

APEKSHA

Fri Apr 05 15:43:43 2019

```
#ggplot2 is used to plot the bar plot
#install.packages("ggplot2")
library("ggplot2")
#corrplot is used to plot the correlation matrix
#install.packages("corrplot")
library("corrplot")
```

```
## corrplot 0.84 loaded
```

```
#It is used to reshape a one-dimensional array into a two-dimensional array with one column and multiple arr
ays.
#install.packages("reshape")
library("reshape")
```

```
## Warning: package 'reshape' was built under R version 3.5.2
```

```
#Reading the dataset
breast_cancer <- read.csv("C:\\Users\\APEKSHA\\Downloads\\wisc_bc_data.csv")
#Displaying the dataset using head function
head(breast_cancer)</pre>
```

```
id diagnosis radius_mean texture_mean perimeter_mean area_mean
## 1 87139402 B 12.32 12.39 78.85 464.1
## 2 8910251 B 10.60 18.95 69.28 346.4
                                 16.83
## 3 905520
                В
                      11.04
                                              70.92
                      11.28
                                 13.39
## 4 868871
                В
                                             73.00
                      15.19
                                 13.21
## 5 9012568
                В
                                             97.65
                                                      711.8
## 6 906539 B 11.57 19.04
                                          74.20
## smoothness_mean compactness_mean concavity_mean points_mean
## 1 0.10280 0.06981 0.03987 0.03700
                      0.11470
        0.09688
                                  0.06387
                                            0.02642
## 2
         0.10770
                                  0.03046
                                            0.02480
## 3
                      0.07804
                                            0.04796
                      0.11360
                                   0.04635
## 4
         0.11640
                  0.11360
0.06934
0.07722
                                  0.03393
0.05485
## 5
          0.07963
                                             0.02657
         0.08546
                                             0.01428
## symmetry mean dimension mean radius se texture se perimeter se area se
     0.1959 0.05955 0.2360 0.6656 1.670 17.43
## 1
                                     1.1970
                   0.06491 0.4505
                                                 3.430 27.10
## 2
        0.1922
        0.1714
## 3
                   0.06340 0.1967 1.3870
                                                1.342 13.54
## 4
        0.1771
                   0.06072 0.3384 1.3430
                                                1.851 26.33
## 5 0.1721 0.05544 0.1783 0.4125 1.338 17.72
## 6 0.2031 0.06267 0.2864 1.4400 2.206 20.30
## smoothness_se compactness_se concavity_se points_se symmetry_se

    ## 1
    0.008045
    0.011800
    0.01683
    0.012410
    0.01924

## 2
      0.007470
                                                0.03504
                   0.035810
                              0.03354 0.013650
                  0.009355
      0.005158
## 3
                                                0.01718
                              0.01056 0.007483
       0.011270
## 4
                   0.034980
                              0.02187 0.019650
                                                0.01580
                               0.01551 0.009155
       0.005012
                   0.014850
## 5
                                                 0.01647
## 6 0.007278 0.020470 0.04447 0.008799
                                              0.01868
## dimension_se radius_worst texture_worst perimeter_worst area_worst
## 1 0.002248 13.50 15.64
                                            86.97 549.1
                              22.94
                                                    424.8
## 2
    0.003318
                   11.88
                                           78.28
## 3 0.002198
                   12.41
                              26.44
                                           79.93
                                                    471.4
## 4 0.003442
                   11.92
                              15.77
                                           76.53
## 5 0.001767
                              15.73
                                          104.50
                                                    819.1
                  16.20
## 6 0.003339 13.07 26.98 86.43
## smoothness_worst compactness_worst concavity_worst points_worst
## 1 0.1385 0.1266 0.12420 0.09391
                         0.2515
                                     0.19160
                                                0.07926
## 2
           0.1213
                         0.1482
                                     0.10670
## 3
           0.1369
                                                0.07431
                         0.1822
## 4
           0.1367
                                      0.08669
                                                 0.08611
                                      0.25600
## 5
           0.1126
                          0.1737
                                                 0.08178
                        0.1937
           0.1249
## 6
                                                0.06664
## symmetry_worst dimension_worst
## 1 0.2827 0.06771
## 2
         0.2940
                     0.07587
## 3
         0.2998
                     0.07881
## 4
        0.2102
                     0.06784
## 5
        0.2487
                     0.06766
## 6
         0.3035
                     0.08284
```

#Displays structure of the dataset
str(breast cancer)

```
## 'data.frame': 569 obs. of 32 variables:
## $ id : int 87139402 8910251 905520 868871 9012568 906539 925291 87880 862989 89827 ... ## $ diagnosis : Factor w/ 2 levels "B", "M": 1 1 1 1 1 1 1 2 1 1 ...
## $ diagnosis : Factor w/ 2 levels "B", "M": 1 1 1 1 1 1 2 1 1 ...
## $ radius_mean : num 12.3 10.6 11 11.3 15.2 ...
## $ texture_mean : num 12.4 18.9 16.8 13.4 13.2 ...
## $ perimeter_mean : num 78.8 69.3 70.9 73 97.7 ...
## $ area mean : num 464 346 373 385 712 ...
## $ smoothness_mean : num 0.1028 0.0969 0.1077 0.1164 0.0796 ...
## $ compactness mean : num   0.0698   0.1147   0.078   0.1136   0.0693   ...
## $ concavity_mean : num 0.0399 0.0639 0.0305 0.0464 0.0339 ...
## $ points_mean : num 0.037 0.0264 0.0248 0.048 0.0266 ...
## $ symmetry_mean : num 0.196 0.192 0.171 0.177 0.172 ...
## $ dimension_mean : num 0.0595 0.0649 0.0634 0.0607 0.0554 ...
## $ area_se : num 0.236 0.451 0.197 0.338 0.412 ...
## $ area_se : num 1.67 3.43 1.34 1.85 1 34
## $ area_se : num 1.7
## $ smoothness_se : num 0.00805 0.00747 0.00516 0.01127 0.00501 ...
## $ compactness_se : num 0.0118 0.03581 0.00936 0.03498 0.01485 ...
## $ concavity_se : num   0.0168   0.0335   0.0106   0.0219   0.0155   ...   ## $ points_se : num   0.01241   0.01365   0.00748   0.01965   0.00915   ...   ## $ symmetry_se : num   0.0192   0.035   0.0172   0.0158   0.0165   ...
## $ dimension_se : num 0.00225 0.00332 0.0022 0.00344 0.00177 ...
## $ radius_worst : num 13.5 11.9 12.4 11.9 16.2 ...
## $ texture_worst : num 15.6 22.9 26.4 15.8 15.7 ...
## $ perimeter_worst : num 87 78.3 79.9 76.5 104.5 ...
## $ area_worst : num 549 425 471 434 819 ...
## $ smoothness_worst : num 0.139 0.121 0.137 0.137 0.113 ...
## $ compactness_worst: num 0.127 0.252 0.148 0.182 0.174 ...
## $ concavity_worst : num 0.1242 0.1916 0.1067 0.0867 0.1362 ...
                          : num 0.0939 0.0793 0.0743 0.0861 0.0818 ...
## $ points_worst
## $ symmetry_worst : num 0.283 0.294 0.3 0.21 0.249 ...
## $ dimension worst : num 0.0677 0.0759 0.0788 0.0678 0.0677 ...
```

#Displays the names of the columns
names(breast_cancer)

```
## [1] "id"
                      "diagnosis"
                                       "radius_mean"
## [4] "texture_mean" "perimeter_mean"
                                       "area_mean"
## [7] "smoothness mean" "compactness mean" "concavity mean"
## [10] "points_mean" "symmetry_mean" "dimension_mean"
## [13] "radius_se"
## [16] "area_se"
                      "texture_se"
                                       "perimeter_se"
                      "smoothness_se" "compactness_se"
## [25] "perimeter_worst" "area_worst"
                                       "smoothness_worst"
## [28] "compactness_worst" "concavity_worst" "points_worst"
                      "dimension_worst"
## [31] "symmetry worst"
```

#Displays the summary of the dataset
summary(breast cancer)

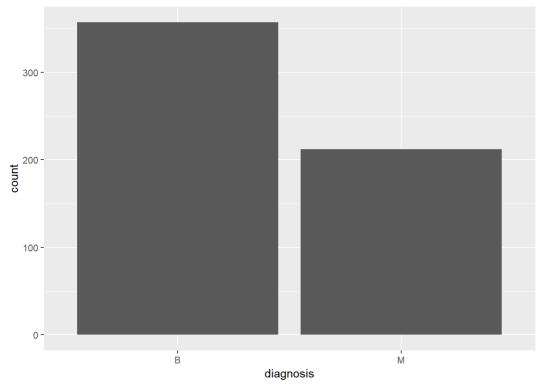
```
diagnosis radius_mean texture_mean
            8670 B:357 Min. : 6.981 Min. : 9.71
## Min. :
                             1st Qu.:11.700 1st Qu.:16.17
##
   1st Qu.:
            869218 M:212
## Median : 906024
                            Median :13.370 Median :18.84
## Mean : 30371831
                            Mean :14.127 Mean :19.29
## 3rd Qu.: 8813129
                            3rd Qu.:15.780 3rd Qu.:21.80
## Max. :911320502
                            Max. :28.110 Max. :39.28
## perimeter_mean area_mean smoothness_mean compactness_mean
## Min. : 43.79 Min. : 143.5 Min. :0.05263 Min. :0.01938
## 1st Qu.: 75.17 1st Qu.: 420.3 1st Qu.:0.08637 1st Qu.:0.06492
## Median: 86.24 Median: 551.1 Median: 0.09587 Median: 0.09263
## Mean : 91.97 Mean : 654.9 Mean :0.09636 Mean :0.10434
##
   3rd Qu.:104.10
                 3rd Qu.: 782.7
                                3rd Qu.:0.10530
                                               3rd Qu.:0.13040
## Max. :188.50
                 Max. :2501.0
                               Max. :0.16340
                                               Max. :0.34540
##
   concavity_mean
                  points_mean
                                symmetry_mean
                                                dimension mean
                 Min. :0.00000 Min. :0.1060 Min. :0.04996
## Min. :0.00000
## 1st Qu.:0.02956 1st Qu.:0.02031 1st Qu.:0.1619 1st Qu.:0.05770
## Median :0.06154
                 Median: 0.03350 Median: 0.1792 Median: 0.06154
## Mean :0.08880 Mean :0.04892 Mean :0.1812 Mean :0.06280
## 3rd Qu.:0.13070 3rd Qu.:0.07400 3rd Qu.:0.1957 3rd Qu.:0.06612
## Max. :0.42680 Max. :0.20120 Max. :0.3040 Max. :0.09744
##
   radius se
                  texture se
                                perimeter_se
                                                area se
## Min. :0.1115 Min. :0.3602 Min. :0.757 Min. : 6.802
## 1st Qu.:0.2324 1st Qu.:0.8339 1st Qu.: 1.606 1st Qu.: 17.850
## Median :0.3242 Median :1.1080 Median : 2.287 Median : 24.530
                               Mean : 2.866
## Mean :0.4052 Mean :1.2169
                                              Mean : 40.337
##
   3rd Qu.:0.4789
                 3rd Qu.:1.4740
                                3rd Qu.: 3.357
                                               3rd Qu.: 45.190
                 Max. :4.8850 Max. :21.980
##
   Max. :2.8730
                                              Max. :542.200
##
   smoothness se
                 compactness se concavity se
## Min. :0.001713 Min. :0.002252 Min. :0.00000
## 1st Qu.:0.005169 1st Qu.:0.013080 1st Qu.:0.01509
## Median: 0.006380 Median: 0.020450 Median: 0.02589
## Mean :0.007041 Mean :0.025478 Mean :0.03189
## 3rd Qu.:0.008146 3rd Qu.:0.032450 3rd Qu.:0.04205
## Max. :0.031130 Max. :0.135400 Max. :0.39600
##
   points se
                   symmetry se
                                   dimension se
                                                     radius worst
## Min. :0.000000 Min. :0.007882 Min. :0.0008948 Min. :7.93
## 1st Qu.:0.007638 1st Qu.:0.015160 1st Qu.:0.0022480 1st Qu.:13.01
## Median :0.010930 Median :0.018730 Median :0.0031870
                                                     Median :14.97
##
   Mean :0.011796 Mean :0.020542
                                   Mean :0.0037949
                                                     Mean :16.27
   3rd Qu.:0.014710
                   3rd Qu.:0.023480
                                   3rd Qu.:0.0045580
                                                     3rd Ou.:18.79
   Max. :0.052790 Max. :0.078950 Max. :0.0298400
##
                                                    Max. :36.04
##
   texture worst perimeter worst area worst smoothness worst
## Min. :12.02 Min. :50.41 Min. :185.2 Min. :0.07117
## 1st Qu.:21.08 1st Qu.: 84.11 1st Qu.: 515.3 1st Qu.:0.11660
## Median: 25.41 Median: 97.66 Median: 686.5 Median: 0.13130
## Mean :25.68 Mean :107.26 Mean :880.6 Mean :0.13237
## 3rd Qu.:29.72 3rd Qu.:125.40 3rd Qu.:1084.0 3rd Qu.:0.14600
## Max. :49.54 Max. :251.20 Max. :4254.0 Max. :0.22260
## compactness_worst concavity_worst points_worst symmetry_worst
## Min. :0.02729 Min. :0.0000 Min. :0.0000 Min. :0.1565
## 1st Qu.:0.14720 1st Qu.:0.1145 1st Qu.:0.06493 1st Qu.:0.2504
## Median :0.21190
                  Median :0.2267
                                 Median :0.09993 Median :0.2822
                                 Mean :0.11461 Mean :0.2901
3rd Qu::0.16140 3rd Qu::0.3179
##
   Mean :0.25427
                  Mean :0.2722
                  3rd Qu.:0.3829 3rd Qu.:0.16140
   3rd Qu.:0.33910
##
                  Max. :1.2520 Max. :0.29100 Max. :0.6638
## Max. :1.05800
## dimension_worst
## Min. :0.05504
## 1st Ou.:0.07146
## Median :0.08004
## Mean :0.08395
## 3rd Qu.:0.09208
## Max. :0.20750
```

```
#To display the frequency table
diagnosis.table <- table(breast_cancer$diagnosis)

#Displays the table
#This shows how many patients are benign and malignant
diagnosis.table</pre>
```

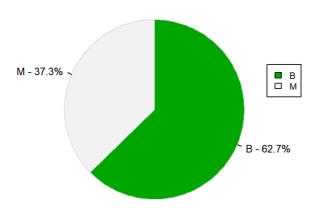
```
##
## B M
## 357 212
```

```
#Generate barplot
ggplot(data=breast_cancer, aes(x=diagnosis)) + geom_bar(stat = "count")
```



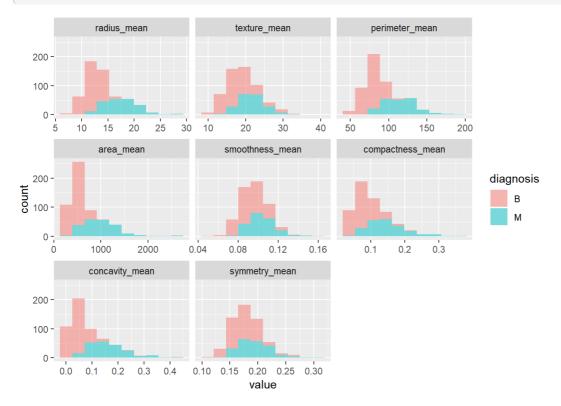
```
#Generate Pie chart represented in frequency
diagnosis.prop.table <- prop.table(diagnosis.table)*100
diagnosis.prop.df <- as.data.frame(diagnosis.prop.table)
pielabels <- sprintf("%s - %3.1f%s", diagnosis.prop.df[,1], diagnosis.prop.table, "%")
colors <- terrain.colors(2)
pie(diagnosis.prop.table,
    labels=pielabels,
    clockwise=TRUE,
    col=colors,
    border="gainsboro",
    radius=0.8,
    cex=0.8,
    main="frequency of cancer diagnosis")
legend(1, .4, legend=diagnosis.prop.df[,1], cex = 0.7, fill = colors)</pre>
```

frequency of cancer diagnosis

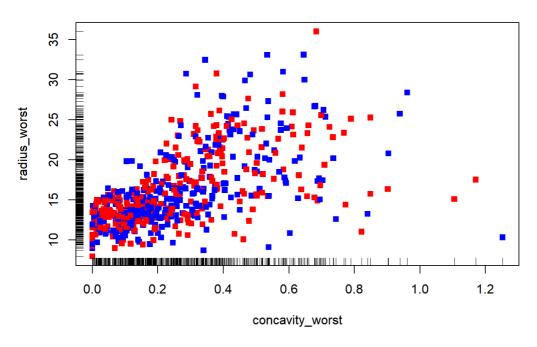


```
#To Plot histograms of "mean" variables group by diagnosis
data_mean <- breast_cancer[ ,c("diagnosis", "radius_mean", "texture_mean", "perimeter_mean", "area_mean", "sm
oothness_mean", "compactness_mean", "concavity_mean", "symmetry_mean" )]

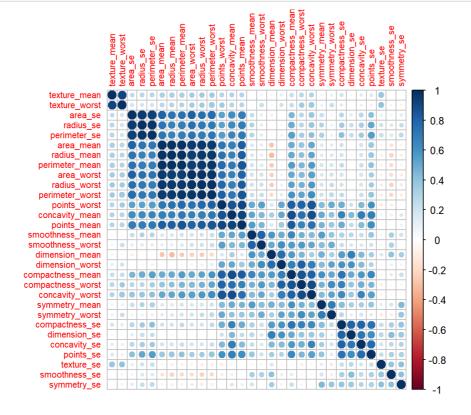
#Plot histograms
ggplot(data = melt(data_mean, id.var = "diagnosis"), mapping = aes(x = value)) +
    geom_histogram(bins = 10, aes(fill=diagnosis), alpha=0.5) + facet_wrap(~variable, scales ='free_x')</pre>
```



Concavity_worst vs radius_worst

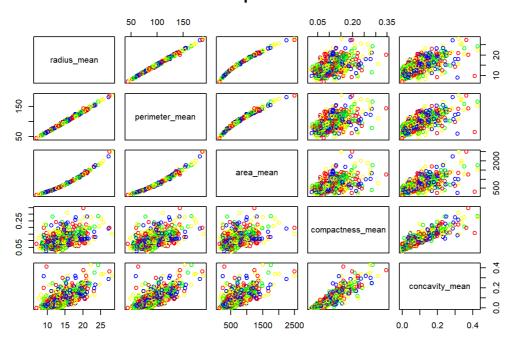


```
#Generate Corelation Matrix of columns
corMatMy <- cor(breast_cancer[,3:32])
corrplot(corMatMy, order = "hclust", tl.cex = 0.7)</pre>
```



```
#Generate Scatterplot Matrix
pairs(~radius_mean+perimeter_mean+area_mean+compactness_mean+concavity_mean,data = breast_cancer,main = "Scatterplot Matrix",col=c("red","blue","green","yellow"))
```

