

Class 9 Mini Project

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#Unsupervised Learning Analysis of Human Breast Cancer Cells

Here we read data from the University of Wisconsin Medical Center on breast cancer patients

```
wisc.df <- read.csv("WisconsinCancer.csv", row.names=1)
head(wisc.df)
```

```
##      diagnosis radius_mean texture_mean perimeter_mean area_mean
## 842302         M      17.99       10.38        122.80     1001.0
## 842517         M      20.57       17.77        132.90     1326.0
## 84300903        M      19.69       21.25        130.00     1203.0
## 84348301         M      11.42       20.38         77.58      386.1
## 84358402         M      20.29       14.34        135.10     1297.0
## 843786         M      12.45       15.70         82.57      477.1
##      smoothness_mean compactness_mean concavity_mean concave.points_mean
## 842302      0.11840      0.27760      0.3001      0.14710
## 842517      0.08474      0.07864      0.0869      0.07017
## 84300903      0.10960      0.15990      0.1974      0.12790
## 84348301      0.14250      0.28390      0.2414      0.10520
## 84358402      0.10030      0.13280      0.1980      0.10430
## 843786      0.12780      0.17000      0.1578      0.08089
##      symmetry_mean fractal_dimension_mean radius_se texture_se perimeter_se
## 842302      0.2419      0.07871      1.0950      0.9053      8.589
## 842517      0.1812      0.05667      0.5435      0.7339      3.398
## 84300903      0.2069      0.05999      0.7456      0.7869      4.585
## 84348301      0.2597      0.09744      0.4956      1.1560      3.445
## 84358402      0.1809      0.05883      0.7572      0.7813      5.438
## 843786      0.2087      0.07613      0.3345      0.8902      2.217
##      area_se smoothness_se compactness_se concavity_se concave.points_se
## 842302    153.40      0.006399      0.04904      0.05373      0.01587
## 842517     74.08      0.005225      0.01308      0.01860      0.01340
## 84300903    94.03      0.006150      0.04006      0.03832      0.02058
## 84348301    27.23      0.009110      0.07458      0.05661      0.01867
## 84358402    94.44      0.011490      0.02461      0.05688      0.01885
## 843786     27.19      0.007510      0.03345      0.03672      0.01137
##      symmetry_se fractal_dimension_se radius_worst texture_worst
## 842302      0.03003      0.006193      25.38      17.33
## 842517      0.01389      0.003532      24.99      23.41
## 84300903      0.02250      0.004571      23.57      25.53
## 84348301      0.05963      0.009208      14.91      26.50
## 84358402      0.01756      0.005115      22.54      16.67
## 843786      0.02165      0.005082      15.47      23.75
```

```
##      perimeter_worst area_worst smoothness_worst compactness_worst
## 842302      184.60    2019.0          0.1622          0.6656
## 842517      158.80    1956.0          0.1238          0.1866
## 84300903     152.50    1709.0          0.1444          0.4245
## 84348301      98.87     567.7          0.2098          0.8663
## 84358402     152.20    1575.0          0.1374          0.2050
## 843786      103.40     741.6          0.1791          0.5249
##      concavity_worst concave.points_worst symmetry_worst
## 842302      0.7119          0.2654          0.4601
## 842517      0.2416          0.1860          0.2750
## 84300903     0.4504          0.2430          0.3613
## 84348301     0.6869          0.2575          0.6638
## 84358402     0.4000          0.1625          0.2364
## 843786      0.5355          0.1741          0.3985
##      fractal_dimension_worst
## 842302      0.11890
## 842517      0.08902
## 84300903     0.08758
## 84348301     0.17300
## 84358402     0.07678
## 843786      0.12440
```

Use -1 to remove the first column

```
wisc.data <- wisc.df[,-1]
head(wisc.data)
```

```
##      radius_mean texture_mean perimeter_mean area_mean smoothness_mean
## 842302      17.99      10.38      122.80    1001.0          0.11840
## 842517      20.57      17.77      132.90    1326.0          0.08474
## 84300903     19.69      21.25      130.00    1203.0          0.10960
## 84348301     11.42      20.38       77.58     386.1          0.14250
## 84358402     20.29      14.34      135.10    1297.0          0.10030
## 843786      12.45      15.70       82.57     477.1          0.12780
##      compactness_mean concavity_mean concave.points_mean symmetry_mean
## 842302      0.27760          0.3001          0.14710          0.2419
## 842517      0.07864          0.0869          0.07017          0.1812
## 84300903     0.15990          0.1974          0.12790          0.2069
## 84348301     0.28390          0.2414          0.10520          0.2597
## 84358402     0.13280          0.1980          0.10430          0.1809
## 843786      0.17000          0.1578          0.08089          0.2087
##      fractal_dimension_mean radius_se texture_se perimeter_se area_se
## 842302      0.07871      1.0950      0.9053      8.589    153.40
## 842517      0.05667      0.5435      0.7339      3.398     74.08
## 84300903     0.05999      0.7456      0.7869      4.585     94.03
## 84348301     0.09744      0.4956      1.1560      3.445     27.23
## 84358402     0.05883      0.7572      0.7813      5.438     94.44
## 843786      0.07613      0.3345      0.8902      2.217     27.19
##      smoothness_se compactness_se concavity_se concave.points_se
## 842302      0.006399          0.04904          0.05373          0.01587
## 842517      0.005225          0.01308          0.01860          0.01340
## 84300903     0.006150          0.04006          0.03832          0.02058
## 84348301     0.009110          0.07458          0.05661          0.01867
```

```
## 84358402      0.011490      0.02461      0.05688      0.01885
## 843786        0.007510      0.03345      0.03672      0.01137
##      symmetry_se fractal_dimension_se radius_worst texture_worst
## 842302      0.03003      0.006193      25.38      17.33
## 842517      0.01389      0.003532      24.99      23.41
## 84300903     0.02250      0.004571      23.57      25.53
## 84348301     0.05963      0.009208      14.91      26.50
## 84358402     0.01756      0.005115      22.54      16.67
## 843786      0.02165      0.005082      15.47      23.75
##      perimeter_worst area_worst smoothness_worst compactness_worst
## 842302      184.60      2019.0      0.1622      0.6656
## 842517      158.80      1956.0      0.1238      0.1866
## 84300903     152.50      1709.0      0.1444      0.4245
## 84348301      98.87      567.7      0.2098      0.8663
## 84358402     152.20      1575.0      0.1374      0.2050
## 843786      103.40      741.6      0.1791      0.5249
##      concavity_worst concave.points_worst symmetry_worst
## 842302      0.7119      0.2654      0.4601
## 842517      0.2416      0.1860      0.2750
## 84300903     0.4504      0.2430      0.3613
## 84348301     0.6869      0.2575      0.6638
## 84358402     0.4000      0.1625      0.2364
## 843786      0.5355      0.1741      0.3985
##      fractal_dimension_worst
## 842302      0.11890
## 842517      0.08902
## 84300903     0.08758
## 84348301     0.17300
## 84358402     0.07678
## 843786      0.12440
```

