04/cluster-analysis-in-r/>

**PCA -**

<http://www.sthda.com/english/articles/31-principal-component-methods-in-r-practical-guide/112-pca-principal-component-analysis-essentials/>