Scatterplot Matrix



```
#Multivariate analysis
#t-tEST
with(data=breast_cancer,t.test(radius_mean[diagnosis=="B"],radius_mean[diagnosis=="M"],var.equal=TRUE))
```

```
##
## Two Sample t-test
##
## data: radius_mean[diagnosis == "B"] and radius_mean[diagnosis == "M"]
## t = -25.436, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -5.726832 -4.905781
## sample estimates:
## mean of x mean of y
## 12.14652 17.46283</pre>
```

with(data=breast_cancer,t.test(texture_mean[diagnosis=="B"],texture_mean[diagnosis=="M"],var.equal=TRUE))

```
##
## Two Sample t-test
##
## data: texture_mean[diagnosis == "B"] and texture_mean[diagnosis == "M"]
## t = -10.867, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -4.357107 -3.023181
## sample estimates:
## mean of x mean of y
## 17.91476 21.60491</pre>
```

```
with(data=breast_cancer,t.test(perimeter_mean[diagnosis=="B"],perimeter_mean[diagnosis=="M"],var.equal=TRUE)
)
```

```
##
## Two Sample t-test
##
## data: perimeter_mean[diagnosis == "B"] and perimeter_mean[diagnosis == "M"]
## t = -26.405, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -40.06379 -34.51615
## sample estimates:
## mean of x mean of y
## 78.07541 115.36538</pre>
```

with(data=breast_cancer,t.test(area_mean[diagnosis=="B"],area_mean[diagnosis=="M"],var.equal=TRUE))

```
##
## Two Sample t-test
##
## data: area_mean[diagnosis == "B"] and area_mean[diagnosis == "M"]
## t = -23.939, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -557.8898 -473.2826
## sample estimates:
## mean of x mean of y
## 462.7902 978.3764</pre>
```

 $\label{lem:with(data=breast_cancer, t.test(smoothness_mean[diagnosis=="B"], smoothness_mean[diagnosis=="M"], var.equal=TRUE))$

```
##
## Two Sample t-test
##
## data: smoothness_mean[diagnosis == "B"] and smoothness_mean[diagnosis == "M"]
## t = -9.1461, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.012658756 -0.008182931
## sample estimates:
## mean of x mean of y
## 0.09247765 0.10289849</pre>
```

 $\label{lem:with(data=breast_cancer, t.test(compactness_mean[diagnosis=="B"], compactness_mean[diagnosis=="M"], var.equal=TRUE))} \\$

```
##
## Two Sample t-test
##
## data: compactness_mean[diagnosis == "B"] and compactness_mean[diagnosis == "M"]
## t = -17.698, df = 567, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.07232827 -0.05787805
## sample estimates:
## mean of x mean of y
## 0.08008462 0.14518778</pre>
```

with(data=breast_cancer,t.test(concavity_mean[diagnosis=="B"],concavity_mean[diagnosis=="M"],var.equal=TRUE)
)

```
## Two Sample t-test
\#\,\#
## data: concavity_mean[diagnosis == "B"] and concavity_mean[diagnosis == "M"]
## t = -23.104, df = 567, p-value < 2.2e-16
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.1244696 -0.1049646
## sample estimates:
## mean of x mean of y
## 0.04605762 0.16077472
with(data=breast_cancer,t.test(points_mean[diagnosis=="B"],points_mean[diagnosis=="M"],var.equal=TRUE))
## Two Sample t-test
##
## data: points mean[diagnosis == "B"] and points mean[diagnosis == "M"]
## t = -29.354, df = 567, p-value < 2.2e-16
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
   -0.06643938 -0.05810581
## sample estimates:
## mean of x mean of y
## 0.02571741 0.08799000
with(data=breast_cancer,t.test(symmetry_mean[diagnosis=="B"],symmetry_mean[diagnosis=="M"],var.equal=TRUE))
##
## Two Sample t-test
##
## data: symmetry_mean[diagnosis == "B"] and symmetry_mean[diagnosis == "M"]
## t = -8.3383, df = 567, p-value = 5.733e-16
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02313331 -0.01431262
## sample estimates:
## mean of x mean of y
## 0.174186 0.192909
with(data=breast_cancer,t.test(dimension_mean[diagnosis=="B"],dimension_mean[diagnosis=="M"],var.equal=TRUE)
)
## Two Sample t-test
##
## data: dimension_mean[diagnosis == "B"] and dimension_mean[diagnosis == "M"]
## t = 0.30571, df = 567, p-value = 0.7599
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.001016083 0.001390684
```

```
#Hotelling's T2 test
#install.packages("Hotelling")
library(Hotelling)
```

sample estimates:
mean of x mean of y
0.06286739 0.06268009

```
## Loading required package: corpcor
```

Warning: package 'Hotelling' was built under R version 3.5.2

```
## Warning: package 'corpcor' was built under R version 3.5.2
```

```
t2testcan <- hotelling.test(radius_mean + texture_mean + perimeter_mean + area_mean + smoothness_mean + comp
actness mean + concavity mean + points mean + symmetry mean + dimension mean ~ diagnosis, data=breast cancer
# Output of the function hotelling.test is given
cat("T2 statistic =",t2testcan$stat[[1]],"\n")
## T2 statistic = 1220.313
print(t2testcan)
## Test stat: 120.09
## Numerator df: 10
## Denominator df: 558
## P-value: 0
# T2 statistic is located in the first element of the list "stat"
#View(t2testcan)
#View(breast cancer)
#Levene's tests based on absolute differences around means using t-tests. Standarizing the data set with sca
matstand <- scale(breast cancer[,3:10])</pre>
head (matstand)
       radius mean texture mean perimeter mean area mean smoothness mean
## [1,] -0.5128453 -1.60418301 -0.5399006 -0.5421468
## [2,] -1.0009202 -0.07896900
                                   -0.9337442 -0.8766033
                                                              0.0369535
## [3,] -0.8760638 -0.57187353 -0.8662517 -0.8004484
                                                             0.8062867
## [4,] -0.8079604 -1.37168088 -0.7806514 -0.7674858
                                                             1.4248817
## [5,] 0.3015589 -1.41353126
                                   0.2337944 0.1617181
                                                             -1.1895712
## [6,] -0.7256686 -0.05804381
                                   -0.7312666 -0.6967299
                                                             -0.7750414
##
       compactness mean concavity mean points mean
## [1,]
           -0.6538379
                         -0.6137661 -0.30717196
## [2,]
             0.1961461
                           -0.3127117 -0.57983238
                          -0.7318045 -0.62158190
## [3,]
             -0.4980044
                          -0.5324814 -0.02471844
## [4,]
             0.1753178
## [5,]
             -0.6627373 -0.6882771 -0.57596668
## [6,]
             -0.5135309
                          -0.4258580 -0.89269604
matben <- matstand[breast_cancer$diagnosis =="B",]</pre>
head (matben)
       radius_mean texture_mean perimeter_mean area_mean smoothness_mean
## [1,] -0.5128453 -1.60418301 -0.5399006 -0.5421468 0.4578825
## [2,] -1.0009202 -0.07896900
                                  -0.9337442 -0.8766033
                                                             0.0369535
## [3,] -0.8760638 -0.57187353
                                  -0.8662517 -0.8004484
                                                             0.8062867
## [4,] -0.8079604 -1.37168088
                                  -0.7806514 -0.7674858
                                                             1.4248817
## [5,] 0.3015589 -1.41353126
                                   0.2337944 0.1617181
                                                             -1.1895712
## [6,] -0.7256686 -0.05804381
                                  -0.7312666 -0.6967299
                                                             -0.7750414
##
       compactness mean concavity mean points mean
## [1,]
          -0.6538379
                         -0.6137661 -0.30717196
## [2,]
              0.1961461
                           -0.3127117 -0.57983238
                           -0.7318045 -0.62158190
             -0.4980044
## [3,]
             0.1753178
                           -0.5324814 -0.02471844
## [4,]
                           -0.6882771 -0.57596668
## [5,]
             -0.6627373
## [6,]
                           -0.4258580 -0.89269604
             -0.5135309
matmalign <- matstand[breast cancer$diagnosis == "M",]</pre>
vecmedianben <- apply(matben, 2, median)</pre>
# in the above 2 represents column. Hence, we are asking for column median
vecmedianben
```

```
##
       radius_mean texture_mean perimeter_mean
                                                          area_mean
##
        -0.5468970
                      -0.4416723
                                      -0.5674737
                                                         -0.5583439
                                    concavity_mean
                                                       points_mean
##
   smoothness_mean compactness_mean
##
        -0.3981961
                      -0.5500751
                                      -0.6486382
                                                         -0.6566309
vecmedianmalign <- apply(matmalign, 2, median)</pre>
matabsdevben <- abs(matben - matrix(rep(vecmedianben,nrow(matben)),nrow=nrow(matben), byrow=TRUE))</pre>
matabsdevmalign <- abs(matmalign - matrix(rep(vecmedianmalign,nrow(matmalign)),nrow=nrow(matmalign), byrow=T</pre>
RUE))
head(matabsdevmalign)
      radius mean texture mean perimeter mean area mean smoothness mean
## [1,] 0.9974323 0.53242989 0.9317263 0.9496635 2.14019666
## [2,] 0.9264911 0.05115047
                                   0.9712341 1.1025417
                                                           0.44794814
## [3,] 0.6427266 0.82305756
                                   0.5555789 0.7359750
                                                           0.07110288
                                   0.8436568 0.8951047
## [4,] 0.7846089 0.55800512
                                                           0.41239670
## [5,] 1.0002699 1.31828711
                                   0.9782303 0.9885934
                                                            0.10665432
## [6,] 0.5746231 0.88583314
                                   0.4856171 0.5671838
                                                            0.42661727
    compactness mean concavity mean points mean
##
## [1,]
          0.84165269
                          0.05582051 0.14122676
## [2,]
             0.60686094
                           0.84608833 0.66541513
                          0.17749666 0.11210518
## [3,]
            0.63696730
                          0.96525570 1.07517890
## [4,]
            0.97419643
## [5,]
                          0.92147737 0.91720079
            0.08236646
## [6,]
           0.70721548
                          0.21261968 0.03169871
matabsdev.all <- rbind(matabsdevben, matabsdevmalign)</pre>
matabsdev.all <- data.frame(breast_cancer$diagnosis, matabsdev.all)</pre>
t.test(matabsdev.all$radius mean[breast cancer$diagnosis == "B"], matabsdev.all$radius mean[breast cancer$dia
gnosis == "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$radius mean[breast cancer$diagnosis == "B"] and matabsdev.all$radius mean[breast can
cer$diagnosis == "M"]
## t = 0.32562, df = 567, p-value = 0.6276
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
       -Inf 0.07485419
##
## sample estimates:
## mean of x mean of y
## 0.5301158 0.5177632
t.test(matabsdev.all$texture_mean[breast_cancer$diagnosis == "B"],matabsdev.all$texture_mean[breast_cancer$d
iagnosis == "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$texture mean[breast cancer$diagnosis == "B"] and matabsdev.all$texture mean[breast c
ancer$diagnosis == "M"]
## t = -2.1618, df = 567, p-value = 0.01553
\#\# alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
          -Inf -0.02723094
## sample estimates:
## mean of x mean of v
## 0.6364762 0.7509490
```

t.test(matabsdev.all\$perimeter_mean[breast_cancer\$diagnosis == "B"],matabsdev.all\$perimeter_mean[breast_cancer\$diagnosis == "M"], alternative="less",var.equal = TRUE)

```
##
## Two Sample t-test
\#\,\#
## data: matabsdev.all$perimeter_mean[breast_cancer$diagnosis == "B"] and matabsdev.all$perimeter_mean[brea
st_cancer$diagnosis == "M"]
## t = 0.2439, df = 567, p-value = 0.5963
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
        -Inf 0.07148672
## sample estimates:
## mean of x mean of v
## 0.5125724 0.5033541
t.test(matabsdev.all$area_mean[breast_cancer$diagnosis == "B"], matabsdev.all$area_mean[breast_cancer$diagnos
is == "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$area mean[breast cancer$diagnosis == "B"] and matabsdev.all$area mean[breast cancer$
diagnosis == "M"]
## t = 0.40112, df = 567, p-value = 0.6558
\#\# alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
        -Inf 0.0909786
##
## sample estimates:
## mean of x mean of y
## 0.4981297 0.4803166
t.test(matabsdev.all$smoothness_mean[breast_cancer$diagnosis == "B"], matabsdev.all$smoothness_mean[breast_ca
ncer$diagnosis == "M"], alternative="less", var.equal = TRUE)
##
##
  Two Sample t-test
##
## data: matabsdev.all$smoothness mean[breast cancer$diagnosis == "B"] and matabsdev.all$smoothness mean[br
east cancer$diagnosis == "M"]
## t = 1.6742, df = 567, p-value = 0.9527
\#\# alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
       -Inf 0.167207
##
## sample estimates:
## mean of x mean of y
## 0.7680704 0.6837950
t.test(matabsdev.all$compactness mean[breast cancer$diagnosis == "B"], matabsdev.all$compactness mean[breast
cancer$diagnosis == "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$compactness mean[breast cancer$diagnosis == "B"] and matabsdev.all$compactness mean[
breast_cancer$diagnosis == "M"]
## t = 1.8406, df = 567, p-value = 0.9669
\#\# alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
       -Inf 0.1710355
## sample estimates:
## mean of x mean of y
## 0.6249227 0.5346711
```

t.test(matabsdev.all\$concavity_mean[breast_cancer\$diagnosis == "B"],matabsdev.all\$concavity_mean[breast_canc
er\$diagnosis == "M"], alternative="less",var.equal = TRUE)

```
## Two Sample t-test
##
## data: matabsdev.all$concavity_mean[breast_cancer$diagnosis == "B"] and matabsdev.all$concavity_mean[brea
st_cancer$diagnosis == "M"]
## t = 1.0995, df = 567, p-value = 0.864
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
## -Inf 0.1302286
## sample estimates:
## mean of x mean of v
## 0.4977532 0.4456302
t.test(matabsdev.all$points_mean[breast_cancer$diagnosis == "B"],matabsdev.all$points_mean[breast_cancer$diagnosis == "B"]
gnosis == "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$points mean[breast cancer$diagnosis == "B"] and matabsdev.all$points mean[breast can
cer$diagnosis == "M"]
## t = 0.31387, df = 567, p-value = 0.6231
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
       -Inf 0.07740908
##
## sample estimates:
## mean of x mean of y
## 0.4434506 0.4310634
head (mat.st.and)
       radius_mean texture_mean perimeter_mean area_mean smoothness_mean
```

```
## [1,] -0.5128453 -1.60418301 -0.5399006 -0.5421468 0.4578825
## [2,] -1.0009202 -0.07896900
                                -0.9337442 -0.8766033
                                                         0.0369535
## [3,] -0.8760638 -0.57187353
                                -0.8662517 -0.8004484
## [4,] -0.8079604 -1.37168088
                                -0.7806514 -0.7674858
                                                         1.4248817
## [5,] 0.3015589 -1.41353126
                                0.2337944 0.1617181
                                                         -1.1895712
## [6,] -0.7256686 -0.05804381
                                -0.7312666 -0.6967299
                                                         -0.7750414
##
     compactness_mean concavity_mean points_mean
## [1,] -0.6538379 -0.6137661 -0.30717196
            0.1961461
                         -0.3127117 -0.57983238
## [2.1
## [3,]
            -0.4980044
                         -0.7318045 -0.62158190
            0.1753178
                         -0.5324814 -0.02471844
## [4,]
                         -0.6882771 -0.57596668
## [5,]
            -0.6627373
           -0.5135309
                        -0.4258580 -0.89269604
## [6,]
```

matstand.all <- data.frame(breast_cancer\$diagnosis, matstand)
head(matstand.all)</pre>

```
## breast cancer.diagnosis radius mean texture mean perimeter mean
## 1
                        B -0.5128453 -1.60418301
                        B -1.0009202 -0.07896900
## 2
                                                    -0.9337442
## 3
                        B -0.8760638 -0.57187353
                                                    -0.8662517
                        B -0.8079604 -1.37168088
## 4
                                                    -0.7806514
## 5
                        B 0.3015589 -1.41353126
                                                    0.2337944
                        B -0.7256686 -0.05804381
                                                  -0.7312666
## area mean smoothness mean compactness mean concavity mean points mean
## 1 -0.5421468 0.4578825 -0.6538379 -0.6137661 -0.30717196
## 2 -0.8766033
                  0.0369535
                                  0.1961461
                                              -0.3127117 -0.57983238
## 3 -0.8004484
                  0.8062867
                                 -0.4980044 -0.7318045 -0.62158190
## 4 -0.7674858
                  1.4248817
                                  0.1753178
                                               -0.5324814 -0.02471844
## 5 0.1617181
                  -1.1895712
                                  -0.6627373
                                                -0.6882771 -0.57596668
## 6 -0.6967299
                                  -0.5135309
                                                -0.4258580 -0.89269604
                  -0.7750414
```

```
colnames(matstand.all) <- colnames(breast_cancer[2:10])</pre>
t2testcan <- hotelling.test(radius mean + texture mean + perimeter mean + area mean + smoothness mean + comp
actness mean + concavity mean + points mean + symmetry mean + dimension mean ~ diagnosis, data=breast_cancer
cat("T2 statistic =",t2testcan$stat[[1]],"\n")
## T2 statistic = 1220.313
print(t2testcan)
## Test stat: 120.09
## Numerator df: 10
## Denominator df: 558
## P-value: 0
# In the above we standardized using scale function
head(matabsdev.all)
## breast cancer.diagnosis radius mean texture mean perimeter mean
                                                 0.02757317
                        B 0.03405174 1.1625107
## 1
## 2
                        B 0.45402322
                                       0.3627033
                                                    0.36627050
                                      0.1302012
## 3
                                                   0.29877796
                        B 0.32916684
                        В 0.26106335 0.9300085
                                                   0.21317766
## 4
## 5
                        В 0.84845589 0.9718589
                                                  0.80126817
## 6
                        B 0.17877164 0.3836285 0.16379288
##
   area mean smoothness mean compactness mean concavity mean points mean
## 2 0.31825946
                  0.4351496
                                 0.74622121
                                              0.33592655 0.07679849
## 3 0.24210452
                  1.2044828
                                0.05207075 0.08316628 0.03504898
## 4 0.20914193
                  1.8230778
                                 0.72539291 0.11615683 0.63191243
                  0.7913750
0.3768453
## 5 0.72006202
                                 0.11266217 0.03963883 0.08066419
                                 ## 6 0.13838603
#install.packages("car")
library(car)
## Warning: package 'car' was built under R version 3.5.2
## Loading required package: carData
#leveneTest() produces a two-sided test
# Leverne test is used to verify Homoscedasticity. It tests if the variance of two samples are # #equal. Lev
ene's test is an inferential statistic used to assess the equality of variances for a #variable calculated f
or two or more groups.[1] Some common statistical procedures assume that #variances of the populations from
which different samples are drawn are equal. Levene's test #assesses this assumption.
leveneTest(radius_mean ~ diagnosis, data=breast_cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 90.477 < 2.2e-16 ***
##
       567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(texture_mean ~ diagnosis, data=breast_cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
            0.684 0.4086
## group 1
```