Diagnosis of vector

```
diagnosis <- as.factor(wisc.df$diagnosis)
head(diagnosis)
```

```
## [1] M M M M M M
## Levels: B M
```

Exploring data analysis

Q1. How many observations are in this dataset?

```
dim(wisc.data)
```

```
## [1] 569 30
```

How many rows

```
nrow(wisc.data)
```

```
## [1] 569
```

How many columns (i.e. variables)

```
ncol(wisc.data)
```

```
## [1] 30
```

Q2. How many of the observations have a malignant diagnosis?

```
table(wisc.df$diagnosis)
```

```
##  
##    B    M  
## 357 212
```

A useful function we will use lots and lots -> table()

Q3. How many variables/features in the data are suffixed with `_mean`?

```
length(grep("_mean", colnames(wisc.df)))
```

```
## [1] 10
```

Principle Component Analysis

Performing PCA

Here we need to scale the data before PCA as the various variable (i.e. columns) have very different scales.

Checking column means and standard deviations

```
colMeans(wisc.data)
```

```
##           radius_mean      texture_mean      perimeter_mean  
##      1.412729e+01      1.928965e+01      9.196903e+01  
##           area_mean      smoothness_mean      compactness_mean  
##      6.548891e+02      9.636028e-02      1.043410e-01  
##      concavity_mean      concave.points_mean      symmetry_mean  
##      8.879932e-02      4.891915e-02      1.811619e-01  
## fractal_dimension_mean      radius_se      texture_se  
##      6.279761e-02      4.051721e-01      1.216853e+00  
##      perimeter_se      area_se      smoothness_se  
##      2.866059e+00      4.033708e+01      7.040979e-03  
##      compactness_se      concavity_se      concave.points_se  
##      2.547814e-02      3.189372e-02      1.179614e-02  
##      symmetry_se      fractal_dimension_se      radius_worst  
##      2.054230e-02      3.794904e-03      1.626919e+01  
##      texture_worst      perimeter_worst      area_worst  
##      2.567722e+01      1.072612e+02      8.805831e+02  
##      smoothness_worst      compactness_worst      concavity_worst  
##      1.323686e-01      2.542650e-01      2.721885e-01  
##      concave.points_worst      symmetry_worst      fractal_dimension_worst  
##      1.146062e-01      2.900756e-01      8.394582e-02
```

```
apply(wisc.data,2,sd)
```

```
##           radius_mean      texture_mean      perimeter_mean
##      3.524049e+00      4.301036e+00      2.429898e+01
##           area_mean      smoothness_mean      compactness_mean
##      3.519141e+02      1.406413e-02      5.281276e-02
##      concavity_mean      concave.points_mean      symmetry_mean
##      7.971981e-02      3.880284e-02      2.741428e-02
## fractal_dimension_mean      radius_se      texture_se
##      7.060363e-03      2.773127e-01      5.516484e-01
##      perimeter_se      area_se      smoothness_se
##      2.021855e+00      4.549101e+01      3.002518e-03
##      compactness_se      concavity_se      concave.points_se
##      1.790818e-02      3.018606e-02      6.170285e-03
##      symmetry_se      fractal_dimension_se      radius_worst
##      8.266372e-03      2.646071e-03      4.833242e+00
##      texture_worst      perimeter_worst      area_worst
##      6.146258e+00      3.360254e+01      5.693570e+02
##      smoothness_worst      compactness_worst      concavity_worst
##      2.283243e-02      1.573365e-01      2.086243e-01
##      concave.points_worst      symmetry_worst      fractal_dimension_worst
##      6.573234e-02      6.186747e-02      1.806127e-02
```

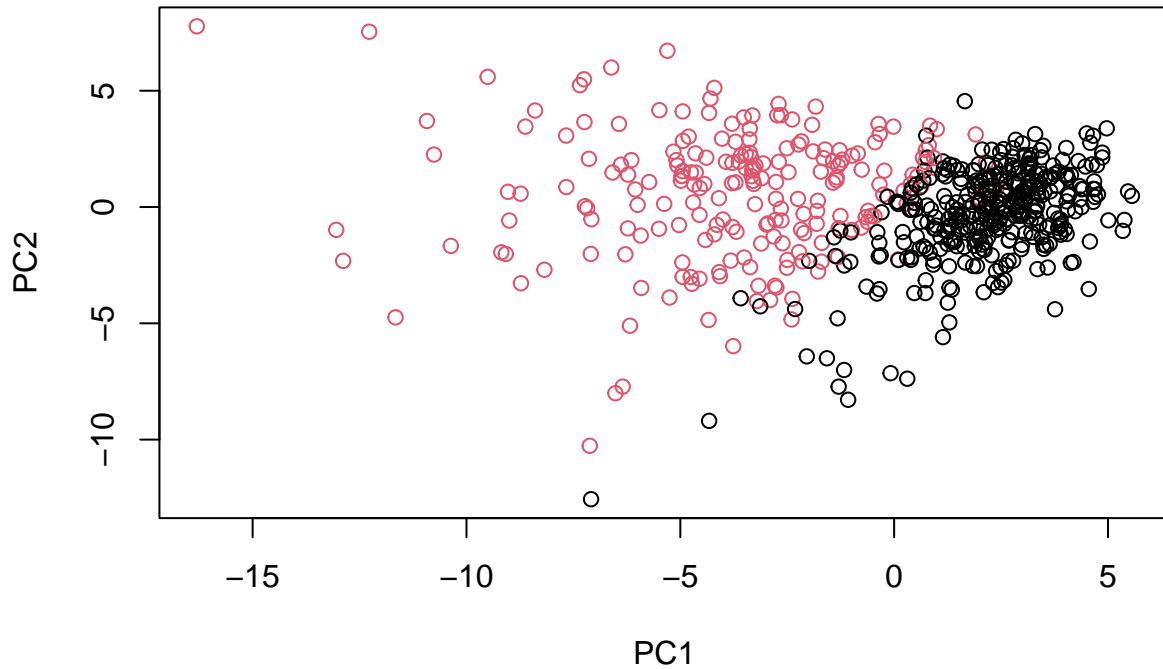
Performing PCA on wisc.data

```
wisc.pr <- prcomp(wisc.data, scale=TRUE)
summary(wisc.pr)
```

```
## Importance of components:
##           PC1      PC2      PC3      PC4      PC5      PC6      PC7
## Standard deviation  3.6444 2.3857 1.67867 1.40735 1.28403 1.09880 0.82172
## Proportion of Variance 0.4427 0.1897 0.09393 0.06602 0.05496 0.04025 0.02251
## Cumulative Proportion 0.4427 0.6324 0.72636 0.79239 0.84734 0.88759 0.91010
##           PC8      PC9      PC10      PC11      PC12      PC13      PC14
## Standard deviation  0.69037 0.6457 0.59219 0.5421 0.51104 0.49128 0.39624
## Proportion of Variance 0.01589 0.0139 0.01169 0.0098 0.00871 0.00805 0.00523
## Cumulative Proportion 0.92598 0.9399 0.95157 0.9614 0.97007 0.97812 0.98335
##           PC15      PC16      PC17      PC18      PC19      PC20      PC21
## Standard deviation  0.30681 0.28260 0.24372 0.22939 0.22244 0.17652 0.1731
## Proportion of Variance 0.00314 0.00266 0.00198 0.00175 0.00165 0.00104 0.0010
## Cumulative Proportion 0.98649 0.98915 0.99113 0.99288 0.99453 0.99557 0.9966
##           PC22      PC23      PC24      PC25      PC26      PC27      PC28
## Standard deviation  0.16565 0.15602 0.1344 0.12442 0.09043 0.08307 0.03987
## Proportion of Variance 0.00091 0.00081 0.0006 0.00052 0.00027 0.00023 0.00005
## Cumulative Proportion 0.99749 0.99830 0.9989 0.99942 0.99969 0.99992 0.99997
##           PC29      PC30
## Standard deviation  0.02736 0.01153
## Proportion of Variance 0.00002 0.00000
## Cumulative Proportion 1.00000 1.00000
```

Now we will make my main result: the “PCA plot” (a.k.a. “score plot”, PC1 vs. PC2 plot)

```
plot(wisc.pr$x[,1:2], col=diagnosis)
```



Q4. From your results, what proportion of the original variance is captured by the first principal components (PC1)?

44.27%

Q5. How many principal components (PCs) are required to describe at least 70% of the original variance in the data?

3 principal components (PC3)

Q6. How many principal components (PCs) are required to describe at least 90% of the original variance in the data?

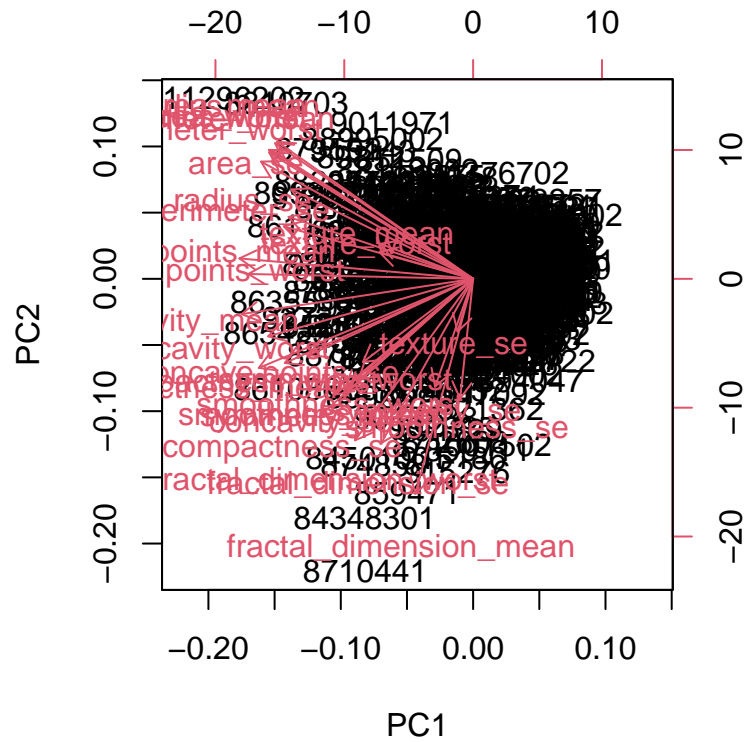
7 principal components (PC7)

Interpreting PCR results

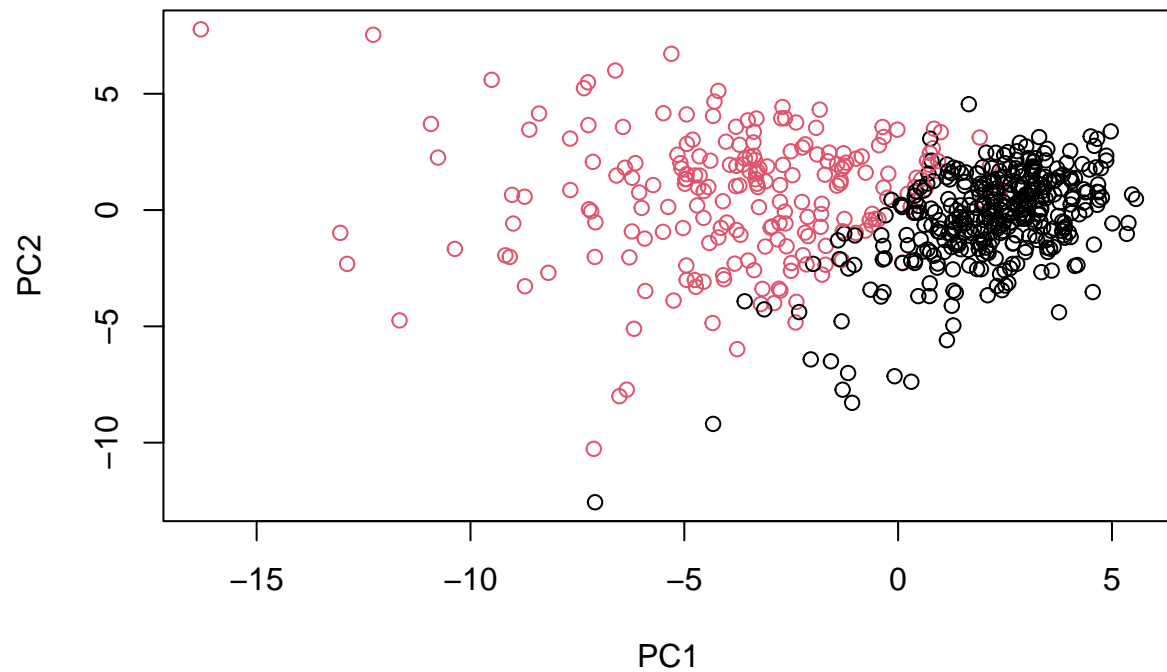
Q7. What stands out to you about this plot? Is it easy or difficult to understand? Why?

The plot is very messy and difficult to understand because all components are being observed on the plot.

```
biplot(wisc.pr)
```



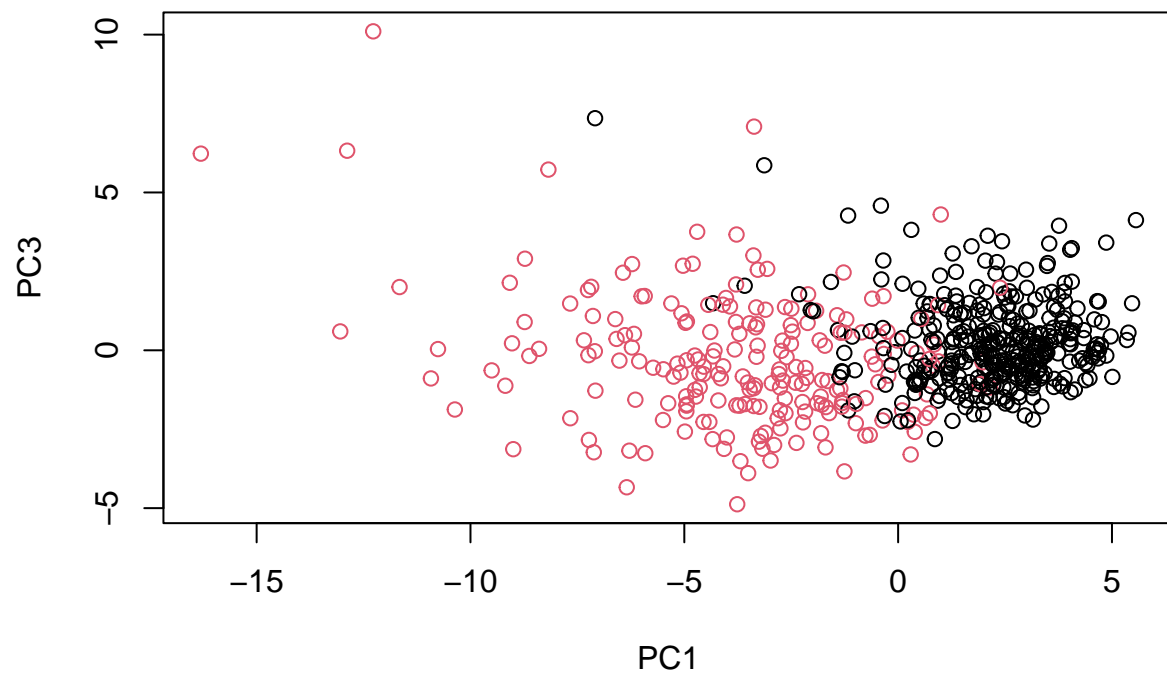
```
plot(wisc.pr$x , col = diagnosis ,
      xlab = "PC1", ylab = "PC2")
```



8. Generate a similar plot for principal components 1 and 3. What do you notice about these plots?

These plots contain less variance than principle component 2 and the principle component 1 is capturing a separation of malignant from benign.

```
plot(wisc.pr$x[, c(1,3) ], col = diagnosis,
     xlab = "PC1", ylab = "PC3")
```

Creating data.frame for ggplot

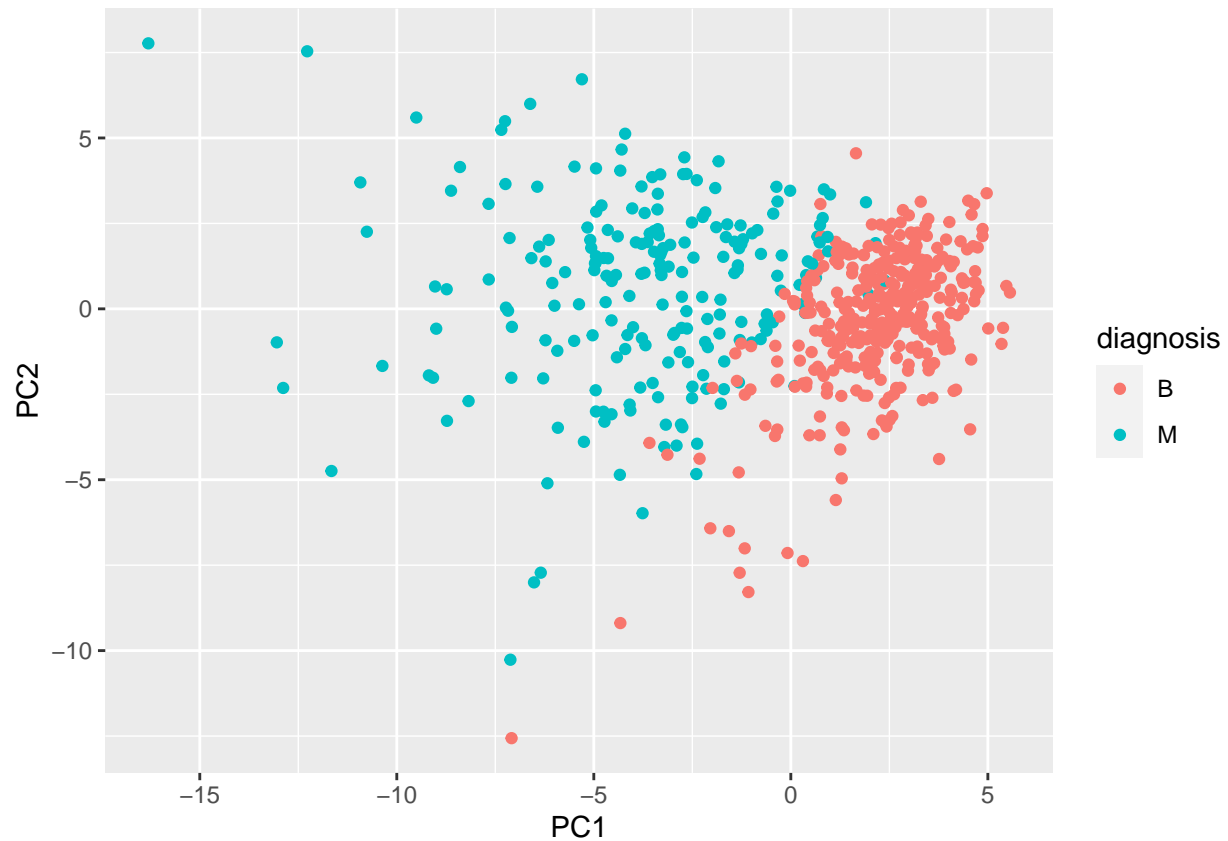
```
df <- as.data.frame(wisc.pr$x)
df$diagnosis <- diagnosis
```

Loading ggplot2 package

```
library(ggplot2)
```

Making a scatter plot colored by diagnosis

```
ggplot(df) +
  aes(PC1, PC2, col= diagnosis) +
  geom_point()
```



Variance explained

Calculating variance of each component

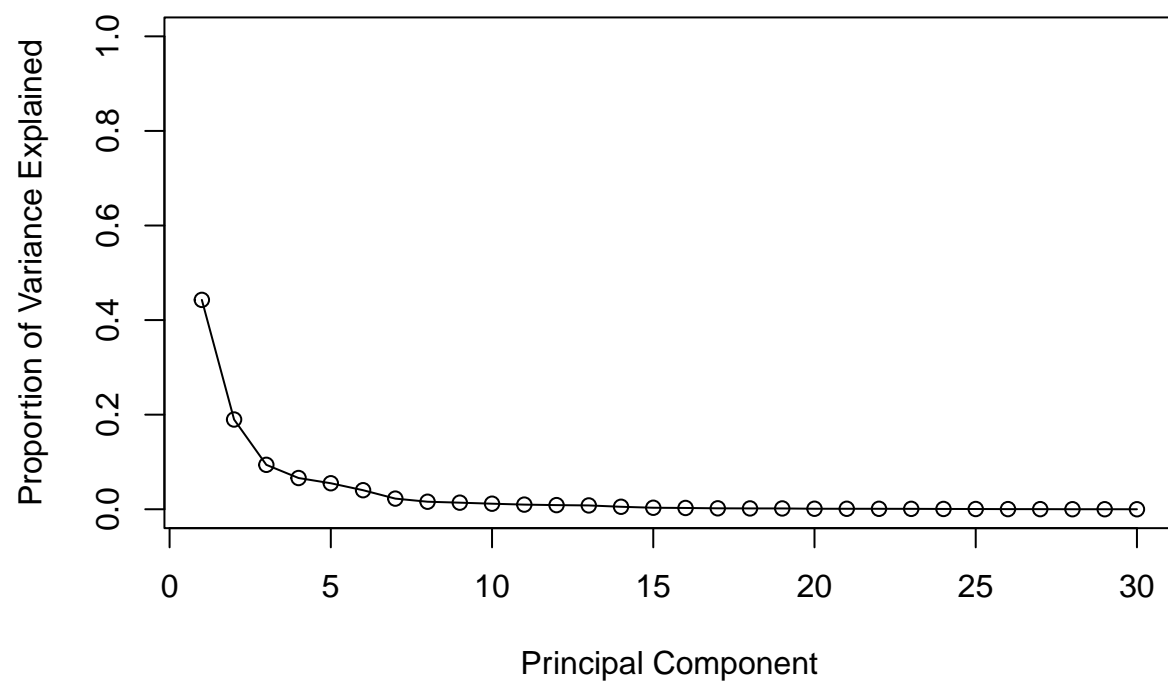
```
pr.var <- wisc.pr$sdev^2
head(pr.var)
```

```
## [1] 13.281608  5.691355  2.817949  1.980640  1.648731  1.207357
```

Calculating variance explained by each principle component : pve

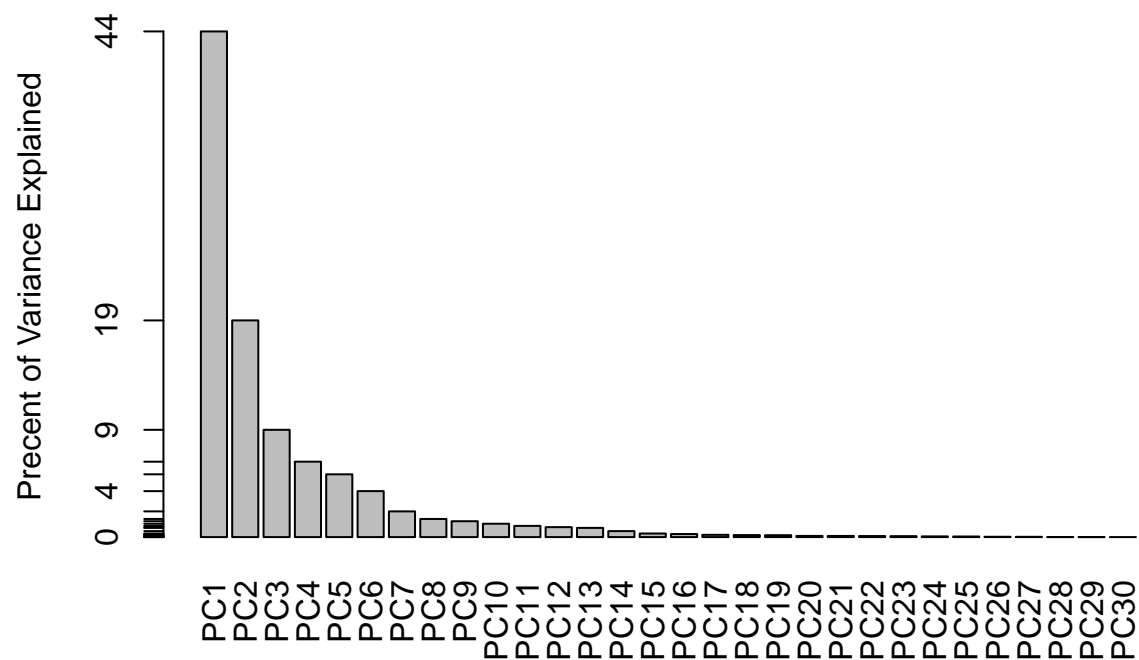
```
pve <- pr.var/ sum(pr.var)

plot(pve, xlab = "Principal Component",
     ylab = "Proportion of Variance Explained",
     ylim = c(0, 1), type = "o")
```



Alternative scree plot of the same data, note data driven y-axis

```
barplot(pve, ylab = "Precent of Variance Explained",
        names.arg=paste0("PC",1:length(pve)), las=2, axes = FALSE)
axis(2, at=pve, labels=round(pve,2)*100 )
```

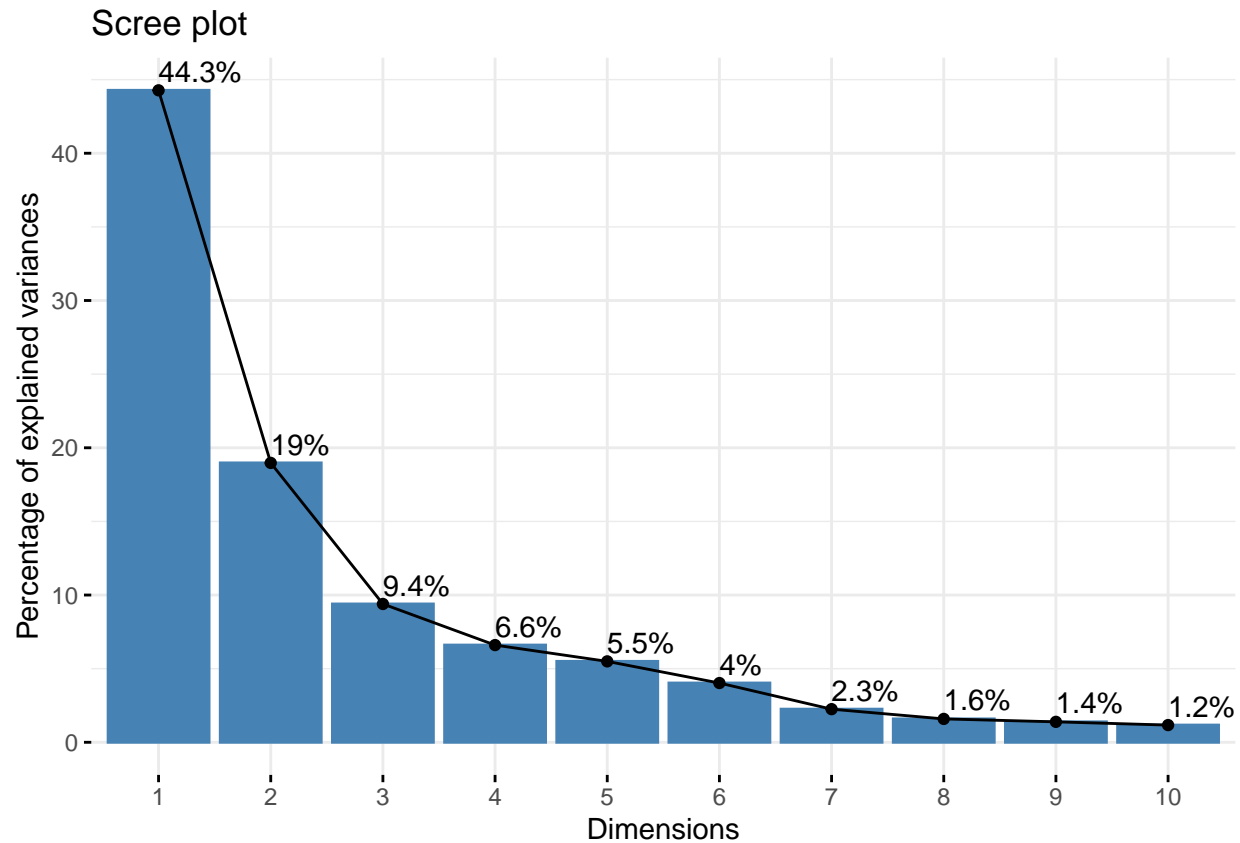


ggplot based on graph

```
#install.packages("factoextra")
library(factoextra)
```

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

```
fviz_eig(wisc.pr, addlabels = TRUE)
```



Communicating PCA results

Q9. For the first principal component, what is the component of the loading vector (i.e. `wisc.pr$rotation[,1]`) for the feature `concave.points_mean`?

```
wisc.pr$rotation[,1]
```

```
##      radius_mean      texture_mean      perimeter_mean
##      -0.21890244      -0.10372458      -0.22753729
##      area_mean      smoothness_mean      compactness_mean
##      -0.22099499      -0.14258969      -0.23928535
##      concavity_mean      concave.points_mean      symmetry_mean
##      -0.25840048      -0.26085376      -0.13816696
##      fractal_dimension_mean      radius_se      texture_se
##      -0.06436335      -0.20597878      -0.01742803
##      perimeter_se      area_se      smoothness_se
##      -0.21132592      -0.20286964      -0.01453145
##      compactness_se      concavity_se      concave.points_se
##      -0.17039345      -0.15358979      -0.18341740
##      symmetry_se      fractal_dimension_se      radius_worst
##      -0.04249842      -0.10256832      -0.22799663
##      texture_worst      perimeter_worst      area_worst
##      -0.10446933      -0.23663968      -0.22487053
##      smoothness_worst      compactness_worst      concavity_worst
```

```
##          -0.12795256          -0.21009588          -0.22876753
## concave.points_worst symmetry_worst fractal_dimension_worst
##          -0.25088597          -0.12290456          -0.13178394
```

The loading vector for the concave.points_mean is -0.26085376.

Q10. What is the minimum number of principal components required to explain 80% of the variance of the data?

```
summary(wisc.pr)
```

```
## Importance of components:
##          PC1      PC2      PC3      PC4      PC5      PC6      PC7
## Standard deviation  3.6444 2.3857 1.67867 1.40735 1.28403 1.09880 0.82172
## Proportion of Variance 0.4427 0.1897 0.09393 0.06602 0.05496 0.04025 0.02251
## Cumulative Proportion 0.4427 0.6324 0.72636 0.79239 0.84734 0.88759 0.91010
##          PC8      PC9      PC10     PC11     PC12     PC13     PC14
## Standard deviation  0.69037 0.6457 0.59219 0.5421 0.51104 0.49128 0.39624
## Proportion of Variance 0.01589 0.0139 0.01169 0.0098 0.00871 0.00805 0.00523
## Cumulative Proportion 0.92598 0.9399 0.95157 0.9614 0.97007 0.97812 0.98335
##          PC15     PC16     PC17     PC18     PC19     PC20     PC21
## Standard deviation  0.30681 0.28260 0.24372 0.22939 0.22244 0.17652 0.1731
## Proportion of Variance 0.00314 0.00266 0.00198 0.00175 0.00165 0.00104 0.0010
## Cumulative Proportion 0.98649 0.98915 0.99113 0.99288 0.99453 0.99557 0.9966
##          PC22     PC23     PC24     PC25     PC26     PC27     PC28
## Standard deviation  0.16565 0.15602 0.1344 0.12442 0.09043 0.08307 0.03987
## Proportion of Variance 0.00091 0.00081 0.0006 0.00052 0.00027 0.00023 0.00005
## Cumulative Proportion 0.99749 0.99830 0.9989 0.99942 0.99969 0.99992 0.99997
##          PC29     PC30
## Standard deviation  0.02736 0.01153
## Proportion of Variance 0.00002 0.00000
## Cumulative Proportion 1.00000 1.00000
```

5 principle component (PC5)

Hierarchical CLustering

Scaling the wisc.data using “scale()” function

```
data.scaled <- scale(wisc.data)
```

Calculating the distances between all pairs of observations and assigning to data.dist

```
data.dist <- dist(data.scaled)
```

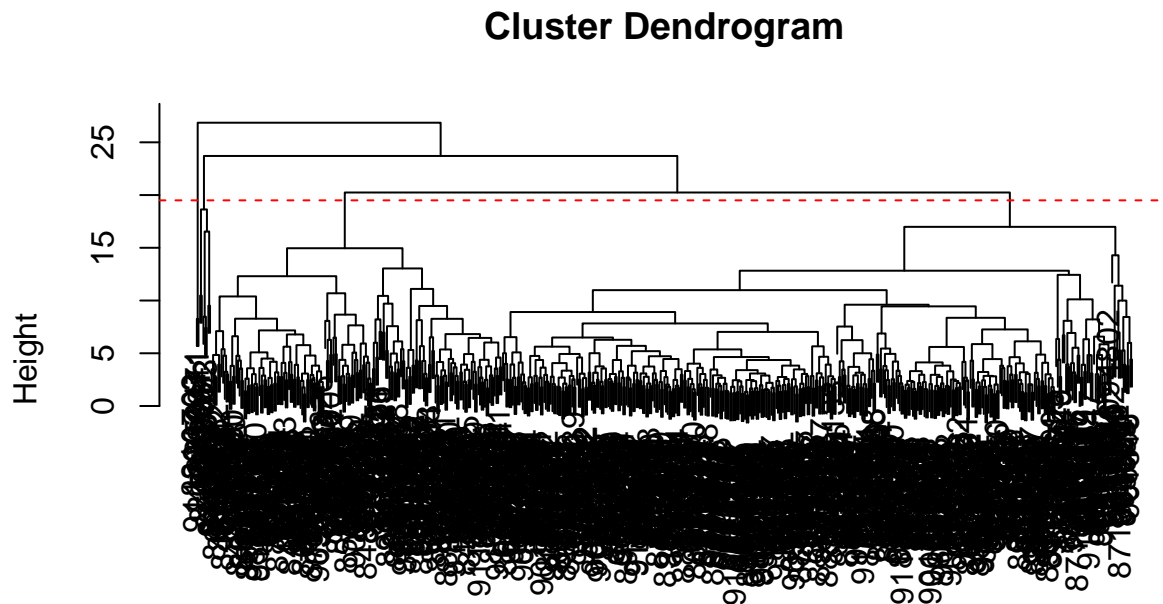
Creating hierarchical clustering model

```
wisc.hclust <- hclust(data.dist, method= "complete" )
```

Results of hierarchical clustering

Q11. Using the `plot()` and `abline()` functions, what is the height at which the clustering model has 4 clusters?

```
plot(wisc.hclust)
abline(h= 19.5, col="red", lty=2)
```



```
data.dist
hclust (*, "complete")
```

Height= 19.5 at which the clustering model has 4 clusters. ### Selecting number of clusters

```
wisc.hclust.clusters <- cutree(wisc.hclust, h=19.5)
```

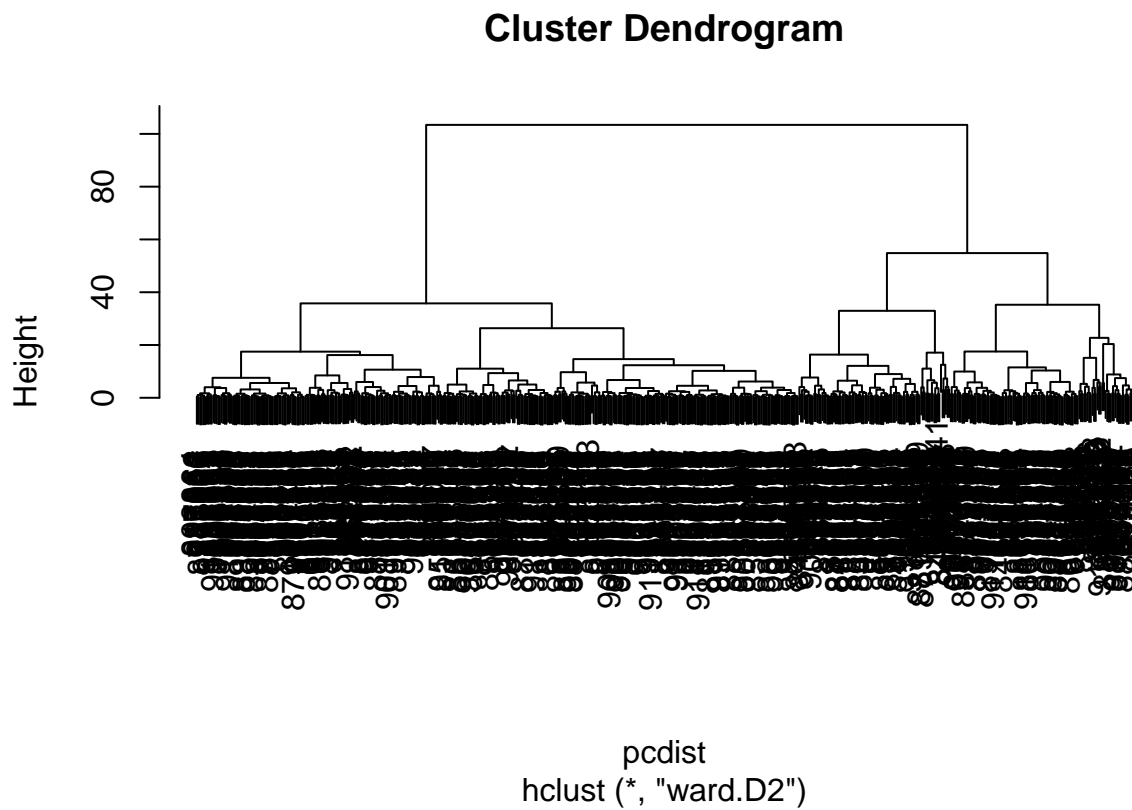
Using “`table()`” function to compare the cluster membership to actual diagnosis.

```
table(wisc.hclust.clusters, diagnosis)
```

```
##              diagnosis
## wisc.hclust.clusters  B  M
##                   1 12 165
##                   2   2   5
##                   3 343  40
##                   4   0   2
```

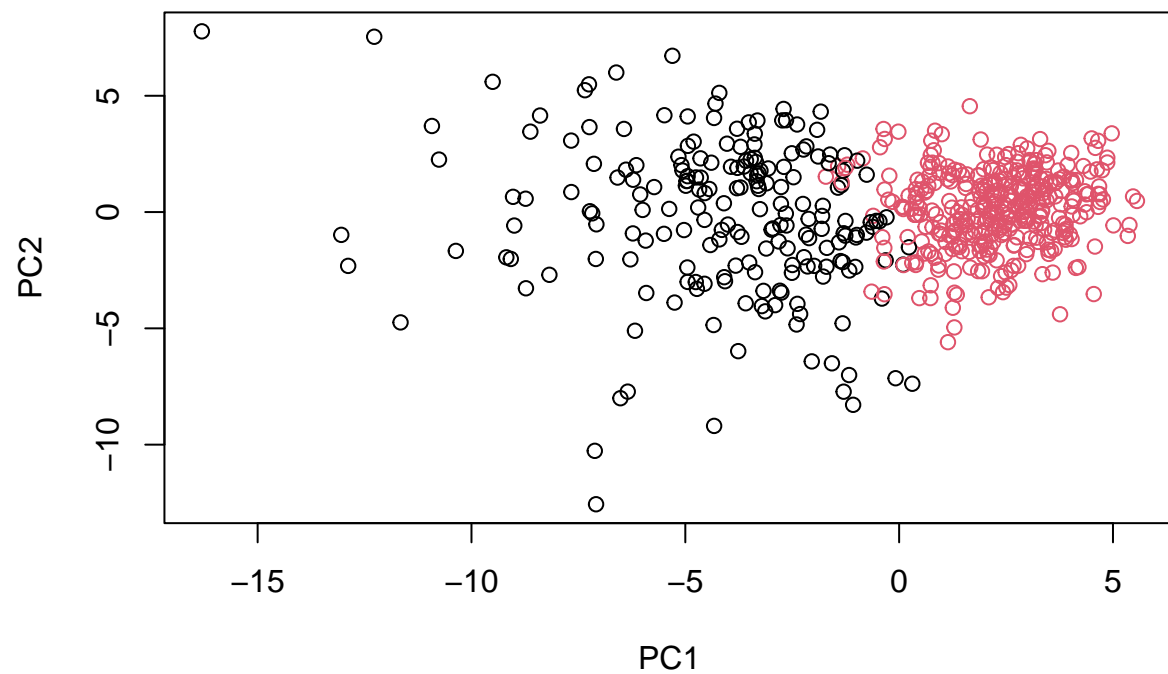
First 3 PCs for clustering

```
pcdist <- dist(wisc.pr$x[,1:3])
wisc.pr.hclust <- hclust(pcdist, method="ward.D2")
plot(wisc.pr.hclust)
```



```
grps <- cutree(wisc.pr.hclust, k= 2 )
```

```
plot(wisc.pr$x[,1:2], col= grps)
```

How well do my clusters agree with the expert M/B values

```
table(diagnosis, grps)
```

```
##          grps
## diagnosis  1  2
##      B   24 333
##      M  179  33
```