```
##-----##
```

library(data.table)

Creating new data frame for numerical values for doing PCA

tablechurnPCA = data.frame(tablechurn\$MonthlyCharges,tablechurn\$TotalCharges,tablechurn\$Tenure) setDT(tablechurnPCA)

##Setting column names for our new dataframe

names(tablechurnPCA) <- c('MonthlyCharges','TotalCharges','Tenure')

##head(tablechurnPCA)

tablechurnPCA

```
MonthlyCharges TotalCharges Tenure

    1:
    29.85
    29.85

    2:
    56.95
    1889.50

    3:
    53.85
    108.15

    4:
    42.30
    1840.75

    5:
    70.70
    151.65

                                                        34
                                                        2
                                                        45
                                      151.65
                                                        2
                    84.80
                                    1990.50
                                                        24
7028:
                               7362.90
346
                 103.20
                                                        72
7029:
7030:
                  29.60
74.40
                                                      11
7031:
                                       306.60
                                                        4
                   105.65
7032:
                                      6844.50
                                                        66
```

####Using prcomp function to compute the principal components (eigenvalues and eigenvectors).

#With scale=TRUE, variable means are set to zero, and variances set to one

ChurnPC <- prcomp(tablechurnPCA[,1:3],scale=TRUE)

ChurnPC

```
Standard deviations (1, .., p=3):
```

```
[1] 1.4764138 0.8722163 0.2438052
Rotation (n \times k) = (3 \times 3):
                        PC1
                                      PC2
                                                   PC3
MonthlyCharges 0.4857136 0.79237469 0.3690862
TotalCharges
                 0.6650968 -0.06101971 -0.7442600
Tenure
                 0.5672112 -0.60697524 0.5566440
summary(ChurnPC)
Importance of components:
                               PC1
                                       PC2
                                                PC3
                           1.4764 0.8722 0.24381
Standard deviation
Proportion of Variance 0.7266 0.2536 0.01981
Cumulative Proportion 0.7266 0.9802 1.00000
##Steps to check variances between the principal components
##Sample scores are stored in ChurnPC$x
# Square roots of eigenvalues are stored in ChurnPC$sdev
# Eigenvectors are stored in ChurnPC$rotation
# variable means stored in ChurnPC$center
# variable standard deviations stored in ChurnPC$scale
# Eigenvalues are sdev^2
##Calculating Eigen values
(eigen_churn <- ChurnPC$sdev^2)</pre>
[1] 2.1797978 0.7607612 0.0594410
names(eigen_churn) <- paste("PC",1:3,sep="")</pre>
names(eigen_churn)
[1] "PC1" "PC2" "PC3"
```

```
eigen_churn
```

```
PC1 PC2 PC3
2.1797978 0.7607612 0.0594410
```

##Taking sum of all eigen values

```
sum_churn <- sum(eigen_churn)
sum_churn</pre>
```

[1] 3

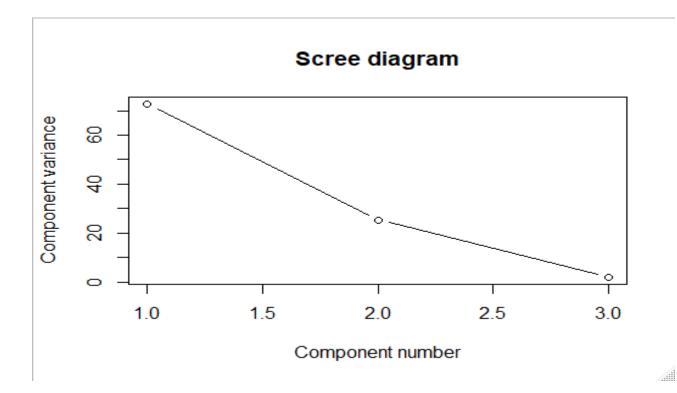
##Calculating percentage of variance

pvarchurn <- (eigen_churn/sum_churn)*100
pvarchurn</pre>

##The results show that PC1 and PC2 columns have good amount of data that is almost 98% variance #is presented with these 2 columns and since there is only 2% information in PC3 component we can afford to lose it

##Plotting Scree Diagram for Principal Components wrt percentage of their variances

plot(pvarchurn, xlab = "Component number", ylab = "Component variance", type = "b", main = "Scree diagram")



##Checking cumulative variances and other values

cumvar_churn <- cumsum(pvarchurn)</pre>

cumvar_churn

matchurn <- rbind(eigen_churn,pvarchurn,cumvar_churn)</pre>

rownames (matchurn) <- c ("Eigenvalues", "Prop. variance", "Cum. prop. variance")

round(matchurn,4)

```
PC1 PC2 PC3
Eigenvalues 2.1798 0.7608 0.0594
Prop. variance 72.6599 25.3587 1.9814
Cum. prop. variance 72.6599 98.0186 100.0000
```

summary(ChurnPC)

```
Importance of components:
                             PC1
                                    PC2
                                             PC3
                         1.4764 0.8722 0.24381
Standard deviation
Proportion of Variance 0.7266 0.2536 0.01981
Cumulative Proportion 0.7266 0.9802 1.00000
ChurnPC$rotation
                    PC1
                                 PC2
                                             PC3
MonthlyCharges 0.4857136 0.79237469 0.3690862
TotalCharges 0.6650968 -0.06101971 -0.7442600
Tenure
                0.5672112 -0.60697524 0.5566440
print(ChurnPC)
Standard deviations (1, .., p=3):
[1] 1.4764138 0.8722163 0.2438052
Rotation (n \times k) = (3 \times 3):
                       PC1
                                    PC2
                                                 PC3
MonthlyCharges 0.4857136 0.79237469 0.3690862
TotalCharges 0.6650968 -0.06101971 -0.7442600
Tenure
                0.5672112 -0.60697524 0.5566440
ChurnPC$x
sparrtyp_pca <- cbind(data.frame(tablechurn$Churn),ChurnPC$x)</pre>
sparrtyp_pca
# Means of scores for all the PC's classified by Churn value
tabmeansPC <- aggregate(sparrtyp_pca[,2:4],by=list(Churn=tablechurn$Churn),mean)
tabmeansPC
tabmeansPC <- tabmeansPC[rev(order(tabmeansPC$Churn)),]
tabmeansPC
tabfmeans <- t(tabmeansPC[,-1])
tabfmeans
colnames(tabfmeans) <- t(as.vector(tabmeansPC[1]))</pre>
```

```
tabfmeans
```

```
# Standard deviations of scores for all the PC's classified by Churn value
tabsdsPC <- aggregate(sparrtyp_pca[,2:4],by=list(Churn=tablechurn$Churn),sd)
tabfsds <- t(tabsdsPC[,-1])
colnames(tabfsds) <- t(as.vector(tabsdsPC[1]))</pre>
tabfsds
#T-Test
t.test(PC1~tablechurn$Churn,data=sparrtyp_pca)
t.test(PC2~tablechurn$Churn,data=sparrtyp_pca)
t.test(PC3~tablechurn$Churn,data=sparrtyp_pca)
# F ratio test
var.test(PC1~tablechurn$Churn,data=sparrtyp_pca)
var.test(PC2~tablechurn$Churn,data=sparrtyp_pca)
var.test(PC3~tablechurn$Churn,data=sparrtyp_pca)
# Levene's tests (one-sided)
library(car)
(LTPC1 <- leveneTest(PC1~tablechurn$Churn,data=sparrtyp_pca))
(p_PC1_1sided <- LTPC1[[3]][1]/2)
(LTPC2 <- leveneTest(PC2~tablechurn$Churn,data=sparrtyp_pca))
(p_PC2_1sided=LTPC2[[3]][1]/2)
(LTPC3 <- leveneTest(PC2~tablechurn$Churn,data=sparrtyp_pca))
```

```
(p_PC3_1sided <- LTPC3[[3]][1]/2)
# Plotting the scores for the first and second PC components
plot(sparrtyp_pca$PC1, sparrtyp_pca$PC2,pch=ifelse(sparrtyp_pca$Churn == "Yes",1,16),xlab="PC1",
ylab="PC2", main="7043 customer values for PC1 & PC2")
abline(h=0)
abline(v=0)
legend("bottomleft", legend=c("Yes","No"), pch=c(1,16))
plot(eigen_churn, xlab = "Component number", ylab = "Component variance", type = "I", main = "Scree
diagram")
plot(log(eigen_churn), xlab = "Component number", ylab = "log(Component variance)", type="l", main =
"Log(eigenvalue) diagram")
print(summary(ChurnPC))
View(ChurnPC)
diag(cov(ChurnPC$x))
xlim <- range(ChurnPC$x[,1])
ChurnPC$x[,1]
ChurnPC$x
plot(ChurnPC$x,xlim=xlim,ylim=xlim)
ChurnPC$rotation[,1]
ChurnPC$rotation
ChurnPC$x
plot(ChurnPC)
#get the original value of the data based on PCA
center <- ChurnPC$center
scale <- ChurnPC$scale
new_ChurnPC <- as.matrix(tablechurnPCA[,-3])</pre>
new ChurnPC
predict(ChurnPC)[,1]
```

```
##-----##
library(cluster)
tablechurnclust = data.frame(
tablechurn$Tenure,
tablechurn$TotalCharges,
tablechurn$MonthlyCharges)
##Making cluster on the basis of Total charges
rownames(tablechurnclust) <- tablechurn$TotalCharges
##Scaling done to make the data on scale
scaleTotalCharges <- scale(tablechurnclust[,1:ncol(tablechurnclust)])</pre>
scaleTotalCharges
#Here we have selected first row to see how our scaled matrix is like
head(scaleTotalCharges,1)
# We will find K-means by taking k=2, 3, 4, 5, 6...
\#For k-means = 2
(kmeans2.scaleTotalCharges <- kmeans(scaleTotalCharges,2,nstart = 10))
# Computing the percentage of variation accounted for two clusters
perc_var_kmeans2 <- round(100*(1 -
kmeans2.scaleTotalCharges$betweenss/kmeans2.scaleTotalCharges$totss),1)
names(perc_var_kmeans2) <- "Perc. 2 clus"</pre>
perc_var_kmeans2
```

```
\#For k-means = 3
(kmeans3.scaleTotalCharges <- kmeans(scaleTotalCharges,3,nstart = 10))
# Computing the percentage of variation accounted for. Two clusters
perc_var_kmeans3 <- round(100*(1 -
kmeans3.scaleTotalCharges$betweenss/kmeans3.scaleTotalCharges$totss),1)
names(perc_var_kmeans3) <- "Perc. 3 clus"</pre>
perc_var_kmeans3
#For k-means = 4
(kmeans4.scaleTotalCharges <- kmeans(scaleTotalCharges,4,nstart = 10))
# Computing the percentage of variation accounted for. Two clusters
perc_var_kmeans4 <- round(100*(1 -
kmeans 4. scale Total Charges \$ betweenss/kmeans 4. scale Total Charges \$ totss), 1)
names(perc_var_kmeans4) <- "Perc. 4 clus"</pre>
perc_var_kmeans4
#Using k means 4 could be good to preseent our data
# Saving above 4 k-means (1,2,3) in a list
##Now we will plot these clusters
library(fpc)
plotcluster(tablechurnclust,kmeans3.scaleTotalCharges$cluster)
##We didnt find any significant result using cluster analysis hence we will not be using it
#for our dataset
```