##

567

leveneTest(perimeter_mean ~ diagnosis, data=breast_cancer)

```
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 91.237 < 2.2e-16 ***
##
       567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(area_mean ~ diagnosis, data=breast_cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 170.21 < 2.2e-16 ***
##
       567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(smoothness_mean ~ diagnosis, data=breast_cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 0.8377 0.3604
leveneTest(compactness mean~ diagnosis, data=breast cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 39.892 5.428e-10 ***
##
       567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(concavity mean~ diagnosis, data=breast cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 70.484 3.723e-16 ***
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(points mean ~ diagnosis, data=breast cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 94.906 < 2.2e-16 ***
##
       567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(symmetry_mean ~ diagnosis, data=breast_cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 2.036 0.1542
##
       567
```

```
leveneTest(dimension_mean ~ diagnosis, data=breast_cancer)
```

```
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 6.113 0.01371 *
## 567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
#PCA
dim(breast cancer)
## [1] 569 32
attach (breast_cancer)
head(breast_cancer)
    id diagnosis radius_mean texture_mean perimeter_mean area_mean
## 1 87139402 B 12.32 12.39 78.85 464.1
## 2 8910251 B 10.60 18.95 69.28 346.4
                В
                                             69.28
## 3 905520
                В
                      11.04
                                 16.83
                                             70.92
                                                     373.2
                                 13.39
## 4 868871
                В
                      11.28
                                             73.00
## 5 9012568
                В
                      15.19
                                 13.21
                                             97.65
                B 11.57 19.04
## 6 906539
                                             74.20
## smoothness_mean compactness_mean concavity_mean points_mean
## 1 0.10280 0.06981 0.03987 0.03700
                                  0.06387
         0.09688
                      0.11470
                                            0.02642
## 2
                                  0.03046
## 3
         0.10770
                      0.07804
                                             0.02480
## 4
         0.11640
                       0.11360
                                   0.04635
                                             0.04796
## 5
          0.07963
                       0.06934
                                   0.03393
                                             0.02657
                      0.07722
## 6
         0.08546
                                   0.05485
                                             0.01428
## symmetry_mean_dimension_mean_radius_se_texture_se_perimeter_se_area_se
     0.1959 0.05955 0.2360 0.6656 1.670 17.43
## 1
                   0.06491 0.4505
                                     1.1970
                                                3.430 27.10
## 2
         0.1922
        0.1714
                   0.06340 0.1967 1.3870
                                                1.342 13.54
## 3
## 4
        0.1771
                   0.06072 0.3384 1.3430
                                                1.851 26.33
                   0.05544 0.1783 0.4125
## 5
        0.1721
                                                1.338 17.72
       0.2031 0.06267 0.2864 1.4400 2.206 20.30
## 6
## smoothness_se compactness_se concavity_se points_se symmetry_se
## 1 0.008045 0.011800 0.01683 0.012410 0.01924
       0.007470
## 2
                   0.035810
                              0.03354 0.013650
                                                0.03504
                              0.01056 0.007483
       0.005158
## 3
                   0.009355
                                                0.01718
       0.011270
                   0.034980
                              0.02187 0.019650
                                                0.01580
                              0.01551 0.009155
## 5
       0.005012
                   0.014850
                                                0.01647
      0.01868
## 6
## dimension_se radius_worst texture_worst perimeter_worst area_worst
## 1 0.002248 13.50 15.64 86.97 549.1
    0.003318
                              22.94
                                           78.28
                                                   424.8
## 2
                   11.88
## 3 0.002198
                  12.41
                              26.44
                                           79.93
                   11.92
                              15.77
                                           76.53
## 4 0.003442
                                                   434.0
## 5
    0.001767
                  16.20
                              15.73
                                          104.50
                                                   819.1
## 6 0.003339 13.07 26.98 86.43
                                                   520.5
## smoothness worst compactness worst concavity worst points worst
## 1 0.1385 0.1266 0.12420 0.09391
## 2
           0.1213
                         0.2515
                                     0.19160
                                                0.07926
## 3
           0.1369
                          0.1482
                                     0.10670
                                                0.07431
## 4
           0.1367
                          0.1822
                                      0.08669
                                                0.08611
## 5
           0.1126
                          0.1737
                                     0.13620
                                                0.08178
                         0.1937
## 6
           0.1249
                                     0.25600
                                                0.06664
## symmetry_worst dimension_worst
## 1
    0.2827 0.06771
## 2
         0.2940
                     0.07587
## 3
         0.2998
                     0.07881
## 4
         0.2102
                     0.06784
## 5
         0.2487
                     0.06766
         0.3035
## 6
                     0.08284
```

#Get the Correlations between the measurements cor(breast_cancer[-2])

```
id radius mean texture mean perimeter mean
                 1.000000000 0.074626470 0.099769891 0.073159412
 ## id
 ## smoothness mean -0.0129681975 0.170581187 -0.023388516 0.207278164
 ## compactness_mean 0.0000957011 0.506123578 0.236702222 0.556936211
 ## concavity_mean 0.0500799532 0.676763550 0.302417828 0.716135650
 ## points_mean
                 0.0441580956 0.822528522 0.293464051 0.850977041
                 -0.0221140609 0.147741242 0.071400980 0.183027212
 ## symmetry_mean
 ## dimension_mean -0.0525114476 -0.311630826 -0.076437183 -0.261476908
                 0.1430475814 0.679090388 0.275868676
 0.691765014
                                                    -0.086761078
 ## perimeter_se
                  0.1373310660 0.674171616 0.281673115
                                                     0.693134890
                                                   0.744982694
                 0.1777419152 0.735863663 0.259844987
 ## area_se
 ## smoothness_se
                 0.0967805739 -0.222600125 0.006613777 -0.202694026
 ## compactness_se 0.0339609721 0.205999980 0.191974611 0.250743681
## concavity_se 0.0552393174 0.194203623 0.143293077 0.228082345
## points se 0.0787680711 0.376168956 0.163851025 0.407216916
 ## points_se
## symmetry_se
                 0.0787680711 0.376168956 0.163851025 0.407216916
                -0.0173062948 -0.104320881 0.009127168 -0.081629327
 ## dimension_se
                 0.0257253243 -0.042641269 0.054457520 -0.005523391
 ## radius_worst
## texture_worst
                 0.0824053373 0.969538973 0.352572947 0.969476363
                 0.0647195454 0.297007644 0.912044589 0.303038372
 ## perimeter_worst 0.0799858731 0.965136514 0.358039575 0.970386887
 ## area_worst 0.1071865233 0.941082460 0.343545947 ## smoothness_worst 0.0103380343 0.119616140 0.077503359
                                                    0.941549808
                                                     0.150549404
 ## compactness_worst -0.0029680998 0.413462823 0.277829592
                                                     0.455774228
 0.563879263
 0.771240789
                                                   0.189115040
 ## dimension_worst -0.0298656360 0.007065886 0.119205351 0.051018530
 ##
                  area mean smoothness mean compactness mean
                 0.096892823 -0.01296820 0.0000957011
 ## id
 ## radius_mean 0.987357170
## texture_mean 0.321085696
                               0.17058119 0.5061235775
                               -0.02338852 0.2367022221
 1.000000000
                               0.17702838 0.4985016822
 ## area_mean
 ## smoothness_mean
                  0.177028377
                               1.00000000
                                            0.6591232152
                               0.65912322
 ## compactness_mean 0.498501682
                                            1.00000000000
 ## concavity_mean
                  0.685982829
                                0.52198377
                                             0.8831206702
 ## points_mean
## symmetry_mean
                               0.55369517
                  0.823268869
                                            0.8311350431
                 0.151293079
                               0.55777479 0.6026410484
## dimension_mean -0.283109812
                 0.37232020
-0.072496588
-0.19886963
0.28360670
0.21312014
                                            0.5353153982
                               0.03607180 0.2481328333
 ## perimeter_worst 0.959119574
                               0.23885263 0.5902104277
 ## compactness_worst 0.390410309
                               0.47246844 0.8658090398
 ## concavity_worst 0.512605920
                               0.43492571 0.8162752498
 ## points_worst 0.722016626
                               ## symmetry_worst 0.143569914 0.39430948 0.5102234299
## dimension_worst 0.003737597 0.49931637 0.6873823228
        concavity_mean points_mean symmetry_mean dimension mean
 ##
                 0.05007995 0.04415810 -0.02211406 -0.0525114476
 ## id
```

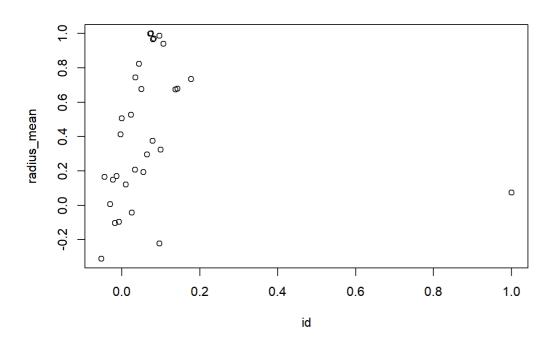
```
1.00000000 0.92139103 0.50066662 0.3367833594
## concavity_mean
## area mean 0.7325622270 -0.06628021 0.72662833 0.80008592
## compactness_mean  0.4974734461  0.04620483  0.54890526  0.45565285
## concavity_mean 0.6319248221 0.07621835 0.66039079 0.61742681
## smoothness_worst 0.1419185529 -0.07365766 0.13005439 0.12538943
## compactness_worst 0.2871031656 -0.09243935 0.34191945 0.28325654
## points_worst 0.5310623278 -0.11963752 0.55489723 0.53816631
## symmetry_worst 0.0945428304 -0.12821476 0.10993043 0.07412629
## dimension_worst 0.0495594325 -0.04565457 0.08543257 0.01753930
  ##
## id
        ## radius_mean
## texture_mean
-0.222600125 0.20599998 0.19420362 0.37616896
## smoothness se 1.000000000 0.33669608 0.26868476 0.32842950
```

```
## concavity_se
                                    ## points_se
                                    0.328429499 0.74408267 0.77180399 1.00000000
## area_worst
 ## smoothness_worst 0.314457456 0.22739423 0.16848132 0.21535060
 ## compactness worst -0.055558139 0.67878035 0.48485780 0.45288838
 ## concavity_worst -0.058298387 0.63914670 0.66256413 0.54959238
## area mean
 ## smoothness_mean 0.200774376 0.283606699 0.21312014 0.036071799
 ## compactness_mean 0.229976591 0.507318127 0.53531540 0.248132833
## dimension_mean 0.345007397 0.688131577 -0.25369149 -0.051269202
## perimeter_worst -0.103753044 -0.001000398 0.99370792 0.365098245
 ## area_worst -0.110342743 -0.022736147 0.98401456 0.345842283
 ## smoothness_worst -0.012661800 0.170568316 0.21657443 0.225429415
 ## compactness_worst 0.060254879 0.390158842 0.47582004 0.360832339
perimeter worst area worst smoothness worst
 ##
## id 0.079985873 0.10718652 0.01033803  
## radius_mean 0.965136514 0.94108246 0.11961614  
## texture_mean 0.358039575 0.34354595 0.07750336  
## perimeter_mean 0.970386887 0.94154981 0.15054940  
## area_mean 0.959119574 0.95921333 0.12352294  
## smoothness_mean 0.238852626 0.20671836 0.80532420  
## compactness_mean 0.590210428 0.50960381 0.56554117
## concavity_mean 0.729564917 0.67598723
## concavity_mean
## points_mean
## symmetry_mean
## dimension_mean
## texture_se
## smoothness_se
## compactness_se
## concavity_se
## goints_se
## concavity_se
## dimension_se
## concavity_se
## texture_se
## compactness_se
## concavity_se
## concavity_se
## symmetry_se
## dimension_se
## symmetry_se
## dimension_se
## symmetry_se
## dimension_se
## concavity_se
## dimension_se
## symmetry_se
## dimension_se
## concavity_se
## dimension_se
## concavity_se
## dimension_se
## dimension_se
## dimension_se
## concavity_se
## dimension_se
## dimension_se
## concavity_se
## dimension_se
## dimension_se
## dimension_se
## dimension_se
## concavity_se
## dimension_se

                                                                                          0.44882204
                                                                                             0.45275305
                                                                                             0.42667503
                                                                                            0.50494208
                                                                                           0.14191855
                                                                                         -0.07365766
                                                                                           0.13005439
                                                                                           0.12538943
                                                                                           0.31445746
                                                                                           0.22739423
                                                                                           0.16848132
                                                                                           0.21535060
                                                                                          -0.01266180
                                                                                          0.17056832
                                                                                           0.21657443
                                                                                           0.22542941
0.23677460
```

```
## smoothness_worst 0.236774604 0.20914533 ## compactness_worst 0.529407690 0.43829628
                                       U. ZUJI4JJJ
                                       1.00000000
                                      0.56818652
0.51852329
## points_worst
## symmetry_worst
                                      0.54769090
## symmetry_worst 0.269492769 0.20914551 0.49383833
## dimension_worst 0.138956862 0.07964703 0.61762419
     compactness_worst concavity_worst points_worst -0.00296810 0.02320274 0.03517350
              ## id
## radius_mean
## texture_mean
## perimeter_mean
                   0.41346282
                              0.52691146 0.74421420
                   0.27782959
                              0.30102522 0.29531584
                  0.45577423
                              0.56387926 0.77124079
## area_mean
                   0.39041031
                              0.51260592 0.72201663
    ##
              -0.04422425 -0.029865636
0.16395333 0.007065886
0.10500791 0.119205351
## id
## radius_mean
## texture_mean
## symmetry_se
## points_worst
## symmetry_worst
## dimension_worst
                            0.511114146
                 1.00000000 0.537848206
                0.53784821 1.000000000
```

```
c <- (cor(breast_cancer[-2]))
plot(c)</pre>
```



```
# Using prcomp to compute the principal components (eigenvalues and eigenvectors). With scale=TRUE, variable
means are set to zero, and variances set to one
breast_cancer_pca <- prcomp(breast_cancer[,-2],scale=TRUE)
breast_cancer_pca</pre>
```

```
## Standard deviations (1, ..., p=31):
## [1] 3.64527878 2.38679814 1.68386313 1.40760690 1.28406203 1.11115827
## [7] 0.98907696 0.81960537 0.67881693 0.63492763 0.59089337 0.54211662
## [13] 0.51102537 0.49125372 0.39619900 0.30680373 0.28250655 0.24299439
## [19] 0.22932770 0.22163467 0.17626907 0.17303527 0.16562163 0.15572098
## [25] 0.13431069 0.12441756 0.09039745 0.08305482 0.03986650 0.02735646
## [31] 0.01153431
##
## Rotation (n x k) = (31 \times 31):
##
                                     PC2
                                                 PC3
## id
                  ## radius mean
                  ## texture mean
                  ## perimeter_mean
                  -0.22753491 0.214589002 -0.012124791 0.042752797
                  -0.22104577 0.230668816 0.026293150 0.054114724
## area_mean
                  -0.14241471 -0.186422211 -0.103182400 0.158098177
## smoothness_mean
## compactness_mean -0.23906730 -0.152454726 -0.074768623 0.031818117
## concavity_mean
                  -0.25828025 -0.060541625 0.001758736 0.019497124
                  -0.26073811 0.034167392 -0.027579607
## points_mean
## symmetry_mean
                  -0.13797774 -0.190684979 -0.040962032 0.067502543
## dimension_mean
                  -0.06414779 -0.366531055 -0.020817875 0.047957856
                  -0.20611747 0.105935702 0.266917221 0.099114446
## radius_se
## texture_se
                  -0.01741339 -0.089547789 0.371439885 -0.356497230
## perimeter_se
                  -0.21144652 0.089807043 0.264925682 0.090293055
## area se
                  -0.20307642 0.152771289 0.215790250 0.108568705
## smoothness_se
                  -0.01467821 -0.203189876 0.311787845 0.044368664
## compactness se
                  -0.17028840 -0.232503362 0.154557465 -0.026425360
## concavity_se
                  -0.15354367 -0.196846081 0.176560052 0.002248291
                  -0.18340675 -0.129965181 0.223850479 0.075252232
## points_se
## symmetry_se
                  -0.04241552 -0.183558627 0.285265066 0.046936126
## dimension_se
                  -0.10249607 -0.279584139 0.211893354 0.016212450
## radius worst
                  ## texture_worst
                  -0.10451545 0.045501223 -0.039828934 -0.633119655
## perimeter_worst
                  -0.23663734 0.199295985 -0.050431945 0.014068572
                  -0.22493214 0.218985461 -0.013188891 0.025970672
## area_worst
## smoothness worst -0.12782441 -0.172562959 -0.255328751 0.014523359
```

```
" " DWOOCIIIICDD WOTDC
                 U.IL.ULIII U.I.LUULJUJ
                                    0.200020,01
## compactness_worst -0.20988456 -0.144253637 -0.234513609 -0.092562168
## concavity_worst -0.22860218 -0.098526524 -0.172024941 -0.074807188
## points worst -0.25074620 0.007534367 -0.170480673 0.005305980
## symmetry worst -0.12267993 -0.142619436 -0.270515902 -0.037129466
## dimension_worst -0.13156024 -0.275702077 -0.229474476 -0.078971489
##
                      PC5 PC6 PC7
## id
                0.011327587 -0.316733438 0.9071156324 -0.096362415
## radius_mean -0.038129861 0.029588521 -0.0422987777 -0.116427419
                0.049091450 -0.031394323 0.0149935618 0.001875482
## texture_mean
## perimeter_mean
                -0.037715592 0.028394008 -0.0435888242 -0.106272097
## area mean
                -0.010562229 0.006113155 -0.0289256668 -0.047414568
               0.365750055 -0.262508993 -0.1403403617 -0.123541189
## smoothness_mean
## compactness mean -0.011786637 -0.004903894 -0.0453031106 0.043145968
## concavity_mean -0.086512506 -0.002356338 -0.0325530646 -0.102436021
                0.043667412 -0.034509273 -0.0814216298 -0.136923237
## points_mean
## symmetry_mean
               0.305378893  0.335082168  0.1182592361 -0.098874531
## compactness_se -0.280298048 0.066788120 0.02222220211 0.021001944
## concavity_se -0.354164595 0.049699104 0.0336810725 -0.219193299
## perimeter worst -0.007599144 0.012921166 -0.0145260986 0.002162488
## area worst 0.027413595 -0.024033338 -0.0007372602 0.066173186
## smoothness_worst 0.325860028 -0.365048687 -0.0670682168 -0.116496117
## compactness worst -0.121503371 0.034042714 0.0507556727 0.136509363
## concavity_worst -0.188280510 0.017962040 0.0352007117 -0.067085744
## points_worst -0.043123573 -0.029549100 -0.0207238959 -0.166500918
                ## symmetry_worst
## dimension worst -0.093699078 -0.092479698 0.0347167538 0.372034479
##
         PC9 FC10 --
0.149115642 -0.16926751 0.058188997 -0.006721252
                 PC9 PC10 PC11
## id
## radius_mean
                -0.046270835 -0.22402704 -0.079466081 -0.042213788
## perimeter_mean -0.036230738 -0.22634517 -0.069865929 -0.017573055
## area mean -0.080649856 -0.18600385 -0.062795372 -0.110760120
## smoothness mean 0.278996404 -0.06133822 0.084661549 0.135321954
## compactness mean 0.099214048 -0.19518602 0.005172841 0.307036205
## concavity_mean 0.075750464 0.03395563 0.134664686 -0.124553100
## dimension_mean 0.130639482 -0.15848117 -0.066456112 0.037318709
-0.030627892 0.24966405 0.071991560 -0.050731496
## area se
## compactness_se -0.148593714 -0.11518343 -0.038615749 0.206959272
-0.008571809 0.10669296 -0.038561250 -0.012490348
## texture_worst
## perimeter_worst -0.058854223 -0.09821693 -0.045750979 -0.051125158
## area_worst -0.097034650 -0.06179787 -0.068822329 -0.184460981
## smoothness worst -0.173257498 0.16912753 0.109278029 -0.142996001
## compactness_worst -0.111218083 -0.06445290 0.175401648 0.196805544
## concavity_worst -0.035467377 0.19661986 0.295581609 -0.184959562
                0.052322473 0.05121611 0.075496752 0.117518361
## points worst
## symmetry_worst -0.188266324 0.10308901 0.019223451 -0.157210098
## dimension_worst -0.087222442 -0.11291399 -0.007071634 -0.118625115
##
                     PC13 PC14 PC15 PC16
## id
               -0.004841084 -0.006500099 0.006885943 -0.002753492
## radius mean
                0.050603927 -0.012496988 -0.059054553 0.050789156
```

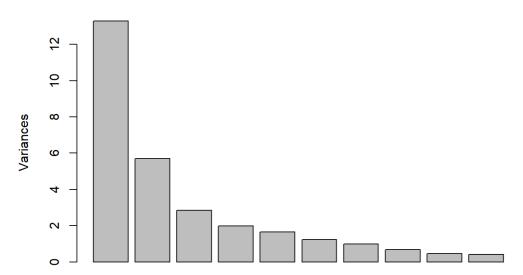
```
0.256273666 -0.201876125 0.020701124 0.108089530
## texture mean
## perimeter mean 0.038470392 -0.044684430 -0.048019221 0.039590476
## area_mean 0.065047550 -0.067879244 -0.010152279 -0.014636050
## compactness_mean -0.104264618 -0.230005458 -0.007661166 -0.230759682
## concavity_mean 0.065723393 -0.387349680 0.189733740 0.128386008
## points_mean
## symmetry_mean
                 0.042253113 -0.132637847 0.245219266 0.217299938
                -0.288054252 -0.189570545 -0.030903840 0.073950596
## smoothness se -0.293287983 -0.149148603 0.200139961 -0.018414232
## compactness se -0.263398426 -0.010320713 -0.491903153 -0.167886977
## concavity_se 0.251864823 -0.157777595 -0.135322845 -0.250292522
## points_se
                -0.006430584 0.494527095 0.199547389 -0.062548716
                0.319874237 -0.010836031 0.047340593 0.113219397
## symmetry_se
## dimension_se
## radius_worst
## texture_worst
                0.275943072 0.240767973 -0.145958050 0.353782637
                 0.080142089 0.080737140 -0.053897961 -0.100862417
## perimeter_worst -0.009084762 0.097004376 -0.012559001 -0.182407021
                0.047986766 0.101235629 0.006646192 -0.315142865
## area_worst
## smoothness_worst 0.056931408 0.206026671 -0.163389545 -0.045226715
## compactness_worst -0.371991007 -0.013117334 -0.165941776 0.049613607
## concavity_worst -0.086870368 -0.218055908 0.066854662 0.204743734
## points worst -0.068367254 0.254345228 0.276401728 0.169597618
## symmetry_worst 0.043937722 0.256766084 -0.005448734 -0.139913723
## dimension_worst -0.035134642 0.172524501 0.212520491 0.255448214
0.159152972 -0.034161758 0.040048687 0.029931705
## compactness_mean -0.170379447 -0.016302860 -0.014259132 0.292092522
## concavity_mean -0.270010606 -0.005071590 0.027973937 0.007197446
## dimension_mean 0.039119713 0.046298986 -0.086779501 -0.062879947
## radius_se -0.055118880 -0.124562479 0.231233991 0.181436577
## compactness_se -0.190065826 -0.015610691 -0.094108380 -0.250216687
## concavity_se
                 0.126034946 0.092345618 -0.005794297 0.119490304
                0.197671940 0.106747906 -0.046944796 -0.015851066
## points se
## points_se
## symmetry_se
                0.158541381 -0.279918359 -0.180195394 -0.084242460
## perimeter worst 0.056649279 -0.228493742 0.189279122 0.105666112
## area_worst 0.090325036 -0.286471546 0.158722686 -0.393681440
## smoothness worst -0.142781922 -0.276751162 -0.504565504 0.228506719
## compactness_worst 0.153347954 -0.003683424 0.073627229 0.025544372
## concavity_worst 0.216302398 -0.190307542 0.107894455 -0.035839305
##
                      PC21 PC22 PC23
                0.009870917 0.006195707 0.003190337 -0.010289027
## id
## id 0.009870917 0.006195707 0.003190337 -0.010289027
## radius_mean 0.046009507 0.070394387 -0.073021974 -0.098704322
## texture_mean 0.264801220 -0.436269565 -0.095890704 0.001311285
## perimeter_mean 0.015122205 0.070963404 -0.074821704 -0.040500943
## area mean 0.087345298 0.021672998 -0.097428804 0.009396470
## smoothness mean -0.023842011 0.117945821 -0.063741313 -0.020088204
## compactness mean -0.476395571 -0.213187888 0.094254664 0.058295270
## concavity_mean 0.037771062 -0.001270114 0.188862925 0.321062737
## points_mean
                0.231546040 0.017493297 0.313280824 -0.057974684
```

```
## radius_se
## texture_se
## perimeter_se
                0.090564458 -0.085660592 0.147793165 -0.263799753
               0.083589382 -0.212168357 -0.048761201 -0.001150858
0.169586632 0.317246026 -0.153859020 0.081384223
## area se
               -0.270679518 -0.207916141 -0.068745790 0.110258620
## smoothness_se
               -0.095370809 0.066602974 -0.051852247 -0.057154068
                0.451033960 0.159332265 0.048970757 0.003993806
## compactness_se
## perimeter_worst -0.010544107 0.094457678 -0.014952244 0.058698441
## compactness worst -0.220884131 -0.191897773 -0.033373706 -0.145389941
## dimension_worst 0.095366679 0.092769737 0.470358007 0.002775112
##
                PC25 PC26 PC27 PC28
               -0.004233388 -0.00132610 -0.002571324 -0.001623875
## id
## perimeter mean -0.117262178 -0.02743488 0.124670225 0.115650274
## area_mean 0.070557041 0.21057100 -0.361014547 -0.467489167
## smoothness_mean 0.068940049 -0.02876100 0.037372832 -0.069482805
## compactness mean -0.102198309 -0.39651346 -0.262695425 -0.098624638
## concavity_mean
                0.045550527 0.09717977 0.550227716 -0.363040016
## points_mean
## symmetry_mean
                0.082349955 0.18630114 -0.389316679 0.453345398
                0.018841491 0.02451053 0.015910368 0.015157593
## dimension_mean -0.134601525 0.20670502 0.096796804 0.101343150
## compactness se -0.122457873 -0.17364984 0.049404535 -0.046651501
## area worst
## smoothness worst -0.011224935 0.04787154 -0.012860335 0.040730207
## compactness_worst 0.185437121 0.62471727 0.100772153 0.071087434
## concavity worst -0.286701322 -0.11586768 -0.267236886 0.142148446
## points worst 0.105286798 -0.26352782 0.133749940 -0.230794105
## symmetry_worst -0.013193455 -0.04505357 -0.027824916 -0.022695808
## dimension_worst 0.037882167 -0.28015574 -0.004500884 -0.060081371
##
                 PC29 PC30
## id
               -1.891724e-05 -0.0006852263 -7.122581e-05
## radius_mean
## texture_mean
                2.111968e-01 -0.2114371011 -7.024325e-01
                -6.362507e-05 0.0106165839 -2.644366e-04
                8.434280e-02 -0.3838889617 6.898676e-01
## perimeter_mean
               -2.725167e-01 0.4227208085 3.297173e-02
## area_mean
## smoothness_mean 1.480038e-03 0.0034638648 4.850746e-03
## compactness_mean -5.466656e-03 0.0409079834 -4.468229e-02
## concavity_mean 4.554138e-02 0.0101122808 -2.512860e-02
## points mean
                -8.885707e-03 0.0041142627 1.067984e-03
## symmetry_mean
               1.432581e-03 0.0075571475 1.279594e-03
## dimension mean -6.312291e-03 -0.0073311823 4.751885e-03
## perimeter_se
                2.631905e-01 0.0060612569 -1.373310e-02
## compactness_se
                -1.539757e-02 -0.0032295613 -1.923037e-03
## concavity_se 5.819985e-03 -0.0161202167 8.921294e-03
## points_se -2.900497e-02 0.0241014722 2.178643e-03
## symmetry_se -7.637856e-03 0.0051771158 -3.338380e-04
```

```
"" OJIHHOCE J_OC
                     ,.00,0000 00
                    1.975791e-02 0.0083971145 -1.792802e-03
## dimension_se
                    4.126296e-01 0.6356796555 1.356846e-01
## radius_worst
                    -3.896988e-04 -0.0172219636 -1.020237e-03
## texture worst
## perimeter worst
                   -7.286790e-01 -0.0228830657 -7.974244e-02
                    2.389679e-01 -0.4448733182 -3.976788e-02
## area_worst
## smoothness_worst -1.535941e-03 -0.0074142082 -4.586820e-03
## compactness_worst 4.869512e-02 0.0001075081 1.285262e-02
## concavity_worst -1.764174e-02 0.0126547542 -4.031809e-04
                   2.247340e-02 -0.0353341030 2.276561e-03
## points_worst
## symmetry_worst
                    4.922100e-03 -0.0133523613 -3.910451e-04
## dimension worst
                   -2.356283e-02 -0.0115053741 -1.897779e-03
```

plot(breast_cancer_pca)

breast cancer pca



summary(breast_cancer_pca)

```
## Importance of components:
                                  PC2
##
                           PC1
                                         PC3
                                                 PC4
                                                         PC5
                        3.6453 2.3868 1.68386 1.40761 1.28406 1.11116
## Standard deviation
## Proportion of Variance 0.4286 0.1838 0.09146 0.06391 0.05319 0.03983
## Cumulative Proportion 0.4286 0.6124 0.70388 0.76779 0.82098 0.86081
##
                            PC7 PC8
                                          PC9 PC10
                                                       PC11
## Standard deviation
                       0.98908 0.81961 0.67882 0.6349 0.59089 0.54212
## Proportion of Variance 0.03156 0.02167 0.01486 0.0130 0.01126 0.00948
## Cumulative Proportion 0.89237 0.91404 0.92890 0.9419 0.95317 0.96265
##
                           PC13 PC14 PC15 PC16 PC17 PC18
## Standard deviation
                        0.51103 0.49125 0.39620 0.30680 0.28251 0.2430
## Proportion of Variance 0.00842 0.00778 0.00506 0.00304 0.00257 0.0019
## Cumulative Proportion 0.97107 0.97886 0.98392 0.98696 0.98953 0.9914
                                PC20 PC21
\# \#
                         PC19
                                              PC22
                                                       PC23
                       0.2293 0.22163 0.1763 0.17304 0.16562 0.15572
## Standard deviation
## Proportion of Variance 0.0017 0.00158 0.0010 0.00097 0.00088 0.00078
## Cumulative Proportion 0.9931 0.99472 0.9957 0.99669 0.99757 0.99835
                           PC25 PC26 PC27 PC28
                                                       PC29
## Standard deviation
                       0.13431 0.1244 0.09040 0.08305 0.03987 0.02736
## Proportion of Variance 0.00058 0.0005 0.00026 0.00022 0.00005 0.00002
## Cumulative Proportion 0.99893 0.9994 0.99970 0.99992 0.99997 1.00000
##
                           PC31
## Standard deviation
                        0.01153
## Proportion of Variance 0.00000
## Cumulative Proportion 1.00000
```

```
PC2
                              PC3
                                       PC4
          PC1
                                                 PC5
## [1,] 2.501946 -0.09694805 -0.4489597 2.3341176 0.69771548 -0.2430058
## [2,] 1.467439 -1.68630059 1.1542039 0.3362109 0.45962538 1.2308248
## [3,] 2.929028 -0.38319924 -0.8955891 -0.1164828 0.98441377 -0.2587872
## [4,] 1.995342 -1.33046592 1.1172876 2.0502761 0.25303846 -1.5539634
## [5,] 2.500252 2.01035097 -0.7584035 1.9862169 -1.13537096 0.5940361
## [6,] 2.018308 -0.78242095 0.1125197 -0.6532280 0.01841577 0.6914453
##
            PC7
                     PC8
                               PC9
                                        PC10
                                                 PC11
## [1,] 0.5092015 -1.11423307 0.2840243 0.32463197 -0.3245353 0.04981306
## [2,] 0.2937434 0.10000461 -0.0668399 0.42612180 0.4564029 1.19357566
## [5,] 0.1198201 -0.48279704 -0.2727816 -0.29439485 -0.3577533 0.03266208
## [6,] 0.1454026 0.06214539 0.2342454 0.73681239 -0.3671239 -0.77029743
##
            PC13
                    PC14
                                PC15
                                        PC16
## [2,] 0.01807424 -0.2824292 -0.204858888 -0.07067959 0.03088787
## [3,] 0.37435458 0.2585457 -0.330274216 -0.13000189 -0.24616091
## [4,] -0.90968379 0.2179117 -0.665825669 0.10213387 -0.10289446
## [5,] -0.35547138 -0.1480140 -0.005540503 -0.06495881 0.22273309
## [6,] -0.49542291 -0.2992431 0.049952835 -0.20161083 0.14920422
                      PC19
             PC18
                              PC20
## [1,] -0.104542766 -0.03484189 -0.09691187 -0.02846306 -0.00673628
## [2,] -0.405534243 -0.02886103 -0.05262226 -0.05987170 0.05868642
## [3,] 0.327711259 0.15937793 -0.13804895 -0.13489743 0.10080029
## [4,] 0.197085181 0.36251771 -0.40018239 -0.10302093 -0.28821708
## [5,] -0.129129156 -0.35877054 0.08515543 -0.08500541 -0.06332008
## [6,] -0.002229379 -0.08178568 0.18970936 -0.06872875 0.09669594
             PC23
                       PC24
                                   PC25
                                             PC26
## [2,] 0.070978613 -0.030822339 -0.016741580 0.04173030 -0.059332996
## [3,] 0.053909008 0.085484364 0.038277664 -0.04151896 -0.035546410
## [4,] 0.182045907 0.222848059 -0.115720065 -0.03676948 -0.148171674
## [5,] 0.043591030 0.008165322 0.002738052 0.05983731 0.046167735
##
            PC28 PC29 PC30
                                             PC31
## [1,] 0.05356131 0.015184882 0.015985406 0.001396101
## [3,] -0.07653067 -0.014640388 0.010307894 0.009074601
## [4,] -0.01711665 -0.047828494 0.023862995 0.000265075
## [5,] 0.03835364 0.032450800 -0.002312178 -0.002563269
## [6,] 0.00955434 -0.004403431 0.003869919 -0.002931194
# sample scores stored in breast_cancer_pca$x
# singular values (square roots of eigenvalues) stored in breast cancer pca$sdev
# loadings (eigenvectors) are stored in breast_cancer_pca$rotation
 variable means stored in breast cancer pca$center
# variable standard deviations stored in sparrows_pca$scale
# A table containing eigenvalues and %'s accounted, follows
# Eigenvalues are sdev^2
(eigen_breast_cancer <- breast_cancer_pca$sdev^2) ## brackets for print</pre>
## [1] 1.328806e+01 5.696805e+00 2.835395e+00 1.981357e+00 1.648815e+00
## [6] 1.234673e+00 9.782732e-01 6.717530e-01 4.607924e-01 4.031331e-01
## [11] 3.491550e-01 2.938904e-01 2.611469e-01 2.413302e-01 1.569736e-01
## [16] 9.412853e-02 7.980995e-02 5.904627e-02 5.259119e-02 4.912193e-02
## [21] 3.107078e-02 2.994121e-02 2.743052e-02 2.424902e-02 1.803936e-02
## [26] 1.547973e-02 8.171699e-03 6.898103e-03 1.589338e-03 7.483761e-04
## [31] 1.330402e-04
```

```
names(eigen_breast_cancer) <- paste("PC",1:31,sep="")
eigen_breast_cancer</pre>
```

```
PC1 PC2 PC3 PC4 PC5
## 1.328806e+01 5.696805e+00 2.835395e+00 1.981357e+00 1.648815e+00
  PC6 PC7 PC8 PC9 PC10
##
## 1.234673e+00 9.782732e-01 6.717530e-01 4.607924e-01 4.031331e-01
       PC11 PC12 PC13 PC14 PC15
##
## 3.491550e-01 2.938904e-01 2.611469e-01 2.413302e-01 1.569736e-01
       PC16 PC17 PC18 PC19
## 9.412853e-02 7.980995e-02 5.904627e-02 5.259119e-02 4.912193e-02
       PC21 PC22 PC23 PC24
## 3.107078e-02 2.994121e-02 2.743052e-02 2.424902e-02 1.803936e-02
       PC26 PC27 PC28 PC29 PC30
##
## 1.547973e-02 8.171699e-03 6.898103e-03 1.589338e-03 7.483761e-04
##
       PC31
## 1.330402e-04
sumlambdas <- sum(eigen breast cancer)</pre>
sumlambdas
```

```
propvar <- eigen_breast_cancer/sumlambdas</pre>
```

[1] 31

```
propvar
       PC1
                 PC2
                           PC3
                                     PC4
## 4.286470e-01 1.837679e-01 9.146436e-02 6.391475e-02 5.318759e-02
       PC6 PC7 PC8 PC9
##
                                             PC10
```

```
## 3.982815e-02 3.155720e-02 2.166945e-02 1.486427e-02 1.300429e-02
##
        PC11
                   PC12
                              PC13
                                         PC14
## 1.126306e-02 9.480337e-03 8.424094e-03 7.784846e-03 5.063666e-03
        PC16
                   PC17
                              PC18
                                         PC19
## 3.036404e-03 2.574514e-03 1.904718e-03 1.696490e-03 1.584578e-03
##
       PC21 PC22 PC23
                                    PC2.4
## 1.002283e-03 9.658453e-04 8.848556e-04 7.822265e-04 5.819149e-04
        PC26 PC27 PC28
                                    PC29
##
## 4.993461e-04 2.636032e-04 2.225194e-04 5.126895e-05 2.414116e-05
##
       PC31
```

summary(eigen breast cancer)

4.291620e-06

```
Min. 1st Qu. Median Mean 3rd Qu.
## 0.000133 0.025840 0.094129 1.000000 0.566273 13.288057
```

summary(breast_cancer_pca)

```
##
                          PC1
                                PC2
                                       PC3
                                                PC4
## Standard deviation
                      3.6453 2.3868 1.68386 1.40761 1.28406 1.11116
## Proportion of Variance 0.4286 0.1838 0.09146 0.06391 0.05319 0.03983
## Cumulative Proportion 0.4286 0.6124 0.70388 0.76779 0.82098 0.86081
##
                           PC7 PC8 PC9 PC10 PC11 PC12
## Standard deviation
                       0.98908 0.81961 0.67882 0.6349 0.59089 0.54212
## Proportion of Variance 0.03156 0.02167 0.01486 0.0130 0.01126 0.00948
## Cumulative Proportion 0.89237 0.91404 0.92890 0.9419 0.95317 0.96265
##
                         PC13 PC14 PC15 PC16 PC17 PC18
                       0.51103 0.49125 0.39620 0.30680 0.28251 0.2430
## Standard deviation
## Proportion of Variance 0.00842 0.00778 0.00506 0.00304 0.00257 0.0019
## Cumulative Proportion 0.97107 0.97886 0.98392 0.98696 0.98953 0.9914
##
                         PC19
                               PC20 PC21 PC22 PC23
## Standard deviation
                       0.2293 0.22163 0.1763 0.17304 0.16562 0.15572
## Proportion of Variance 0.0017 0.00158 0.0010 0.00097 0.00088 0.00078
## Cumulative Proportion 0.9931 0.99472 0.9957 0.99669 0.99757 0.99835
                         PC25 PC26 PC27 PC28 PC29 PC30
##
## Standard deviation
                      0.13431 0.1244 0.09040 0.08305 0.03987 0.02736
## Proportion of Variance 0.00058 0.0005 0.00026 0.00022 0.00005 0.00002
## Cumulative Proportion 0.99893 0.9994 0.99970 0.99992 0.99997 1.00000
##
                          PC31
## Standard deviation
                    0.01153
## Proportion of Variance 0.00000
## Cumulative Proportion 1.00000
cumvar breast cancer <- cumsum(propvar)</pre>
cumvar_breast_cancer
        PC1
               PC2
                        PC3
                                  PC4
                                            PC5
                                                      PC6
## 0.4286470 0.6124149 0.7038793 0.7677940 0.8209816 0.8608098 0.8923670
## PC8 PC9 PC10 PC11 PC12 PC13 PC14
## 0.9140364 0.9289007 0.9419050 0.9531681 0.9626484 0.9710725 0.9788573
##
      PC15 PC16 PC17 PC18 PC19 PC20 PC21
## 0.9839210 0.9869574 0.9895319 0.9914366 0.9931331 0.9947177 0.9957200
## PC22 PC23 PC24 PC25 PC26 PC27 PC28
## 0.9966858 0.9975707 0.9983529 0.9989348 0.9994342 0.9996978 0.9999203
## PC29 PC30 PC31
## 0.9999716 0.9999957 1.0000000
matlambdas <- rbind(eigen breast cancer,propvar,cumvar breast cancer)</pre>
rownames(matlambdas) <- c("Eigenvalues", "Prop. variance", "Cum. prop. variance")
round (matlambdas, 4)
##
                        PC1 PC2 PC3 PC4 PC5
                   13.2881 5.6968 2.8354 1.9814 1.6488 1.2347 0.9783
## Eigenvalues
## Prop. variance 0.4286 0.1838 0.0915 0.0639 0.0532 0.0398 0.0316
## Cum. prop. variance 0.4286 0.6124 0.7039 0.7678 0.8210 0.8608 0.8924
                      PC8 PC9 PC10 PC11 PC12 PC13 PC14
##
                   0.6718 0.4608 0.4031 0.3492 0.2939 0.2611 0.2413
## Eigenvalues
## Prop. variance 0.0217 0.0149 0.0130 0.0113 0.0095 0.0084 0.0078
## Cum. prop. variance 0.9140 0.9289 0.9419 0.9532 0.9626 0.9711 0.9789
                     PC15 PC16 PC17 PC18 PC19 PC20 PC21
##
## Eigenvalues
                    0.1570 0.0941 0.0798 0.0590 0.0526 0.0491 0.0311
## Prop. variance
                  0.0051 0.0030 0.0026 0.0019 0.0017 0.0016 0.0010
## Cum. prop. variance 0.9839 0.9870 0.9895 0.9914 0.9931 0.9947 0.9957
##
                      PC22 PC23 PC24 PC25 PC26 PC27 PC28
## Eigenvalues
                    0.0299 0.0274 0.0242 0.0180 0.0155 0.0082 0.0069
                   0.0010 0.0009 0.0008 0.0006 0.0005 0.0003 0.0002
## Prop. variance
## Cum. prop. variance 0.9967 0.9976 0.9984 0.9989 0.9994 0.9997 0.9999
##
                     PC29 PC30 PC31
## Eigenvalues
                    0.0016 7e-04 1e-04
## Prop. variance
                   0.0001 0e+00 0e+00
## Cum. prop. variance 1.0000 1e+00 1e+00
summary (breast cancer pca)
```

Importance of components:

```
## Importance of components:
##
                          PC1 PC2 PC3
                                               PC4
                                                      PC5
                     3.6453 2.3868 1.68386 1.40761 1.28406 1.11116
## Standard deviation
## Proportion of Variance 0.4286 0.1838 0.09146 0.06391 0.05319 0.03983
## Cumulative Proportion 0.4286 0.6124 0.70388 0.76779 0.82098 0.86081
##
                           PC7 PC8 PC9 PC10 PC11 PC12
                     0.98908 0.81961 0.67882 0.6349 0.59089 0.54212
## Standard deviation
## Proportion of Variance 0.03156 0.02167 0.01486 0.0130 0.01126 0.00948
## Cumulative Proportion 0.89237 0.91404 0.92890 0.9419 0.95317 0.96265
                        PC13 PC14 PC15 PC16 PC17 PC18
##
## Standard deviation 0.51103 0.49125 0.39620 0.30680 0.28251 0.2430
## Proportion of Variance 0.00842 0.00778 0.00506 0.00304 0.00257 0.0019
## Cumulative Proportion 0.97107 0.97886 0.98392 0.98696 0.98953 0.9914
                        PC19 PC20 PC21 PC22 PC23
## Standard deviation 0.2293 0.22163 0.1763 0.17304 0.16562 0.15572
## Proportion of Variance 0.0017 0.00158 0.0010 0.00097 0.00088 0.00078
## Cumulative Proportion 0.9931 0.99472 0.9957 0.99669 0.99757 0.99835
##
                         PC25 PC26 PC27 PC28 PC29 PC30
## Standard deviation 0.13431 0.1244 0.09040 0.08305 0.03987 0.02736
## Proportion of Variance 0.00058 0.0005 0.00026 0.00022 0.00005 0.00002
## Cumulative Proportion 0.99893 0.9994 0.99970 0.99992 0.99997 1.00000
##
                         PC31
## Standard deviation 0.01153
## Proportion of Variance 0.00000
## Cumulative Proportion 1.00000
```

breast cancer pca\$rotation

```
PC1
                              PC2
                                        PC3
## perimeter_mean -0.22753491 0.214589002 -0.012124791 0.042752797
## area_mean -0.22104577 0.230668816 0.026293150 0.054114724
## smoothness_mean -0.14241471 -0.186422211 -0.103182400 0.158098177
               -0.22104577 0.230668816 0.026293150 0.054114724
## compactness_mean -0.23906730 -0.152454726 -0.074768623 0.031818117
## concavity_mean -0.25828025 -0.060541625 0.001758736 0.019497124
## dimension_mean -0.06414779 -0.366531055 -0.020817875 0.047957856
## smoothness_se
               -0.01467821 -0.203189876 0.311787845 0.044368664
## compactness_se -0.17028840 -0.232503362 0.154557465 -0.026425360
## concavity_se
               -0.15354367 -0.196846081 0.176560052 0.002248291
## perimeter worst   -0.23663734   0.199295985   -0.050431945   0.014068572
## area worst -0.22493214 0.218985461 -0.013188891 0.025970672
## smoothness worst -0.12782441 -0.172562959 -0.255328751 0.014523359
## compactness_worst -0.20988456 -0.144253637 -0.234513609 -0.092562168
## concavity_worst -0.22860218 -0.098526524 -0.172024941 -0.074807188
## points_worst -0.25074620 0.007534367 -0.170480673 0.005305980
               -0.12267993 -0.142619436 -0.270515902 -0.037129466
## symmetry_worst
## dimension worst -0.13156024 -0.275702077 -0.229474476 -0.078971489
##
                      PC5 PC6 PC7
              0.011327587 -0.316733438 0.9071156324 -0.096362415
## id
## perimeter_mean
## area_mean -0.010562229 0.006113155 -0.0289256668 -0.047414568
## smoothness mean 0.365750055 -0.262508993 -0.1403403617 -0.123541189
## compactness mean -0.011786637 -0.004903894 -0.0453031106 0.043145968
## concavity_mean -0.086512506 -0.002356338 -0.0325530646 -0.102436021
```

```
## radius_se
           0.154254367 -0.023261199 0.0167882718 0.307415709
## texture_se
           0.190001500 0.022856912 -0.1902676469 -0.052632477
## concavity_se
           -0.195758558 -0.023197526 -0.0378517870 -0.370217167
## points_se
## perimeter worst -0.007599144 0.012921166 -0.0145260986 0.002162488
## area_worst 0.027413595 -0.024033338 -0.0007372602 0.066173186
## smoothness_worst 0.325860028 -0.365048687 -0.0670682168 -0.116496117
## compactness_worst -0.121503371 0.034042714 0.0507556727 0.136509363
## concavity_worst -0.188280510 0.017962040 0.0352007117 -0.067085744
## dimension worst -0.093699078 -0.092479698 0.0347167538 0.372034479
##
           PC9 PC10 PC11 PC12
          0.149115642 -0.16926751 0.058188997 -0.006721252
## id
## perimeter mean -0.036230738 -0.22634517 -0.069865929 -0.017573055
## area mean -0.080649856 -0.18600385 -0.062795372 -0.110760120
## concavity_mean 0.075750464 0.03395563 0.134664686 -0.124553100
           0.116569072 -0.14261678 0.006124860 0.071564686
## points_mean
## symmetry_mean
           ## compactness se -0.148593714 -0.11518343 -0.038615749 0.206959272
-0.058854223 -0.09821693 -0.045750979 -0.051125158
## perimeter worst
## area worst -0.097034650 -0.06179787 -0.068822329 -0.184460981
## smoothness_worst -0.173257498 0.16912753 0.109278029 -0.142996001
## compactness_worst -0.111218083 -0.06445290 0.175401648 0.196805544
## concavity_worst -0.035467377 0.19661986 0.295581609 -0.184959562
## points worst 0.052322473 0.05121611 0.075496752 0.117518361
## symmetry worst -0.188266324 0.10308901 0.019223451 -0.157210098
## dimension_worst -0.087222442 -0.11291399 -0.007071634 -0.118625115
##
            PC13 PC14 PC15 PC16
## id
           -0.004841084 -0.006500099 0.006885943 -0.002753492
## compactness mean -0.104264618 -0.230005458 -0.007661166 -0.230759682
## dimension mean 0.236120382 -0.106390748 0.377436108 -0.518333769
```

```
-U.UUU43UJO4 U.474JZ/U7J U.177J4/JO7 -U.UUZJ40/10
## Ъотиго ос
## symmetry_se
             0.319874237 -0.010836031 0.047340593 0.113219397
0.275943072 0.240767973 -0.145958050 0.353782637
## dimension se
## radius_worst
              ## perimeter_worst -0.009084762 0.097004376 -0.012559001 -0.182407021
## area worst 0.047986766 0.101235629 0.006646192 -0.315142865
## smoothness worst 0.056931408 0.206026671 -0.163389545 -0.045226715
## compactness_worst -0.371991007 -0.013117334 -0.165941776 0.049613607
## concavity_worst -0.086870368 -0.218055908 0.066854662 0.204743734
## points_worst -0.068367254 0.254345228 0.276401728 0.169597618
              0.043937722 0.256766084 -0.005448734 -0.139913723
## symmetry_worst
## dimension_worst -0.035134642 0.172524501 0.212520491 0.255448214
##
               PC17 PC18 PC19
       PC2.0
## id
## radius_mean
              0.150008977 0.209908003 -0.156773206 0.211821385
## texture_mean
              0.159152972 -0.034161758 0.040048687 0.029931705
## smoothness_mean 0.203117911 0.168171613 0.354463321 -0.160358262
## compactness_mean -0.170379447 -0.016302860 -0.014259132 0.292092522
## concavity mean -0.270010606 -0.005071590 0.027973937 0.007197446
## points_mean -0.381111880 0.028741889 0.087065594 -0.153991624
             0.165691481 -0.194702559 -0.169168737 -0.058503329
## symmetry_mean
              ## dimension_mean
## texture_worst
             -0.185972310 0.065992656 -0.057250572 -0.080880841
## perimeter_worst 0.056649279 -0.228493742 0.189279122 0.105666112
## area_worst
              0.090325036 -0.286471546 0.158722686 -0.393681440
## smoothness_worst -0.142781922 -0.276751162 -0.504565504 0.228506719
## compactness_worst 0.153347954 -0.003683424 0.073627229 0.025544372
## concavity_worst 0.216302398 -0.190307542 0.107894455 -0.035839305
## points_worst
              -0.178353485 -0.085180057 -0.067182996 -0.261323873
## dimension_worst 0.404957673 0.162920272 -0.026674889 -0.022516600
##
     PCZ1 FCZ2

0.009870917 0.006195707 0.003190337 -0.010289027
                  PC21 PC22 PC23 PC24
## id
## perimeter_mean 0.015122205 0.070963404 -0.074821704 -0.040500943
## area_mean 0.087345298 0.021672998 -0.097428804 0.009396470
## smoothness_mean -0.023842011 0.117945821 -0.063741313 -0.020088204
## compactness_mean -0.476395571 -0.213187888 0.094254664 0.058295270
## concavity_mean 0.037771062 -0.001270114 0.188862925 0.321062737
## smoothness se -0.095370809 0.066602974 -0.051852247 -0.057154068
## concavity_se
             -0.070203251 -0.071023842 0.200850815 -0.388573085
## area_worst
## smoothness_worst 0.140127867 -0.156936236 0.069660581 0.091134610
## compactness_worst -0.220884131 -0.191897773 -0.033373706 -0.145389941
```

```
## concavity_worst 0.047166544 0.139729448 -0.456817799 0.290302924
## points worst -0.039740929 -0.006870640 -0.305694162 -0.563297713
## symmetry worst 0.125617213 -0.155827542 -0.096426675 0.122996111
## dimension worst 0.095366679 0.092769737 0.470358007 0.002775112
               -0.117262178 -0.02743488 0.124670225 0.115650274
## perimeter mean
## compactness_mean -0.102198309 -0.39651346 -0.262695425 -0.098624638
## dimension mean -0.134601525 0.20670502 0.096796804 0.101343150
## concavity_se
                0.186159613 -0.01600952 -0.091931364 0.083824645
## points_se
## symmetry_se
               -0.107166573 0.12999049 0.018674110 0.011675700
               0.002613811 0.01936313 0.016991197 0.019891112
## area worst 0.146339946 0.03921251 -0.229813188 -0.237323945
## smoothness_worst -0.011224935 0.04787154 -0.012860335 0.040730207
## compactness_worst 0.185437121 0.62471727 0.100772153 0.071087434
## concavity_worst -0.286701322 -0.11586768 -0.267236886 0.142148446
## points_worst 0.105286798 -0.26352782 0.133749940 -0.230794105
               -0.013193455 -0.04505357 -0.027824916 -0.022695808
## symmetry worst
## dimension_worst 0.037882167 -0.28015574 -0.004500884 -0.060081371
##
                 PC29 PC30 PC31
               -1.891724e-05 -0.0006852263 -7.122581e-05
## id
## perimeter_mean 8.434280e-02 -0.3838889617 6.898676e-01
## area mean
               -2.725167e-01 0.4227208085 3.297173e-02
## smoothness mean 1.480038e-03 0.0034638648 4.850746e-03
## compactness_mean -5.466656e-03 0.0409079834 -4.468229e-02
## concavity mean 4.554138e-02 0.0101122808 -2.512860e-02
## points_mean
               -8.885707e-03 0.0041142627 1.067984e-03
               1.432581e-03 0.0075571475 1.279594e-03
## symmetry_mean
## dimension_mean
               -6.312291e-03 -0.0073311823 4.751885e-03
## radius_se
## texture_se
               -1.922290e-01 -0.1186768422 8.679321e-03
               -5.624974e-03 0.0086942153 1.063104e-03
## texture_se
## perimeter_se
               2.631905e-01 0.0060612569 -1.373310e-02
               -4.205668e-02 0.0863645419 -1.054698e-03
## area se
## smoothness_se
               9.795835e-03 -0.0016737982 1.618711e-03
## compactness_se -1.539757e-02 -0.0032295613 -1.923037e-03
## concavity_se
               5.819985e-03 -0.0161202167 8.921294e-03
## dimension_se
               1.975791e-02 0.0083971145 -1.792802e-03
                4.126296e-01 0.6356796555 1.356846e-01
## radius_worst
               -3.896988e-04 -0.0172219636 -1.020237e-03
## texture worst
## perimeter_worst -7.286790e-01 -0.0228830657 -7.974244e-02
                2.389679e-01 -0.4448733182 -3.976788e-02
## area worst
## smoothness worst -1.535941e-03 -0.0074142082 -4.586820e-03
## compactness_worst 4.869512e-02 0.0001075081 1.285262e-02
## concavity_worst -1.764174e-02 0.0126547542 -4.031809e-04
## points_worst 2.247340e-02 -0.0353341030 2.276561e-03
## symmetry_worst 4.922100e-03 -0.0133523613 -3.910451e-04
## dimension_worst -2.356283e-02 -0.0115053741 -1.897779e-03
```

```
## Stanuaru deviations (i, .., p-si).
  [1] 3.64527878 2.38679814 1.68386313 1.40760690 1.28406203 1.11115827
  [7] 0.98907696 0.81960537 0.67881693 0.63492763 0.59089337 0.54211662
## [13] 0.51102537 0.49125372 0.39619900 0.30680373 0.28250655 0.24299439
## [19] 0.22932770 0.22163467 0.17626907 0.17303527 0.16562163 0.15572098
## [25] 0.13431069 0.12441756 0.09039745 0.08305482 0.03986650 0.02735646
## [31] 0.01153431
## Rotation (n x k) = (31 \times 31):
                               PC2 PC3
##
                     PC1
               -0.02291216 0.034068491 0.096938436 -0.026598045
## id
## radius_mean
                -0.21891302 0.233271401 -0.011393786 0.042187950
## texture_mean
               -0.10384388 0.060044199 0.066892342 -0.602954308
## perimeter_mean -0.22753491 0.214589002 -0.012124791 0.042752797
## area_mean
                -0.22104577 0.230668816 0.026293150 0.054114724
## smoothness_mean -0.14241471 -0.186422211 -0.103182400 0.158098177
## compactness_mean -0.23906730 -0.152454726 -0.074768623 0.031818117
## concavity_mean -0.25828025 -0.060541625 0.001758736 0.019497124
## points_mean
               -0.26073811 0.034167392 -0.027579607 0.065785353
## dimension_mean -0.06414779 -0.366531055 -0.020817875 0.047957856
-0.18340675 -0.129965181 0.223850479 0.075252232
## perimeter_worst    -0.23663734    0.199295985    -0.050431945    0.014068572
## area worst -0.22493214 0.218985461 -0.013188891 0.025970672
## smoothness_worst -0.12782441 -0.172562959 -0.255328751 0.014523359
## compactness_worst -0.20988456 -0.144253637 -0.234513609 -0.092562168
## concavity_worst -0.22860218 -0.098526524 -0.172024941 -0.074807188
## points_worst -0.25074620 0.007534367 -0.170480673 0.005305980
## symmetry_worst
                -0.12267993 -0.142619436 -0.270515902 -0.037129466
## dimension_worst -0.13156024 -0.275702077 -0.229474476 -0.078971489
##
                       PC5 PC6 PC7
                0.011327587 -0.316733438 0.9071156324 -0.096362415
## id
## radius_mean
## texture_mean
               -0.038129861 0.029588521 -0.0422987777 -0.116427419
                0.049091450 -0.031394323 0.0149935618 0.001875482
## perimeter_mean -0.037715592 0.028394008 -0.0435888242 -0.106272097
## area mean -0.010562229 0.006113155 -0.0289256668 -0.047414568
## smoothness mean 0.365750055 -0.262508993 -0.1403403617 -0.123541189
## compactness mean -0.011786637 -0.004903894 -0.0453031106 0.043145968
## concavity_mean -0.086512506 -0.002356338 -0.0325530646 -0.102436021
## compactness_se -0.280298048 0.066788120 0.0222220211 0.021001944
## perimeter worst -0.007599144 0.012921166 -0.0145260986 0.002162488
## area worst
                0.027413595 -0.024033338 -0.0007372602 0.066173186
## smoothness_worst 0.325860028 -0.365048687 -0.0670682168 -0.116496117
## compactness_worst -0.121503371 0.034042714 0.0507556727 0.136509363
## concavity_worst -0.188280510 0.017962040 0.0352007117 -0.067085744
                -0.043123573 -0.029549100 -0.0207238959 -0.166500918
## points worst
## symmetry_worst 0.244245936 0.451404312 0.2340143294 -0.041439633
## dimension_worst -0.093699078 -0.092479698 0.0347167538 0.372034479
```

```
PC9
                             PC10
                                        PC11
##
               0.149115642 -0.16926751 0.058188997 -0.006721252
## id
               -0.046270835 -0.22402704 -0.079466081 -0.042213788
## radius mean
## texture mean -0.088727168 0.11945674 -0.253258091 0.304032359
## perimeter_mean -0.036230738 -0.22634517 -0.069865929 -0.017573055
## area mean -0.080649856 -0.18600385 -0.062795372 -0.110760120
## compactness_mean 0.099214048 -0.19518602 0.005172841 0.307036205
## concavity_mean 0.075750464 0.03395563 0.134664686 -0.124553100
## smoothness_se -0.580789293 -0.01015980 -0.179568831 -0.081753374
## concavity_se 0.034715098 0.36592141 0.113536362 -0.348342358
## points_se
## symmetry_se
## dimension_se
## radius_worst
## texture_worst
               -0.292785738 -0.22049558 0.328314881 0.185998712
               -0.060203202 -0.22637997 -0.353844543 -0.250428852
               -0.070224590 -0.09981025 -0.073013014 -0.105030701
               -0.008571809 0.10669296 -0.038561250 -0.012490348
## perimeter_worst -0.058854223 -0.09821693 -0.045750979 -0.051125158
               -0.097034650 -0.06179787 -0.068822329 -0.184460981
## area worst
## smoothness_worst -0.173257498 0.16912753 0.109278029 -0.142996001
## compactness_worst -0.111218083 -0.06445290 0.175401648 0.196805544
## concavity_worst -0.035467377 0.19661986 0.295581609 -0.184959562
## points worst 0.052322473 0.05121611 0.075496752 0.117518361
## symmetry_worst -0.188266324 0.10308901 0.019223451 -0.157210098
## dimension_worst -0.087222442 -0.11291399 -0.007071634 -0.118625115
PC13 PC14 PC15 PC16
               0.256273666 -0.201876125 0.020701124 0.108089530
              0.038470392 -0.044684430 -0.046013221 0...
0.065047550 -0.067879244 -0.010152279 -0.014636050
## perimeter_mean
## compactness_mean -0.104264618 -0.230005458 -0.007661166 -0.230759682
## area_se
## smoothness_se
               -0.017226446 -0.055687709 -0.083203050 0.045171638
               -0.293287983 -0.149148603 0.200139961 -0.018414232
## compactness_se -0.263398426 -0.010320713 -0.491903153 -0.167886977
## concavity_se
               0.251864823 -0.157777595 -0.135322845 -0.250292522
## points_se
## symmetry_se
               -0.006430584 0.494527095 0.199547389 -0.062548716
0.319874237 -0.010836031 0.047340593 0.113219397
## perimeter worst -0.009084762 0.097004376 -0.012559001 -0.182407021
## area_worst 0.047986766 0.101235629 0.006646192 -0.315142865
## smoothness_worst 0.056931408 0.206026671 -0.163389545 -0.045226715
## compactness_worst -0.371991007 -0.013117334 -0.165941776 0.049613607
## concavity_worst -0.086870368 -0.218055908 0.066854662 0.204743734
## points_worst -0.068367254 0.254345228 0.276401728 0.169597618
## symmetry_worst
               0.043937722 0.256766084 -0.005448734 -0.139913723
## dimension_worst -0.035134642 0.172524501 0.212520491 0.255448214
                     PC17 PC18 PC19 PC20
\# \#
## perimeter mean 0.113792993 0.201233658 -0.168413120 0.227079273
## area mean 0.130173978 0.251460456 -0.269145594 -0.045499625
## smoothness_mean 0.203117911 0.168171613 0.354463321 -0.160358262
## compactness mean -0.170379447 -0.016302860 -0.014259132 0.292092522
## concavity mean -0.270010606 -0.005071590 0.027973937 0.007197446
```

```
-0.381111880 0.028741889 0.087065594 -0.153991624
## points mean
              0.165691481 -0.194702559 -0.169168737 -0.058503329
## symmetry_mean
## dimension_mean
              0.039119713 0.046298986 -0.086779501 -0.062879947
## radius_se
              -0.055118880 -0.124562479 0.231233991 0.181436577
              0.032768777 0.041652813 0.009177450 0.038681291
## texture se
## perimeter_se
              -0.023929011 -0.009084130 0.014508488 0.364045783
## compactness_se -0.190065826 -0.015610691 -0.094108380 -0.250216687
## perimeter_worst 0.056649279 -0.228493742 0.189279122 0.105666112
## area_worst 0.090325036 -0.286471546 0.158722686 -0.393681440
## smoothness worst -0.142781922 -0.276751162 -0.504565504 0.228506719
## compactness_worst 0.153347954 -0.003683424 0.073627229 0.025544372
## concavity_worst 0.216302398 -0.190307542 0.107894455 -0.035839305
              -0.178353485 -0.085180057 -0.067182996 -0.261323873
## points worst
## dimension_worst 0.404957673 0.162920272 -0.026674889 -0.022516600
                   PC21 PC22 PC23 PC24
              0.009870917 0.006195707 0.003190337 -0.010289027
## id
0.087345298 0.021672998 -0.097428804 0.009396470
## area_mean
## smoothness_mean -0.023842011 0.117945821 -0.063741313 -0.020088204
## compactness mean -0.476395571 -0.213187888 0.094254664 0.058295270
## points_mean
## symmetry_mean
              -0.030776761 0.085067786 0.018331111 -0.052004767
## compactness se 0.451033960 0.159332265 0.048970757 0.003993806
## perimeter_worst -0.010544107 0.094457678 -0.014952244 0.058698441
## area_worst -0.053510824 -0.149328216 0.096798702 0.193293235
## smoothness worst 0.140127867 -0.156936236 0.069660581 0.091134610
## compactness worst -0.220884131 -0.191897773 -0.033373706 -0.145389941
## concavity worst 0.047166544 0.139729448 -0.456817799 0.290302924
## points worst -0.039740929 -0.006870640 -0.305694162 -0.563297713
## symmetry worst 0.125617213 -0.155827542 -0.096426675 0.122996111
## dimension worst 0.095366679 0.092769737 0.470358007 0.002775112
##
                PC25 PC26 PC27 PC28
## id
              -0.004233388 -0.00132610 -0.002571324 -0.001623875
## radius_mean
## texture_mean
              -0.183664583 0.01859418 0.128713229 0.131697326
              0.099441545 -0.08442059 0.024821224 0.017622634
              -0.117262178 -0.02743488 0.124670225 0.115650274
## perimeter mean
              0.070557041 0.21057100 -0.361014547 -0.467489167
## area_mean
## smoothness_mean 0.068940049 -0.02876100 0.037372832 -0.069482805
## compactness_mean -0.102198309 -0.39651346 -0.262695425 -0.098624638
## concavity_mean 0.045550527 0.09717977 0.550227716 -0.363040016
## dimension mean -0.134601525 0.20670502 0.096796804 0.101343150
0.023938591 -0.05709165 0.010893175 0.009925699
## texture_se
## perimeter_se
              0.516048248 -0.07217201 -0.103485879 -0.041989200
```

```
## CUMPACTIESS SE
                 -U.1224J/0/3 -U.1704264 U.0424U4JJJ -U.0400J1JU-
                  0.186159613 -0.01600952 -0.091931364 0.083824645
## concavity_se
                  -0.107166573 0.12999049 0.018674110 0.011675700
## points se
                 0.002613811 0.01936313 0.016991197 0.019891112
0.076177800 0.08458109 -0.035156906 0.012141785
## symmetry_se
## perimeter worst 0.236463109 -0.11790535 0.243266456 0.241658719
## area_worst 0.146339946 0.03921251 -0.229813188 -0.237323945
## smoothness worst -0.011224935 0.04787154 -0.012860335 0.040730207
## compactness_worst 0.185437121 0.62471727 0.100772153 0.071087434
## concavity_worst -0.286701322 -0.11586768 -0.267236886 0.142148446
## points_worst
                  0.105286798 -0.26352782 0.133749940 -0.230794105
## symmetry_worst
                  -0.013193455 -0.04505357 -0.027824916 -0.022695808
## dimension_worst 0.037882167 -0.28015574 -0.004500884 -0.060081371
##
                    PC29 PC30 PC31
                 -1.891724e-05 -0.0006852263 -7.122581e-05
## id
## radius_mean
                  2.111968e-01 -0.2114371011 -7.024325e-01
## texture_mean
                  -6.362507e-05 0.0106165839 -2.644366e-04
## perimeter_mean
                  8.434280e-02 -0.3838889617 6.898676e-01
## area mean
                 -2.725167e-01 0.4227208085 3.297173e-02
## smoothness mean 1.480038e-03 0.0034638648 4.850746e-03
## compactness mean -5.466656e-03 0.0409079834 -4.468229e-02
## concavity_mean 4.554138e-02 0.0101122808 -2.512860e-02
## points_mean
                  -8.885707e-03 0.0041142627 1.067984e-03
## symmetry_mean
                  1.432581e-03 0.0075571475 1.279594e-03
                  -6.312291e-03 -0.0073311823 4.751885e-03
## dimension mean
## radius_se
                  -1.922290e-01 -0.1186768422 8.679321e-03
                 -5.624974e-03 0.0086942153 1.063104e-03
2.631905e-01 0.0060612569 -1.373310e-02
## texture_se
## perimeter se
                  -4.205668e-02 0.0863645419 -1.054698e-03
## area_se
## smoothness_se
                  9.795835e-03 -0.0016737982 1.618711e-03
## compactness_se -1.539757e-02 -0.0032295613 -1.923037e-03
## concavity_se
                  5.819985e-03 -0.0161202167 8.921294e-03
                  -2.900497e-02 0.0241014722 2.178643e-03
## points se
## symmetry_se
                 -7.637856e-03 0.0051771158 -3.338380e-04
## dimension se
                  1.975791e-02 0.0083971145 -1.792802e-03
                  4.126296e-01 0.6356796555 1.356846e-01
## radius_worst
                  -3.896988e-04 -0.0172219636 -1.020237e-03
## texture_worst
## perimeter_worst -7.286790e-01 -0.0228830657 -7.974244e-02
## area worst
                   2.389679e-01 -0.4448733182 -3.976788e-02
## smoothness worst -1.535941e-03 -0.0074142082 -4.586820e-03
## compactness_worst 4.869512e-02 0.0001075081 1.285262e-02
## concavity_worst -1.764174e-02 0.0126547542 -4.031809e-04
                  2.247340e-02 -0.0353341030 2.276561e-03
## points_worst
## symmetry_worst
                  4.922100e-03 -0.0133523613 -3.910451e-04
## dimension worst -2.356283e-02 -0.0115053741 -1.897779e-03
```

Sample scores stored in breast_cancer_pca\$x
head(breast cancer pca\$x)

```
PC2 PC3
                                PC4
## [1,] 2.501946 -0.09694805 -0.4489597 2.3341176 0.69771548 -0.2430058
## [2,] 1.467439 -1.68630059 1.1542039 0.3362109 0.45962538 1.2308248
## [3,] 2.929028 -0.38319924 -0.8955891 -0.1164828 0.98441377 -0.2587872
## [4,] 1.995342 -1.33046592 1.1172876 2.0502761 0.25303846 -1.5539634
## [5,] 2.500252 2.01035097 -0.7584035 1.9862169 -1.13537096 0.5940361
## [6,] 2.018308 -0.78242095 0.1125197 -0.6532280 0.01841577 0.6914453
                 PC8 PC9
          PC7
                                    PC10
## [1,] 0.5092015 -1.11423307 0.2840243 0.32463197 -0.3245353 0.04981306
## [2,] 0.2937434 0.10000461 -0.0668399 0.42612180 0.4564029 1.19357566
## [3,] -0.3303385 0.03599041 0.8734350 0.02222192 0.4208602 -0.06687286
## [5,] 0.1198201 -0.48279704 -0.2727816 -0.29439485 -0.3577533 0.03266208
## [6,] 0.1454026 0.06214539 0.2342454 0.73681239 -0.3671239 -0.77029743
##
           PC13 PC14
                         PC15 PC16
## [2,] 0.01807424 -0.2824292 -0.204858888 -0.07067959 0.03088787
## [3,] 0.37435458 0.2585457 -0.330274216 -0.13000189 -0.24616091
## [4,] -0.90968379 0.2179117 -0.665825669 0.10213387 -0.10289446
## [5,] -0.35547138 -0.1480140 -0.005540503 -0.06495881 0.22273309
## [6,] -0.49542291 -0.2992431 0.049952835 -0.20161083 0.14920422
##
            PC18 PC19 PC20 PC21
## [1,] -0.104542766 -0.03484189 -0.09691187 -0.02846306 -0.00673628
## [2,] -0.405534243 -0.02886103 -0.05262226 -0.05987170 0.05868642
## [3,] 0.327711259 0.15937793 -0.13804895 -0.13489743 0.10080029
## [4,] 0.197085181 0.36251771 -0.40018239 -0.10302093 -0.28821708
## [5,] -0.129129156 -0.35877054 0.08515543 -0.08500541 -0.06332008
## [6,] -0.002229379 -0.08178568 0.18970936 -0.06872875 0.09669594
##
           PC23
                 PC24 PC25 PC26 PC27
## [2,] 0.070978613 -0.030822339 -0.016741580 0.04173030 -0.059332996
## [3,] 0.053909008 0.085484364 0.038277664 -0.04151896 -0.035546410
## [4,] 0.182045907 0.222848059 -0.115720065 -0.03676948 -0.148171674
## [5,] 0.043591030 0.008165322 0.002738052 0.05983731 0.046167735
##
          PC28 PC29 PC30 PC31
## [1,] 0.05356131 0.015184882 0.015985406 0.001396101
## [3,] -0.07653067 -0.014640388 0.010307894 0.009074601
## [4,] -0.01711665 -0.047828494 0.023862995 0.000265075
## [5,1
      0.03835364 0.032450800 -0.002312178 -0.002563269
## [6,] 0.00955434 -0.004403431 0.003869919 -0.002931194
```

```
# Identifying the scores by their diagnosis
diag_pca <- cbind(data.frame(diagnosis),breast_cancer_pca$x)
head(diag_pca)</pre>
```

```
diagnosis PC1
                     PC2 PC3
       B 2.501946 -0.09694805 -0.4489597 2.3341176 0.69771548
## 1
          B 1.467439 -1.68630059 1.1542039 0.3362109 0.45962538
## 2
          B 2.929028 -0.38319924 -0.8955891 -0.1164828 0.98441377
## 3
          B 1.995342 -1.33046592 1.1172876 2.0502761 0.25303846
## 4
## 5
          B 2.500252 2.01035097 -0.7584035 1.9862169 -1.13537096
          B 2.018308 -0.78242095 0.1125197 -0.6532280 0.01841577
          PC6 PC7 PC8 PC9 PC10
##
## 1 -0.2430058 0.5092015 -1.11423307 0.2840243 0.32463197 -0.3245353
## 2 1.2308248 0.2937434 0.10000461 -0.0668399 0.42612180 0.4564029
## 3 -0.2587872 -0.3303385 0.03599041 0.8734350 0.02222192 0.4208602
## 4 -1.5539634 -0.9692185 -1.31852134 0.6254396 0.05666470 -0.0691646
## 5 0.5940361 0.1198201 -0.48279704 -0.2727816 -0.29439485 -0.3577533
## 6 0.6914453 0.1454026 0.06214539 0.2342454 0.73681239 -0.3671239
##
     PC12 PC13 PC14 PC15
                                               PC16
## 1 0.04981306 -0.19760220 0.1134403 -0.059302558 0.16637723 -0.04286656
## 2 1.19357566 0.01807424 -0.2824292 -0.204858888 -0.07067959 0.03088787
## 3 -0.06687286 0.37435458 0.2585457 -0.330274216 -0.13000189 -0.24616091
## 4 0.97082409 -0.90968379 0.2179117 -0.665825669 0.10213387 -0.10289446
## 5 0.03266208 -0.35547138 -0.1480140 -0.005540503 -0.06495881 0.22273309
## 6 -0.77029743 -0.49542291 -0.2992431 0.049952835 -0.20161083 0.14920422
\# \#
          PC18
                   PC19 PC20 PC21 PC22
## 1 -0.104542766 -0.03484189 -0.09691187 -0.02846306 -0.00673628
## 2 -0.405534243 -0.02886103 -0.05262226 -0.05987170 0.05868642
## 3 0.327711259 0.15937793 -0.13804895 -0.13489743 0.10080029
## 4 0.197085181 0.36251771 -0.40018239 -0.10302093 -0.28821708
## 5 -0.129129156 -0.35877054 0.08515543 -0.08500541 -0.06332008
## 6 -0.002229379 -0.08178568 0.18970936 -0.06872875 0.09669594
##
           PC23 PC24 PC25 PC26 PC27
## 1 -0.038971937 0.062212075 0.088438866 0.04872948 -0.007000724
## 2 0.070978613 -0.030822339 -0.016741580 0.04173030 -0.059332996
## 3 0.053909008 0.085484364 0.038277664 -0.04151896 -0.035546410
## 4 0.182045907 0.222848059 -0.115720065 -0.03676948 -0.148171674
## 5 0.043591030 0.008165322 0.002738052 0.05983731 0.046167735
## 6 -0.001458054 -0.031338348 0.042784223 -0.08646068 -0.030944690
##
        PC28 PC29 PC30 PC31
## 1 0.05356131 0.015184882 0.015985406 0.001396101
## 3 -0.07653067 -0.014640388 0.010307894 0.009074601
## 4 -0.01711665 -0.047828494 0.023862995 0.000265075
    0.03835364 0.032450800 -0.002312178 -0.002563269
## 6 0.00955434 -0.004403431 0.003869919 -0.002931194
```

```
# Means of scores for all the PC's classified by diagnosis status
tabmeansPC <- aggregate(diag_pca[,2:31],by=list(diagnosis=breast_cancer$diagnosis),mean)
tabmeansPC</pre>
```

```
## diagnosis
                PC1
                         PC2
                                   PC3
                                            PC4
    B 2.204253 -0.3436398 0.2160542 0.1384470 -0.09800974
## 2
          M -3.711879 0.5786765 -0.3638272 -0.2331395 0.16504470
##
          PC.6
                     PC7
                              PC8
                                        PC9
## 1 0.004373132 0.01691799 0.04905754 0.03377092 -0.01002704
## 2 -0.007364189 -0.02848926 -0.08261104 -0.05686895 0.01688516
##
                   PC12 PC13 PC14
          PC11
## 1 -0.0007888591 0.006017621 0.003305282 -0.03736471 -0.02453152
## 2 0.0013284090 -0.010133446 -0.005565970 0.06292076 0.04131016
##
         PC16
                    PC17 PC18 PC19
## 1 -0.02516699 0.0001112917 0.006103777 -0.01307695 -0.009733891
## 2 0.04238026 -0.0001874111 -0.010278530 0.02202108 0.016391505
##
          PC21 PC22 PC23
                                           PC24
## 1 -0.008359794 -0.006225063 -0.003024993 0.002609428 0.007813479
## 2 0.014077577 0.010482771 0.005093974 -0.004394179 -0.013157604
##
               PC27
                          PC28 PC29 PC30
          PC26
## 1 -0.000879209 -0.003967725 -0.001909259 -0.0003418423 -0.0009838392
## 2 0.001480555 0.006681499 0.003215121 0.0005756496 0.0016567480
```

```
## diagnosis PC1 PC2 PC3 PC4
## 2 M -3.711879 0.5786765 -0.3638272 -0.2331395 0.16504470
         B 2.204253 -0.3436398 0.2160542 0.1384470 -0.09800974
## 1
        PC6 PC7 PC8
##
                                  PC9
## 2 -0.007364189 -0.02848926 -0.08261104 -0.05686895 0.01688516
## 1 0.004373132 0.01691799 0.04905754 0.03377092 -0.01002704
                PC12
                         PC13 PC14
## 2 0.0013284090 -0.010133446 -0.005565970 0.06292076 0.04131016
## 1 -0.0007888591 0.006017621 0.003305282 -0.03736471 -0.02453152
##
       PC16 PC17 PC18 PC19
## 2 0.04238026 -0.0001874111 -0.010278530 0.02202108 0.016391505
##
        PC21
             PC22
                          PC23
                                    PC24
## 2 0.014077577 0.010482771 0.005093974 -0.004394179 -0.013157604
##
       PC26
             PC27
                      PC28
                                PC29
## 2 0.001480555 0.006681499 0.003215121 0.0005756496 0.0016567480
## 1 -0.000879209 -0.003967725 -0.001909259 -0.0003418423 -0.0009838392
```

```
tabfmeans <- t(tabmeansPC[,-1])
tabfmeans</pre>
```

```
## PC1 -3.7118786952 2.2042528946
## PC2 0.5786764540 -0.3436397990
## PC3 -0.3638271826 0.2160542373
## PC4 -0.2331394896 0.1384469798
## PC5 0.1650447018 -0.0980097389
## PC6 -0.0073641886 0.0043731316
## PC7 -0.0284892608 0.0169179924
## PC8 -0.0826110415 0.0490575373
## PC9 -0.0568689505 0.0337709174
## PC10 0.0168851623 -0.0100270432
## PC11 0.0013284090 -0.0007888591
## PC12 -0.0101334459 0.0060176205
## PC13 -0.0055659702 0.0033052820
## PC14 0.0629207582 -0.0373647080
## PC15 0.0413101623 -0.0245315249
## PC16 0.0423802589 -0.0251669885
## PC17 -0.0001874111 0.0001112917
## PC18 -0.0102785304 0.0061037771
## PC19 0.0220210837 -0.0130769461
## PC20 0.0163915046 -0.0097338907
## PC21 0.0140775772 -0.0083597937
## PC22 0.0104827709 -0.0062250628
## PC23 0.0050939739 -0.0030249929
## PC24 -0.0043941787 0.0026094282
## PC25 -0.0131576035 0.0078134789
## PC26 0.0014805547 -0.0008792090
## PC27 0.0066814986 -0.0039677247
## PC28 0.0032151208 -0.0019092594
## PC29 0.0005756496 -0.0003418423
## PC30 0.0016567480 -0.0009838392
```

```
colnames(tabfmeans) <- t(as.vector(tabmeansPC[1]))
tabfmeans</pre>
```

```
##
## PC1
       -3.7118786952 2.2042528946
## PC2
        0.5786764540 -0.3436397990
## PC3 -0.3638271826 0.2160542373
       -0.2331394896 0.1384469798
## PC4
## PC5
       0.1650447018 -0.0980097389
## PC6 -0.0073641886 0.0043731316
## PC7 -0.0284892608 0.0169179924
## PC8 -0.0826110415 0.0490575373
## PC9 -0.0568689505 0.0337709174
## PC10 0.0168851623 -0.0100270432
## PC11 0.0013284090 -0.0007888591
## PC12 -0.0101334459 0.0060176205
## PC13 -0.0055659702 0.0033052820
## PC14 0.0629207582 -0.0373647080
## PC15
        0.0413101623 -0.0245315249
## PC16 0.0423802589 -0.0251669885
## PC17 -0.0001874111 0.0001112917
## PC18 -0.0102785304 0.0061037771
## PC19 0.0220210837 -0.0130769461
## PC20 0.0163915046 -0.0097338907
## PC21 0.0140775772 -0.0083597937
## PC22 0.0104827709 -0.0062250628
## PC23 0.0050939739 -0.0030249929
## PC24 -0.0043941787 0.0026094282
## PC25 -0.0131576035 0.0078134789
## PC26 0.0014805547 -0.0008792090
        0.0066814986 -0.0039677247
## PC27
## PC28 0.0032151208 -0.0019092594
## PC29 0.0005756496 -0.0003418423
## PC30 0.0016567480 -0.0009838392
```

```
# Standard deviations of scores for all the PC's classified by diagnosis status
tabsdsPC <- aggregate(diag_pca[,2:31],by=list(breast_cancer$diagnosis),sd)
tabfsds <- t(tabsdsPC[,-1])
colnames(tabfsds) <- t(as.vector(tabsdsPC[1]))
tabfsds</pre>
```

```
##
## PC1 1.63956487 3.02839244
## PC2 2.08691418 2.72966952
## PC3 1.36038677 2.07323861
## PC4 1.39613539 1.39927526
## PC5 1.39777817 1.04807636
## PC6 0.95340249 1.33789147
## PC7
       0.95140466 1.05116532
## PC8 0.64932787 1.04212370
## PC9 0.65881989 0.70917409
## PC10 0.57998076 0.71917808
## PC11 0.60269891 0.57186757
## PC12 0.54006502 0.54668527
## PC13 0.51222488 0.51016086
## PC14 0.37926601 0.63294179
## PC15 0.38070917 0.41867516
## PC16 0.26760334 0.36012377
## PC17 0.26627925 0.30856638
## PC18 0.17881651 0.32377631
## PC19 0.17193548 0.30152802
## PC20 0.17202337 0.28613872
## PC21 0.15015083 0.21280261
## PC22 0.15982321 0.19320204
## PC23 0.14579937 0.19476702
## PC24 0.12247334 0.19984615
## PC25 0.11050781 0.16637188
## PC26 0.09806381 0.15950557
## PC27 0.07202620 0.11477556
## PC28 0.06617618 0.10565982
## PC29 0.02588741 0.05609976
## PC30 0.01834069 0.03798219
```

```
t.test(PC1~breast_cancer$diagnosis,data=diag_pca)
```

```
##
## Welch Two Sample t-test
##
## data: PC1 by breast_cancer$diagnosis
## t = 26.251, df = 285.72, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 5.472542 6.359721
## sample estimates:
## mean in group B mean in group M
## 2.204253 -3.711879</pre>
```

t.test(PC2~breast cancer\$diagnosis,data=diag pca)

```
##
## Welch Two Sample t-test
##
## data: PC2 by breast_cancer$diagnosis
## t = -4.2387, df = 357.38, p-value = 2.865e-05
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.3502373 -0.4943952
## sample estimates:
## mean in group B mean in group M
## -0.3436398     0.5786765
```

t.test(PC3~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC3 by breast_cancer$diagnosis
## t = 3.6343, df = 320.28, p-value = 0.0003246
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.2659658 0.8937970
## sample estimates:
## mean in group B mean in group M
## 0.2160542 -0.3638272
```

t.test(PC4~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC4 by breast_cancer$diagnosis
## t = 3.0652, df = 442.55, p-value = 0.002308
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.1333371 0.6098358
## sample estimates:
## mean in group B mean in group M
## 0.1384470 -0.2331395
```

t.test(PC5~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC5 by breast_cancer$diagnosis
## t = -2.5485, df = 537.03, p-value = 0.0111
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.46581747 -0.06029141
## sample estimates:
## mean in group B mean in group M
## -0.09800974     0.16504470
```

t.test(PC6~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC6 by breast_cancer$diagnosis
## t = 0.11197, df = 339.17, p-value = 0.9109
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.1944615  0.2179362
## sample estimates:
## mean in group B mean in group M
## 0.004373132  -0.007364189
```

t.test(PC7~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC7 by breast_cancer$diagnosis
## t = 0.51587, df = 408.87, p-value = 0.6062
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.1276209 0.2184354
## sample estimates:
## mean in group B mean in group M
## 0.01691799 -0.02848926
```

t.test(PC8~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC8 by breast_cancer$diagnosis
## t = 1.6584, df = 309.75, p-value = 0.09825
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02455572 0.28789287
## sample estimates:
## mean in group B mean in group M
## 0.04905754 -0.08261104
```

t.test(PC9~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC9 by breast_cancer$diagnosis
## t = 1.5132, df = 417.67, p-value = 0.131
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.0271048 0.2083845
## sample estimates:
## mean in group B mean in group M
## 0.03377092 -0.05686895
```

t.test(PC10~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC10 by breast_cancer$diagnosis
## t = -0.46277, df = 372.52, p-value = 0.6438
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.14126432  0.08743991
## sample estimates:
## mean in group B mean in group M
## -0.01002704  0.01688516
```

t.test(PC11~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC11 by breast_cancer$diagnosis
## t = -0.041845, df = 462.01, p-value = 0.9666
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.10154692 0.09731238
## sample estimates:
## mean in group B mean in group M
## -0.0007888591 0.0013284090
```

t.test(PC12~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC12 by breast_cancer$diagnosis
## t = 0.34227, df = 439.04, p-value = 0.7323
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.07659212 0.10889425
## sample estimates:
## mean in group B mean in group M
## 0.006017621 -0.010133446
```

t.test(PC13~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC13 by breast_cancer$diagnosis
## t = 0.20025, df = 444.77, p-value = 0.8414
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.07819457 0.09593708
## sample estimates:
## mean in group B mean in group M
## 0.003305282 -0.005565970
```

t.test(PC14~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC14 by breast_cancer$diagnosis
## t = -2.0945, df = 302.42, p-value = 0.03705
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.194508096 -0.006062836
## sample estimates:
## mean in group B mean in group M
## -0.03736471  0.06292076
```

t.test(PC15~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC15 by breast_cancer$diagnosis
## t = -1.8752, df = 410.43, p-value = 0.06147
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.134862771 0.003179396
## sample estimates:
## mean in group B mean in group M
## -0.02453152 0.04131016
```

t.test(PC16~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC16 by breast_cancer$diagnosis
## t = -2.37, df = 349.77, p-value = 0.01833
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.12360307 -0.01149143
## sample estimates:
## mean in group B mean in group M
## -0.02516699    0.04238026
```

t.test(PC17~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC17 by breast_cancer$diagnosis
## t = 0.011737, df = 393.3, p-value = 0.9906
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.04973747 0.05033487
## sample estimates:
## mean in group B mean in group M
## 0.0001112917 -0.0001874111
```

t.test(PC18~breast_cancer\$diagnosis,data=diag_pca)

t.test(PC19~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC19 by breast_cancer$diagnosis
## t = -1.5516, df = 293.85, p-value = 0.1218
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.079616015 0.009419955
## sample estimates:
## mean in group B mean in group M
## -0.01307695 0.02202108
```

t.test(PC20~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC20 by breast_cancer$diagnosis
## t = -1.2062, df = 303.02, p-value = 0.2287
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.06874572 0.01649493
## sample estimates:
## mean in group B mean in group M
## -0.009733891 0.016391505
```

t.test(PC21~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC21 by breast_cancer$diagnosis
## t = -1.3487, df = 336.76, p-value = 0.1783
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.05516116  0.01028642
## sample estimates:
## mean in group B mean in group M
## -0.008359794  0.014077577
```

t.test(PC22~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC22 by breast_cancer$diagnosis
## t = -1.0618, df = 380.13, p-value = 0.289
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.04764831 0.01423264
## sample estimates:
## mean in group B mean in group M
## -0.006225063 0.010482771
```

t.test(PC23~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC23 by breast_cancer$diagnosis
## t = -0.52575, df = 351.71, p-value = 0.5994
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03849078 0.02225285
## sample estimates:
## mean in group B mean in group M
## -0.003024993 0.005093974
```

t.test(PC24~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC24 by breast_cancer$diagnosis
## t = 0.4614, df = 306.57, p-value = 0.6448
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02286480    0.03687201
## sample estimates:
## mean in group B mean in group M
##    0.002609428    -0.004394179
```

t.test(PC25~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC25 by breast_cancer$diagnosis
## t = 1.6337, df = 322.91, p-value = 0.1033
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.004282315  0.046224480
## sample estimates:
## mean in group B mean in group M
## 0.007813479  -0.013157604
```

t.test(PC26~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC26 by breast_cancer$diagnosis
## t = -0.19467, df = 307.18, p-value = 0.8458
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02621273  0.02149320
## sample estimates:
## mean in group B mean in group M
## -0.000879209  0.001480555
```

t.test(PC27~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC27 by breast_cancer$diagnosis
## t = -1.2162, df = 311.14, p-value = 0.2248
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.027878004 0.006579557
## sample estimates:
## mean in group B mean in group M
## -0.003967725 0.006681499
```

t.test(PC28~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC28 by breast_cancer$diagnosis
## t = -0.63596, df = 310.76, p-value = 0.5253
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02097902 0.01073026
## sample estimates:
## mean in group B mean in group M
## -0.001909259 0.003215121
```

t.test(PC29~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC29 by breast_cancer$diagnosis
## t = -0.22436, df = 265.22, p-value = 0.8226
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.008969119 0.007134135
## sample estimates:
## mean in group B mean in group M
## -0.0003418423 0.0005756496
```

t.test(PC30~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC30 by breast_cancer$diagnosis
## t = -0.9487, df = 270.4, p-value = 0.3436
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.008120430  0.002839256
## sample estimates:
## mean in group B mean in group M
## -0.0009838392  0.0016567480
```

t.test(PC31~breast_cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC31 by breast_cancer$diagnosis
## t = -0.54256, df = 278.74, p-value = 0.5879
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.002917016 0.001656473
## sample estimates:
## mean in group B mean in group M
## -0.0002348289 0.0003954429
```

F ratio tests
var.test(PC1~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC1 by breast_cancer$diagnosis
## F = 0.29311, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2293890 0.3717204
## sample estimates:
## ratio of variances
## ratio of variances
## 0.2931115</pre>
```

var.test(PC2~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC2 by breast_cancer$diagnosis
## F = 0.58451, num df = 356, denom df = 211, p-value = 8.474e-06
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.4574343 0.7412633
## sample estimates:
## ratio of variances
## ratio of variances
## 0.5845061
```

var.test(PC3~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC3 by breast_cancer$diagnosis
## F = 0.43055, num df = 356, denom df = 211, p-value = 2.327e-12
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3369504 0.5460214
## sample estimates:
## ratio of variances
## 0.4305526
```

var.test(PC4~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC4 by breast_cancer$diagnosis
## F = 0.99552, num df = 356, denom df = 211, p-value = 0.9625
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.7790915 1.2625024
## sample estimates:
## ratio of variances
## 0.9955172
```

var.test(PC5~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC5 by breast_cancer$diagnosis
## F = 1.7787, num df = 356, denom df = 211, p-value = 5.82e-06
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 1.391972 2.255662
## sample estimates:
## ratio of variances
## 1.778651
```

var.test(PC6~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC6 by breast_cancer$diagnosis
## F = 0.50782, num df = 356, denom df = 211, p-value = 1.795e-08
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3974207 0.6440124
## sample estimates:
## ratio of variances
## ratio of variances
## 0.5078212
```

var.test(PC7~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC7 by breast_cancer$diagnosis
## F = 0.8192, num df = 356, denom df = 211, p-value = 0.1
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.6411036 1.0388957
## sample estimates:
## ratio of variances
## 0.8191973
```

var.test(PC8~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC8 by breast_cancer$diagnosis
## F = 0.38823, num df = 356, denom df = 211, p-value = 3.319e-15
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3038290 0.4923489
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3882304
```

var.test(PC9~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC9 by breast_cancer$diagnosis
## F = 0.86303, num df = 356, denom df = 211, p-value = 0.2243
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.6754099 1.0944883
## sample estimates:
## ratio of variances
## 0.8630336
```

var.test(PC10~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC10 by breast_cancer$diagnosis
## F = 0.65036, num df = 356, denom df = 211, p-value = 0.0003698
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.5089722 0.8247793
## sample estimates:
## ratio of variances
## ratio of variances
## 0.6503607
```

var.test(PC11~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC11 by breast_cancer$diagnosis
## F = 1.1107, num df = 356, denom df = 211, p-value = 0.4012
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.8692598 1.4086183
## sample estimates:
## ratio of variances
## ratio of variances
```

var.test(PC12~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC12 by breast_cancer$diagnosis
## F = 0.97593, num df = 356, denom df = 211, p-value = 0.8346
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.7637603 1.2376584
## sample estimates:
## ratio of variances
## ratio of variances
## 0.975927
```

var.test(PC13~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC13 by breast_cancer$diagnosis
## F = 1.0081, num df = 356, denom df = 211, p-value = 0.956
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.7889451 1.2784699
## sample estimates:
## ratio of variances
## ratio of variances
## 1.008108
```

var.test(PC14~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC14 by breast_cancer$diagnosis
## F = 0.35905, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2809959 0.4553483
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3590544</pre>
```

var.test(PC15~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC15 by breast_cancer$diagnosis
## F = 0.82686, num df = 356, denom df = 211, p-value = 0.1169
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.6471009 1.0486142
## sample estimates:
## ratio of variances
## ratio of variances
## 0.8268605
```

var.test(PC16~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC16 by breast_cancer$diagnosis
## F = 0.55218, num df = 356, denom df = 211, p-value = 8.26e-07
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.4321348 0.7002658
## sample estimates:
## ratio of variances
## ratio of variances
## 0.5521785
```

var.test(PC17~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC17 by breast_cancer$diagnosis
## F = 0.74469, num df = 356, denom df = 211, p-value = 0.01494
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.5827968 0.9444106
## sample estimates:
## ratio of variances
## 0.7446933
```

var.test(PC18~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC18 by breast_cancer$diagnosis
## F = 0.30502, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2387068 0.3868197
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3050177</pre>
```

var.test(PC19~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC19 by breast_cancer$diagnosis
## F = 0.32514, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2544576 0.4123434
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3251439</pre>
```

var.test(PC20~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC20 by breast_cancer$diagnosis
## F = 0.36143, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2828534 0.4583583
## sample estimates:
## ratio of variances
## 0.3614279</pre>
```

var.test(PC21~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC21 by breast_cancer$diagnosis
## F = 0.49785, num df = 356, denom df = 211, p-value = 6.758e-09
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3896201 0.6313716
## sample estimates:
## ratio of variances
## 0.4978535
```

var.test(PC22~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC22 by breast_cancer$diagnosis
## F = 0.68432, num df = 356, denom df = 211, p-value = 0.001709
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.5355451 0.8678402
## sample estimates:
## ratio of variances
## ratio of variances
## 0.6843154
```

var.test(PC23~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC23 by breast_cancer$diagnosis
## F = 0.56038, num df = 356, denom df = 211, p-value = 1.542e-06
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.4385511 0.7106634
## sample estimates:
## ratio of variances
## ratio of variances
## 0.5603772
```

var.test(PC24~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC24 by breast_cancer$diagnosis
## F = 0.37557, num df = 356, denom df = 211, p-value = 3.495e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2939215 0.4762939
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3755706
```

var.test(PC25~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC25 by breast_cancer$diagnosis
## F = 0.44119, num df = 356, denom df = 211, p-value = 9.824e-12
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3452756 0.5595122
## sample estimates:
## ratio of variances
## 0.4411905
```

var.test(PC26~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC26 by breast_cancer$diagnosis
## F = 0.37798, num df = 356, denom df = 211, p-value = 5.423e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2958050 0.4793461
## sample estimates:
## ratio of variances
## 0.3779774
```

var.test(PC27~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC27 by breast_cancer$diagnosis
## F = 0.39381, num df = 356, denom df = 211, p-value = 8.544e-15
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3081924 0.4994196
## sample estimates:
## ratio of variances
## ratio of variances
## 0.3938058
```

var.test(PC28~breast_cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC28 by breast_cancer$diagnosis
## F = 0.39227, num df = 356, denom df = 211, p-value = 6.601e-15
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.3069894 0.4974701
## sample estimates:
## ratio of variances
## 0.3922686
```

var.test(PC29~breast_cancer\$diagnosis,data=diag_pca)

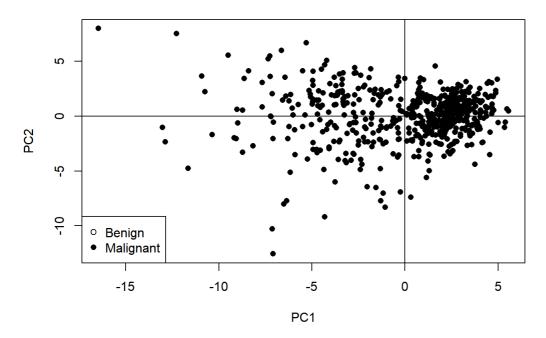
```
##
## F test to compare two variances
##
## data: PC29 by breast_cancer$diagnosis
## F = 0.21294, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.1666460 0.2700465
## sample estimates:
## ratio of variances
## ratio of variances
## 0.2129389</pre>
```

var.test(PC30~breast_cancer\$diagnosis,data=diag_pca)

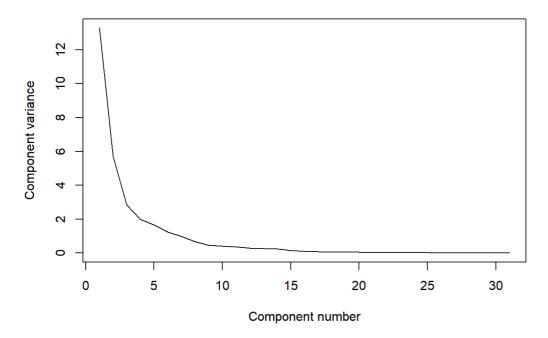
```
## F test to compare two variances
\# \#
## data: PC30 by breast_cancer$diagnosis
## F = 0.23317, num df = 356, denom df = 211, p-value < 2.2e-16
\#\# alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.1824782 0.2957024
## sample estimates:
## ratio of variances
           0.2331693
var.test(PC31~breast_cancer$diagnosis,data=diag_pca)
## F test to compare two variances
##
## data: PC31 by breast_cancer$diagnosis
\#\# F = 0.26577, num df = 356, denom df = 211, p-value < 2.2e-16
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.2079951 0.3370519
## sample estimates:
## ratio of variances
##
          0.2657746
# Levene's tests (one-sided)
library(car)
(LTPC1 <- leveneTest(PC1~breast_cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
   Df F value Pr(>F)
##
## group 1 62.132 1.654e-14 ***
##
        567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(p_PC1_1sided <- LTPC1[[3]][1]/2)</pre>
## [1] 8.268824e-15
(LTPC2 <- leveneTest(PC2~breast_cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
    Df F value
                     Pr(>F)
## group 1 18.786 1.73e-05 ***
\# \#
        567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(p_PC2_1sided=LTPC2[[3]][1]/2)
## [1] 8.651441e-06
(LTPC3 <- leveneTest(PC3~breast_cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
              27.65 2.063e-07 ***
## group 1
##
        567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(p_PC3_1sided <- LTPC3[[3]][1]/2)</pre>
## [1] 1.031266e-07
(LTPC4 <- leveneTest(PC4~breast_cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
     Df F value Pr(>F)
## group 1 2e-04 0.989
        567
##
(p_PC4_1sided <- LTPC4[[3]][1]/2)</pre>
## [1] 0.4944984
(LTPC5 <- leveneTest(PC5~breast_cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
##
        Df F value Pr(>F)
## group 1 6.8535 0.009083 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(p_PC5_1sided <- LTPC5[[3]][1]/2)</pre>
## [1] 0.004541533
# Plotting the scores for the first and second components
\verb|plot(diag_pca\$PC1, diag_pca\$PC2, pch=ifelse(diag_pca\$diagnosis == "S", 1, 16), \verb|xlab="PC1", ylab="PC2", main="569]|
entries against values for PC1 & PC2")
abline (h=0)
abline (v=0)
legend("bottomleft", legend=c("Benign", "Malignant"), pch=c(1,16))
```

569 entries against values for PC1 & PC2

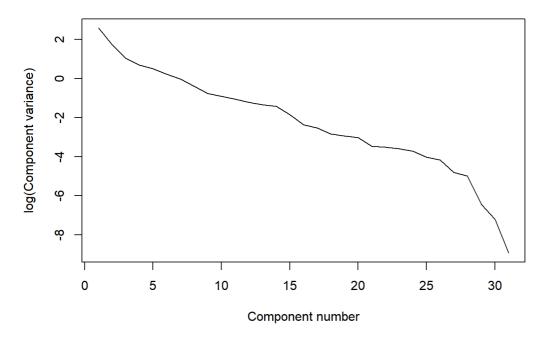


Scree diagram



plot(log(eigen_breast_cancer), xlab = "Component number", ylab = "log(Component variance)", type="l", main = "
Log(eigenvalue) diagram")

Log(eigenvalue) diagram



print(summary(breast_cancer_pca))

```
## Importance of components:
##
                          PC1
                                PC2
                                       PC3
                                                PC4
                                                       PC5
                      3.6453 2.3868 1.68386 1.40761 1.28406 1.11116
## Standard deviation
## Proportion of Variance 0.4286 0.1838 0.09146 0.06391 0.05319 0.03983
## Cumulative Proportion 0.4286 0.6124 0.70388 0.76779 0.82098 0.86081
##
                           PC7 PC8 PC9 PC10 PC11 PC12
                      0.98908 0.81961 0.67882 0.6349 0.59089 0.54212
## Standard deviation
## Proportion of Variance 0.03156 0.02167 0.01486 0.0130 0.01126 0.00948
## Cumulative Proportion 0.89237 0.91404 0.92890 0.9419 0.95317 0.96265
##
                         PC13 PC14 PC15 PC16 PC17 PC18
## Standard deviation 0.51103 0.49125 0.39620 0.30680 0.28251 0.2430
## Proportion of Variance 0.00842 0.00778 0.00506 0.00304 0.00257 0.0019
## Cumulative Proportion 0.97107 0.97886 0.98392 0.98696 0.98953 0.9914
##
                         PC19
                               PC20 PC21
                                             PC22 PC23
## Standard deviation
                      0.2293 0.22163 0.1763 0.17304 0.16562 0.15572
## Proportion of Variance 0.0017 0.00158 0.0010 0.00097 0.00088 0.00078
## Cumulative Proportion 0.9931 0.99472 0.9957 0.99669 0.99757 0.99835
                         PC25 PC26 PC27 PC28 PC29 PC30
##
## Standard deviation 0.13431 0.1244 0.09040 0.08305 0.03987 0.02736
## Proportion of Variance 0.00058 0.0005 0.00026 0.00022 0.00005 0.00002
## Cumulative Proportion 0.99893 0.9994 0.99970 0.99992 0.99997 1.00000
##
                         PC31
## Standard deviation 0.01153
## Proportion of Variance 0.00000
## Cumulative Proportion 1.00000
#View(breast cancer pca)
```

```
diag(cov(breast cancer pca$x))
```

```
PC1 PC2
                          PC3
                                   PC4
## 1.328806e+01 5.696805e+00 2.835395e+00 1.981357e+00 1.648815e+00
## PC6 PC7 PC8 PC9 PC10
## 1.234673e+00 9.782732e-01 6.717530e-01 4.607924e-01 4.031331e-01
## PC11 PC12 PC13 PC14 PC15
## 3.491550e-01 2.938904e-01 2.611469e-01 2.413302e-01 1.569736e-01
## PC16 PC17 PC18 PC19 PC20
## 9.412853e-02 7.980995e-02 5.904627e-02 5.259119e-02 4.912193e-02
  PC21 PC22 PC23 PC24 PC25
##
## 3.107078e-02 2.994121e-02 2.743052e-02 2.424902e-02 1.803936e-02
## PC26 PC27 PC28 PC29 PC30
## 1.547973e-02 8.171699e-03 6.898103e-03 1.589338e-03 7.483761e-04
    PC31
##
## 1.330402e-04
```

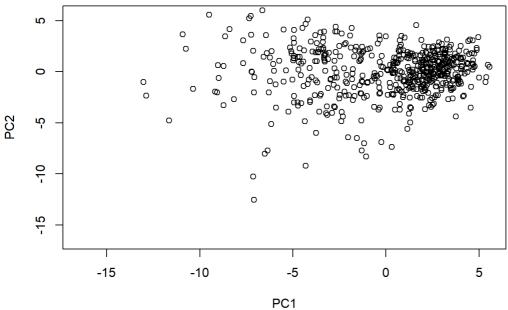
```
xlim <- range(breast cancer pca$x[,1])</pre>
head(breast cancer pca$x[,1])
```

```
## [1] 2.501946 1.467439 2.929028 1.995342 2.500252 2.018308
```

head(breast_cancer_pca\$x)

```
PC2
                        PC3
                                  PC4
                                                PC5
## [1,] 2.501946 -0.09694805 -0.4489597 2.3341176 0.69771548 -0.2430058
## [2,] 1.467439 -1.68630059 1.1542039 0.3362109 0.45962538 1.2308248
## [3,] 2.929028 -0.38319924 -0.8955891 -0.1164828 0.98441377 -0.2587872
## [4,] 1.995342 -1.33046592 1.1172876 2.0502761 0.25303846 -1.5539634
## [5,] 2.500252 2.01035097 -0.7584035 1.9862169 -1.13537096 0.5940361
## [6,] 2.018308 -0.78242095 0.1125197 -0.6532280 0.01841577 0.6914453
                            PC9
           PC7
                  PC8
                                       PC10
## [1,] 0.5092015 -1.11423307 0.2840243 0.32463197 -0.3245353 0.04981306
## [2,] 0.2937434 0.10000461 -0.0668399 0.42612180 0.4564029 1.19357566
## [3,] -0.3303385 0.03599041 0.8734350 0.02222192 0.4208602 -0.06687286
## [5,] 0.1198201 -0.48279704 -0.2727816 -0.29439485 -0.3577533 0.03266208
## [6,] 0.1454026 0.06214539 0.2342454 0.73681239 -0.3671239 -0.77029743
##
           PC13
                 PC14
                          PC15
                                     PC16
## [1,] -0.19760220 0.1134403 -0.059302558 0.16637723 -0.04286656
## [2,] 0.01807424 -0.2824292 -0.204858888 -0.07067959 0.03088787
## [3,] 0.37435458 0.2585457 -0.330274216 -0.13000189 -0.24616091
## [4,] -0.90968379 0.2179117 -0.665825669 0.10213387 -0.10289446
## [5,] -0.35547138 -0.1480140 -0.005540503 -0.06495881 0.22273309
## [6,] -0.49542291 -0.2992431  0.049952835 -0.20161083  0.14920422
##
            PC18 PC19 PC20 PC21
## [1,] -0.104542766 -0.03484189 -0.09691187 -0.02846306 -0.00673628
## [2,] -0.405534243 -0.02886103 -0.05262226 -0.05987170 0.05868642
## [3,] 0.327711259 0.15937793 -0.13804895 -0.13489743 0.10080029
## [4,] 0.197085181 0.36251771 -0.40018239 -0.10302093 -0.28821708
## [5,] -0.129129156 -0.35877054 0.08515543 -0.08500541 -0.06332008
## [6,] -0.002229379 -0.08178568 0.18970936 -0.06872875 0.09669594
\# \#
            PC23
                 PC24 PC25 PC26 PC27
## [2,] 0.070978613 -0.030822339 -0.016741580 0.04173030 -0.059332996
## [3,] 0.053909008 0.085484364 0.038277664 -0.04151896 -0.035546410
## [4,] 0.182045907 0.222848059 -0.115720065 -0.03676948 -0.148171674
## [5,] 0.043591030 0.008165322 0.002738052 0.05983731 0.046167735
##
           PC28 PC29 PC30 PC31
## [1,] 0.05356131 0.015184882 0.015985406 0.001396101
## [3,] -0.07653067 -0.014640388 0.010307894 0.009074601
## [4,] -0.01711665 -0.047828494 0.023862995 0.000265075
## [5,1
      0.03835364 0.032450800 -0.002312178 -0.002563269
## [6,] 0.00955434 -0.004403431 0.003869919 -0.002931194
```

plot(breast cancer pca\$x,xlim=xlim,ylim=xlim)



```
## Principal Components Analysis
## Call: principal(r = breast cancer[-2], nfactors = 4, rotate = "varimax")
## Standardized loadings (pattern matrix) based upon correlation matrix
##
                RC1 RC2 RC3 RC4 h2 u2 com
## id
               0.13 -0.10 0.10 0.07 0.042 0.958 3.3
## perimeter_mean 0.95 0.17 -0.11 0.10 0.954 0.046 1.1
## area mean 0.97 0.10 -0.08 0.09 0.960 0.040 1.1
## smoothness_mean 0.16 0.65 0.26 -0.19 0.547 0.453 1.6
## compactness_mean 0.46 0.77 0.32 0.05 0.910 0.090 2.0
## concavity_mean 0.66 0.61 0.31 0.10 0.908 0.092 2.5
## points_mean
               0.80 0.51 0.15 0.05 0.921 0.079 1.8
## perimeter_se
## area se
               0.82 0.04 0.42 0.04 0.855 0.145 1.5
## area_se
               0.88 -0.02 0.26 0.01 0.836 0.164 1.2
## smoothness_se -0.14 0.00 0.70 -0.05 0.518 0.482 1.1
0.94 0.16 -0.12 0.12 0.947 0.053 1.1
## area worst
## smoothness_worst 0.06 0.75 0.01 -0.01 0.572 0.428 1.0
## compactness_worst 0.31 0.86 0.06 0.19 0.877 0.123 1.4
## concavity_worst 0.45 0.77 0.10 0.19 0.845 0.155 1.8
## dimension_worst -0.09 0.88 0.18 0.10 0.825 0.175 1.1
##
                   RC1 RC2 RC3 RC4
##
## SS loadings
                 10.15 7.04 4.36 2.25
                  0.33 0.23 0.14 0.07
## Proportion Var
## Cumulative Var
                   0.33 0.55 0.70 0.77
## Proportion Explained 0.43 0.30 0.18 0.09
## Cumulative Proportion 0.43 0.72 0.91 1.00
\# \#
## Mean item complexity = 1.6
## Test of the hypothesis that 4 components are sufficient.
##
\#\# The root mean square of the residuals (RMSR) is 0.06
## with the empirical chi square 1923.24 with prob < 3.8e-216
##
## Fit based upon off diagonal values = 0.98
```

```
round(fit.pc$values, 3)
```

```
## [1] 13.288 5.697 2.835 1.981 1.649 1.235 0.978 0.672 0.461 0.403
## [11] 0.349 0.294 0.261 0.241 0.157 0.094 0.080 0.059 0.053 0.049
## [21] 0.031 0.030 0.027 0.024 0.018 0.015 0.008 0.007 0.002 0.001
## [31] 0.000
```

fit.pc\$loadings

```
## Loadings:
                 RC1 RC2 RC3 RC4
##
## id
                  0.133
## id 0.133
## radius_mean 0.951 0.131 -0.139
## texture_mean 0.252
## perimeter_mean 0.950 0.175 -0.107
## area_mean 0.967 0.101
## smoothness mean 0.159 0.649 0.255 -0.187
## compactness_mean 0.459 0.771 0.320
## concavity_mean 0.659 0.606 0.311 0.102
## points_mean
                  0.798 0.508 0.149
## symmetry_mean
                   0.137 0.585 0.332
## dimension_mean -0.314 0.660 0.532
## radius_se 0.832
## texture_se
## perimeter_se 0.823
                                0.395
                   -0.209 0.597 0.538
                  0.823 0.418
                  0.876
                               0.262
## area_se
## smoothness_se -0.138 0.705
## compactness_se 0.204 0.514 0.667 0.111
## perimeter_worst
## area_worst 0.944 0.165
## smoothness_worst 0.754
                   0.944 0.165 -0.118 0.120
## compactness_worst 0.312 0.861 0.186
## concavity_worst 0.450 0.773 0.100 0.187
## points_worst
                  0.678 0.668 0.106
## dimension_worst 0.717
## ##
                        0.880 0.181 0.104
##
##
                  RC1 RC2 RC3 RC4
## SS loadings 10.151 7.037 4.361 2.252
## Proportion Var 0.327 0.227 0.141 0.073
## Cumulative Var 0.327 0.554 0.695 0.768
```

```
# Loadings with more digits
for (i in c(1,3,2,4)) { print(fit.pc$loadings[[1,i]])}
```

```
## [1] 0.1330256

## [1] 0.0991346

## [1] -0.09752479

## [1] 0.06781887
```

```
# Communalities
fit.pc$communality
```

```
id radius_mean texture_mean perimeter_mean 0.04163396 0.95069170 0.89684853 0.95431848
##
##
         area_mean smoothness_mean compactness_mean concavity_mean
                                                           0.90807129
##
         0.96014950 0.54720158 0.90971908
                      symmetry_mean
                                        dimension_mean
      points_mean
##
                                                                radius_se
                                                            0.84993832
                                         0.82580280
##
         0.92076209
                         0.47390203
                      perimeter_se
\#\,\#
         texture_se
                                               area_se
                                                            smoothness_se
                                           0.83634402
        0.69271471
##
                          0.85520824
                                                              0.51759585
                        concavity_se
                                          points_se symmetry_se 0.69650774 0.45095156
##
    compactness se
##
       0.76240129
                          0.62241576

        dimension_se
        radius_worst
        texture_worst
        perimeter_worst

        0.71272740
        0.97219376
        0.95565236
        0.97796884

##
##
        area_worst smoothness_worst compactness_worst concavity_worst
                                                          0.84471615
##
        0.94731995 0.57201913 0.87681767
##
      points_worst symmetry_worst dimension_worst
##
        0.91825491
```

Rotated factor scores, Notice the columns ordering: RC1, RC3, RC2 and RC4
head(fit.pc\$scores)

```
## [1,] -0.3200066 -0.20898001 -0.25160464 -1.75616620

## [2,] -0.5649931 -0.22081178    0.85830109 -0.28887483

## [3,] -0.8242652 -0.03734588 -0.52158508 -0.09098986

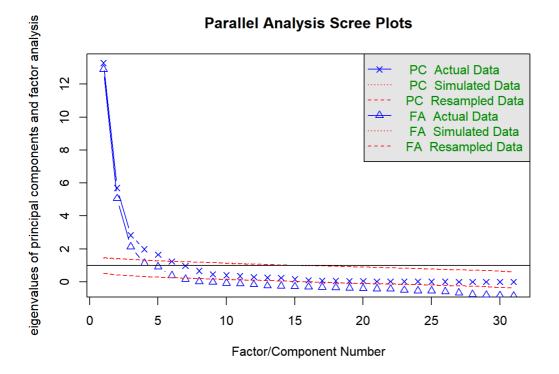
## [4,] -0.3869942 -0.38180634    0.79329588 -1.49911551

## [5,]    0.1145874 -0.61206123 -0.91421184 -1.46638605

## [6,] -0.6900800 -0.15424020    0.07996063    0.35700314
```

```
# Play with FA utilities
fa.parallel(breast_cancer[-2]) # See factor recommendation
```

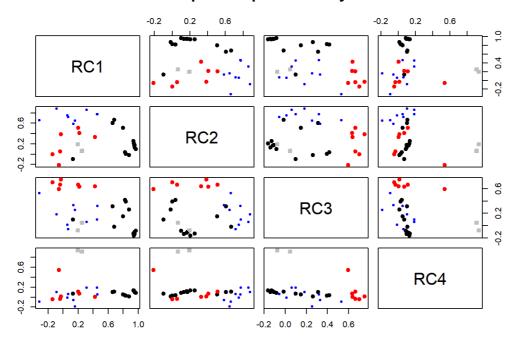
```
## Warning in fa.stats(r = r, f = f, phi = phi, n.obs = n.obs, np.obs
## = np.obs, : The estimated weights for the factor scores are probably
## incorrect. Try a different factor extraction method.
```



```
## Parallel analysis suggests that the number of factors = 6 and the number of components = 5
```

fa.plot(fit.pc) # See Correlations within Factors

Principal Component Analysis



fa.diagram(fit.pc) # Visualize the relationship

Components Analysis

