

R project

Source code:

```
> RawData <- read.table("Desktop/pima-indians
diabetes.data", sep = ",", header=FALSE)

> RowData

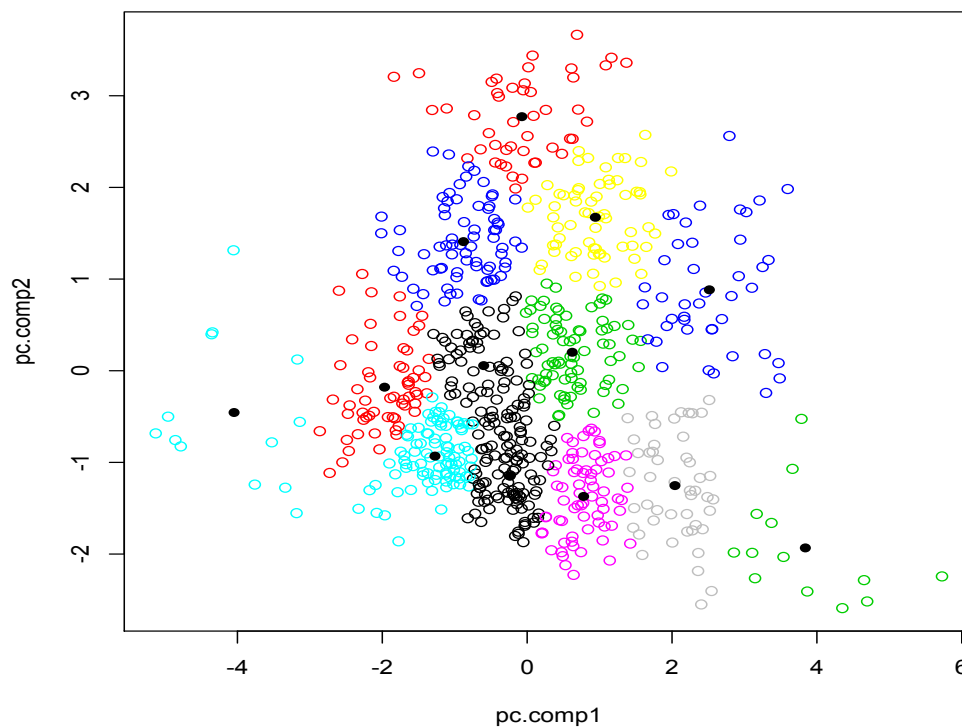
> responseY <- RawData[,dim(RawData)[2]]
> predictorX <- RawData[,1:(dim(RawData)[2]-1)]
> pca <- princomp(predictorX, cor=T) # principal
components analysis using correlation matrix
> pc.comp <- pca$scores
> pc.comp1 <- -1*pc.comp[,1] # principal component 1
scores (negated for convenience)
> pc.comp2 <- -1*pc.comp[,2] # principal component 2
scores (negated for convenience)
>
>
> X <- cbind(pc.comp1, pc.comp2)
> cl <- kmeans(X,13)
> cl$cluster
> plot(pc.comp1, pc.comp2,col=cl$cluster)
> points(cl$centers, pch=16)
```

Screenshot:

```

> plot(pc.comp1, pc.comp2,col=cl$cluster)
> points(cl$centers, pch=16)
> X <- cbind(pc.comp1, pc.comp2)
> cl <- kmeans(X,13)
> cl$cluster
 [1] 2 1 3 1 5 8 1 8 9 7 3 2 2 5 2 8 5 8 6 6 4 7 2 13 2 2 2 1 7 3 2 4 12 8 2
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Android voice recognition Screenshot:

