MVA3.r

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```
wisc_bc_df <- read.csv("C://Users//Yshah//Downloads//Rutgers Sem 2//MVA//wisc_bc_data.csv")
head(wisc_bc_df)</pre>
```

```
id diagnosis radius_mean texture_mean perimeter_mean area_mean
## 1 87139402 B 12.32 12.39 78.85 464.1
## 2 8910251
                  В
                         10.60
                                     18.95
                                                   69.28
            B 11.04
B 11.28
B 15.19
B 11.57
## 3 905520
                                     16.83
                                                  70.92
## 4 868871
                                     13.39
                                                   73.00
                                13.21 97.65
19.04 74.20
## 5 9012568
                                                            711.8
## 6 906539
                                                           409.7
## smoothness_mean compactness_mean concavity_mean points_mean
     0.10280 0.06981 0.03987 0.03700
## 1
                        0.11470 0.06387
0.07804 0.03046
0.11360 0.04635
           0.09688
## 2
                                                  0.02642
## 3
          0.10770
                                                  0.02480
          0.11640
                                                 0.04796
## 4
                         0.06934
          0.07963
                                      0.03393
                                                 0.02657
## 5
         0.08546 0.07722 0.05485 0.01428
## 6
## symmetry mean dimension mean radius se texture se perimeter se area se
## 1 0.1959 0.05955 0.2360 0.6656 1.670 17.43
## 2
         0.1922
                     0.06491 0.4505 1.1970
                                                      3.430 27.10
## 3
         0.1714
                     0.06340 0.1967 1.3870
                                                     1.342 13.54
         0.1771
                     0.06072 0.3384 1.3430
                                                     1.851 26.33
## 4
                     0.06267 0.2864 1.4400 2.206 20.30 actness se concevity
         0.1721
                     0.05544 0.1783 0.4125
## 5
         0.2031
## 6
## smoothness_se compactness_se concavity_se points_se symmetry_se
## 1
     0.008045 0.011800 0.01683 0.012410 0.01924
## 2
        0.007470
                      0.035810
                                  0.03354 0.013650
                                                      0.03504
                               0.01056 0.007483
       0.005158
## 3
                     0.009355
                                                      0.01718
                    0.034980
                                 0.02187 0.019650
       0.011270
                                                     0.01580
## 4

    0.005012
    0.014850
    0.01551
    0.009155
    0.01647

    0.007278
    0.020470
    0.04447
    0.008799
    0.01868

## 5
## 6
## dimension se radius worst texture worst perimeter worst area worst
## 1 0.002248 13.50 15.64 86.97 549.1
## 2 0.003318
                    11.88
                                 22.94
                                                78.28
                                                          424.8
## 3 0.002198
                    12.41
                                 26.44
                                                79.93
                                                          471.4
    0.003442
                    11.92
                                 15.77
## 4
                                                76.53
                                                          434.0
     0.001767
## 5 0.001767 16.20 15.73 104.50
## 6 0.003339 13.07 26.98 86.43
                                                          819.1
                                                          520.5
## smoothness worst compactness worst concavity worst points worst

      0.1385
      0.1266
      0.12420
      0.09391

      0.1213
      0.2515
      0.19160
      0.07926

## 1
## 2
                                                      0.07926
                            0.1482
                                         0.10670
                                                     0.07431
## 3
            0.1369
                            0.1822
                                         0.08669
                                                     0.08611
## 4
            0.1367
                                        0.13620 0.08178
0.25600 0.06664
                            0.1737
## 5
            0.1126
## 6
           0.1249
                           0.1937
## symmetry worst dimension worst
## 1 0.2827 0.06771
## 2
          0.2940
                       0.07587
          0.2998
                       0.07881
## 3
          0.2102
## 4
                       0.06784
          0.2487
## 5
                        0.06766
## 6
          0.3035
                        0.08284
```

```
#Renmaing the dataset
cancer<-wisc_bc_df
library("ggplot2")
library("corrplot")</pre>
```

```
## corrplot 0.84 loaded
```

```
library("reshape")

#number of obsevations and data type
str(cancer)
```

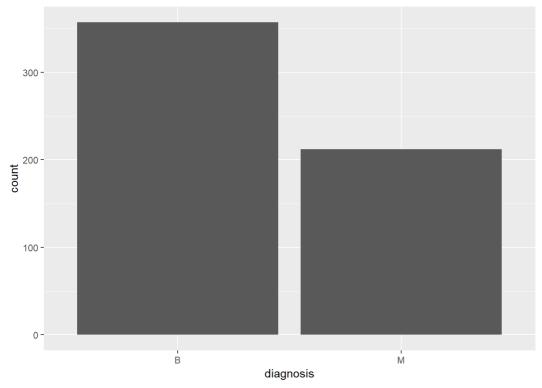
```
## 'data.frame': 569 obs. of 32 variables:
## $ radius_mean
                     : num 12.3 10.6 11 11.3 15.2 ...
## $ texture mean : num 12.4 18.9 16.8 13.4 13.2 ...
## $ perimeter_mean : num 78.8 69.3 70.9 73 97.7 ...
## $ area_mean : num 464 346 373 385 712 ...
## $ smoothness_mean : num 0.1028 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 ...
## $ radius_se : num 0.236 0.451 0.197 0.338 0.178 ...
## $ texture_se : num 0.666 1.197 1.387 1.343 0.412 ...
## $ perimeter_se : num 1.67 3.43 1.34 1.85 1.34 ...
## $ area_se : num 17.4 27.1 13.5 26.3 17.7 ...
## $ 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 ...
                     : num 0.0192 0.035 0.0172 0.0158 0.0165 ...
## $ symmetry_se
## $ 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 ...
                      : num 549 425 471 434 819 ...
## $ area_worst
## $ 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 ...
## $ points worst : num 0.0939 0.0793 0.0743 0.0861 0.0818 ...
## $ 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 ...
```

#gives you the summary of the dataset
summary(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
                 Median :0.03350 Median :0.1792 Median :0.06154
## 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.:0.16140
   3rd Qu.:0.33910
                  3rd Qu.:0.3829
##
                 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
```

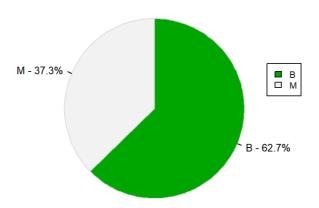
```
##
## B M
## 357 212
```

```
#Bar plot
ggplot(data=cancer, aes(x=diagnosis)) + geom_bar(stat = "count")
```



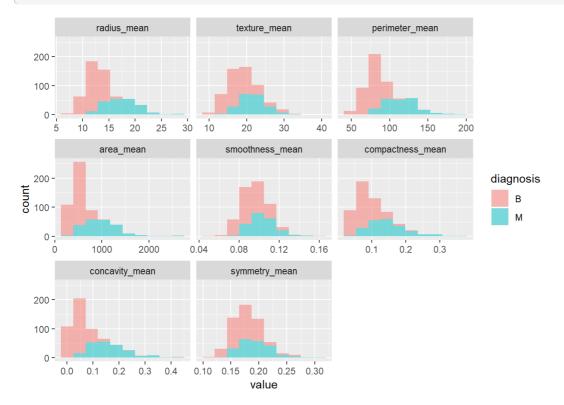
```
#Pie chart represented in frequency
diagnosis.prop.table <- prop.table(diagnosis.table)*100
diagnosis.prop.df <- as.data.frame(diagnosis.prop.table)
pielabels <- sprintf("%s - %3.1f%s", diagnosis.prop.df[,1], diagnosis.prop.table, "%")
colors <- terrain.colors(2)
pie(diagnosis.prop.table,
    labels=pielabels,
    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

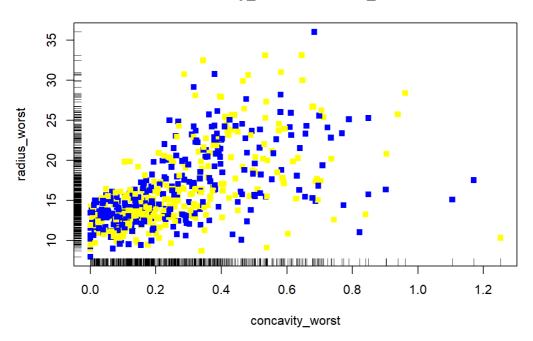


```
#Plot histograms of "mean" variables group by diagnosis
data_mean <- cancer[,c("diagnosis", "radius_mean", "texture_mean", "perimeter_mean", "area_mean", "smoothnes
s_mean", "compactness_mean", "concavity_mean", "symmetry_mean")]

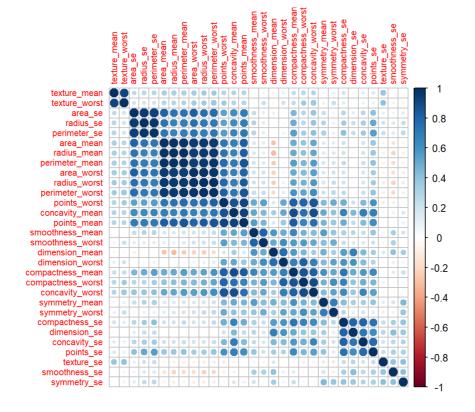
#Plot histograms
ggplot(data = melt(data_mean, id.var = "diagnosis"), mapping = aes(x = value)) +
geom_histogram(bins = 10, aes(fill=diagnosis), alpha=0.5) + facet_wrap(~variable, scales ='free_x')</pre>
```



Concavity_worst vs radius_worst

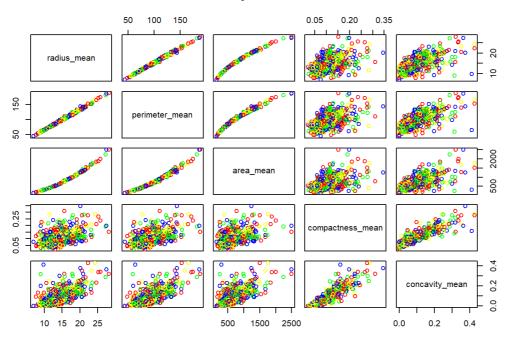


```
#Corelation Matrix of columns
corMatMy <- cor(cancer[,3:32])
corrplot(corMatMy, order = "hclust", tl.cex = 0.7)</pre>
```



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

Scatterplot Matrix



```
names (cancer)
```

```
## [1] "id"
                           "diagnosis"
                                               "radius mean"
## [4] "texture mean"
                           "perimeter mean"
                                               "area mean"
## [7] "smoothness_mean"
                           "compactness_mean"
                                               "concavity_mean"
                           "symmetry_mean"
                                               "dimension_mean"
## [10] "points_mean"
## [13] "radius se"
                           "texture_se"
                                               "perimeter_se"
## [16] "area se"
                           "smoothness_se"
                                               "compactness se"
## [19] "concavity_se"
                           "points se"
                                               "symmetry se"
## [22] "dimension se"
                           "radius worst"
                                               "texture worst"
                           "area_worst"
## [25] "perimeter_worst"
                                               "smoothness_worst"
## [28] "compactness_worst" "concavity_worst"
                                               "points worst"
## [31] "symmetry_worst"
                           "dimension_worst"
```

```
#Multivariate analysis
#T TEST
with(data=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=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=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=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>
```

 $\\ with (\texttt{data} = \texttt{cancer}, \texttt{t.test} (\texttt{smoothness}_\texttt{mean}[\texttt{diagnosis} = \texttt{"B"}], \texttt{smoothness}_\texttt{mean}[\texttt{diagnosis} = \texttt{"M"}], \texttt{var.equal} = \texttt{TRUE}))\\$

```
##
## 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>
```

with(data=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=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</pre>
```

with(data=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</pre>
```

with(data=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=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
## sample estimates:
## mean of x mean of y
## 0.06286739 0.06268009
#Hotelling's T2 test
#install.packages("Hotelling")
library (Hotelling)
## Loading required package: corpcor
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=cancer)
# Output of the function hotelling.test is given
cat("T2 statistic =",t2testcan$stat[[1]],"\n")
## T2 statistic = 1220.313
#print(t2testcan)
# T2 statistic is located in the first element of the list "stat"
#View(t2testcan)
## Levene's tests based on absolute differences around means using t-tests. Standarizing the sparrows data s
matstand <- scale(cancer[,3:10])</pre>
head(matstand)
      radius mean texture mean perimeter mean area mean smoothness mean
## [1,] -0.5128453 -1.60418301 -0.5399006 -0.5421468
                                                          0.4578825
        -1.0009202 -0.07896900
## [2,]
                                   -0.9337442 -0.8766033
                                                              0.0369535
## [3,] -0.8760638 -0.57187353
                                                             0.8062867
                                   -0.8662517 -0.8004484
## [4,] -0.8079604 -1.37168088
                                   -0.7806514 -0.7674858
                                                              1.4248817
## [5,] 0.3015589 -1.41353126
                                   0.2337944 0.1617181
                                                             -1.1895712
## [6,] -0.7256686 -0.05804381
                                  -0.7312666 -0.6967299
                                                             -0.7750414
      compactness mean concavity mean points mean
## [1,] -0.6538379 -0.6137661 -0.30717196
             0.1961461
## [2,]
                          -0.3127117 -0.57983238
           -0.4980044 -0.7318045 -0.62158190
## [3,]
             0.1753178 -0.5324814 -0.02471844
## [4,]
## [5,]
             -0.6627373
                           -0.6882771 -0.57596668
## [6,]
            -0.5135309
                           -0.4258580 -0.89269604
matsurv <- matstand[cancer$diagnosis =="B",]</pre>
head(matsurv)
```

```
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
## [3,]
             -0.4980044 -0.7318045 -0.62158190
                          -0.5324814 -0.02471844
             0.1753178
## [4,]
             -0.6627373
                           -0.6882771 -0.57596668
## [5,]
## [6,]
             -0.5135309
                           -0.4258580 -0.89269604
matnosurv <- matstand[cancer$diagnosis == "M",]</pre>
vecmediansurv <- apply(matsurv, 2, median)</pre>
# in the above 2 represents column. Hence, we are asking for column median
vecmediansurv
##
       radius mean
                     texture mean perimeter mean
                                                          area mean
##
        -0.5468970
                       -0.4416723
                                      -0.5674737
                                                          -0.5583439
                                                       points_mean
## smoothness mean compactness mean
                                    concavity mean
##
       -0.3981961
                   -0.5500751
                                       -0.6486382
                                                          -0.6566309
vecmediannosurv <- apply(matnosurv, 2, median)</pre>
matabsdevsurv <- abs(matsurv - matrix(rep(vecmediansurv,nrow(matsurv)),nrow=nrow(matsurv), byrow=TRUE))</pre>
matabsdevnosurv <- abs(matnosurv - matrix(rep(vecmediannosurv,nrow(matnosurv)),nrow=nrow(matnosurv), byrow=T</pre>
RUE))
head (matabsdevnosurv)
##
       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
## [4,] 0.7846089 0.55800512
                                   0.8436568 0.8951047
                                                           0.41239670
                                  0.9782303 0.9885934
                                                           0.10665432
## [5,] 1.0002699 1.31828711
                                   0.4856171 0.5671838
## [6,] 0.5746231 0.88583314
                                                           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
            0.97419643
                          0.96525570 1.07517890
## [4,]
## [5,]
           0.08236646 0.92147737 0.91720079
           ## [6,]
matabsdev.all <- rbind(matabsdevsurv, matabsdevnosurv)</pre>
matabsdev.all <- data.frame(cancer$diagnosis, matabsdev.all)</pre>
t.test(matabsdev.all$radius_mean[cancer$diagnosis == "B"], matabsdev.all$radius_mean[cancer$diagnosis == "M"]
, alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$radius mean[cancer$diagnosis == "B"] and matabsdev.all$radius mean[cancer$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$texture mean[cancer$diagnosis == "B"], matabsdev.all$texture mean[cancer$diagnosis == "M"]
"], alternative="less", var.equal = TRUE)
##
   Two Sample t-test
##
## data: matabsdev.all$texture_mean[cancer$diagnosis == "B"] and matabsdev.all$texture_mean[cancer$diagnosi
s == "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 y
## 0.6364762 0.7509490
t.test(matabsdev.all$perimeter mean[cancer$diagnosis == "B"],matabsdev.all$perimeter mean[cancer$diagnosis =
= "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$perimeter_mean[cancer$diagnosis == "B"] and matabsdev.all$perimeter_mean[cancer$diag
nosis == "M"l
## 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 y
## 0.5125724 0.5033541
t.test(matabsdev.all$area mean[cancer$diagnosis == "B"], matabsdev.all$area mean[cancer$diagnosis == "M"], al
ternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$area mean[cancer$diagnosis == "B"] and matabsdev.all$area mean[cancer$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$smoothness mean[cancer$diagnosis == "B"], matabsdev.all$smoothness mean[cancer$diagnosis
== "M"], alternative="less", var.equal = TRUE)
##
##
   Two Sample t-test
## data: matabsdev.all$smoothness_mean[cancer$diagnosis == "B"] and matabsdev.all$smoothness_mean[cancer$di
agnosis == "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$compactness_mean[cancer$diagnosis == "B"], matabsdev.all$compactness_mean[cancer$diagnos
```

t.test(matabsdev.all\$compactness_mean[cancer\$diagnosis == "B"], matabsdev.all\$compactness_mean[cancer\$diagnos
is == "M"], alternative="less", var.equal = TRUE)

```
## Two Sample t-test
##
## data: matabsdev.all$compactness_mean[cancer$diagnosis == "B"] and matabsdev.all$compactness_mean[cancer$
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$concavity_mean[cancer$diagnosis == "B"], matabsdev.all$concavity_mean[cancer$diagnosis =
= "M"], alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$concavity mean[cancer$diagnosis == "B"] and matabsdev.all$concavity mean[cancer$diag
nosis == "M"1
## 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 y
## 0.4977532 0.4456302
t.test(matabsdev.all$points_mean[cancer$diagnosis == "B"], matabsdev.all$points_mean[cancer$diagnosis == "M"]
, alternative="less", var.equal = TRUE)
##
## Two Sample t-test
##
## data: matabsdev.all$points mean[cancer$diagnosis == "B"] and matabsdev.all$points mean[cancer$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 (matstand)

```
##
      radius_mean texture_mean perimeter_mean area_mean smoothness_mean
                                                      0.4578825
## [1,] -0.5128453 -1.60418301 -0.5399006 -0.5421468
## [2,] -1.0009202 -0.07896900
                                 -0.9337442 -0.8766033
                                                           0.0369535
       -0.8760638 -0.57187353
                                 -0.8662517 -0.8004484
## [3,1
                                                           0.8062867
## [4,] -0.8079604 -1.37168088
                                                          1.4248817
                                 -0.7806514 -0.7674858
       0.3015589 -1.41353126
                                 0.2337944 0.1617181
                                                         -1.1895712
## [5,]
## [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,]
## [3,]
           -0.4980044
                         -0.7318045 -0.62158190
            0.1753178
## [4,]
                         -0.5324814 -0.02471844
            -0.6627373
                         -0.6882771 -0.57596668
## [5,]
            -0.5135309
                          -0.4258580 -0.89269604
## [6,]
```

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

```
cancer.diagnosis radius_mean texture_mean perimeter_mean area_mean
                  B -0.5128453 -1.60418301 -0.5399006 -0.5421468
## 1
                  В -1.0009202 -0.07896900
## 2
                                               -0.9337442 -0.8766033
                  В -0.8760638 -0.57187353
                                              -0.8662517 -0.8004484
## 3
## 4
                  B -0.8079604 -1.37168088
                                              -0.7806514 -0.7674858
## 5
                  B 0.3015589 -1.41353126
                                               0.2337944 0.1617181
                  B -0.7256686 -0.05804381 -0.7312666 -0.6967299
## 6
## smoothness_mean compactness_mean concavity_mean points_mean
## 1
        0.4578825
                      -0.6538379 -0.6137661 -0.30717196
## 2
         0.0369535
                        0.1961461
                                      -0.3127117 -0.57983238
                        -0.4980044
## 3
         0.8062867
                                      -0.7318045 -0.62158190
## 4
                        0.1753178
                                      -0.5324814 -0.02471844
         1.4248817
## 5
        -1.1895712
                        -0.6627373
                                      -0.6882771 -0.57596668
## 6
         -0.7750414
                         -0.5135309
                                       -0.4258580 -0.89269604
colnames(matstand.all) <- colnames(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=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)
## cancer.diagnosis radius mean texture mean perimeter mean area mean
                 B 0.03405174 1.1625107 0.02757317 0.01619713
## 1
                  B 0.45402322
                                              0.36627050 0.31825946
## 2
                                 0.3627033
## 3
                  B 0.32916684
                                  0.1302012
                                               0.29877796 0.24210452
## 4
                  B 0.26106335
                                  0.9300085
                                               0.21317766 0.20914193
## 5
                  B 0.84845589
                                 0.9718589
                                               0.80126817 0.72006202
                                             0.16379288 0.13838603
                                0.3836285
## 6
                  B 0.17877164
## smoothness_mean compactness_mean concavity_mean points_mean
       0.8560787 0.10376281 0.03487214 0.34945891
## 1
## 2
         0.4351496
                       0.74622121
                                     0.33592655 0.07679849
## 3
        1.2044828
                       0.05207075 0.08316628 0.03504898
                       0.72539291
                                     0.11615683 0.63191243
## 4
         1.8230778
## 5
         0.7913750
                       0.11266217 0.03963883 0.08066419
## 6
         0.3768453
                       0.03654420 0.22278026 0.23606517
#install.packages("car")
library(car)
```

```
## 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=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=cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 0.684 0.4086
##
       567
leveneTest(perimeter_mean ~ diagnosis, data=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=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=cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 0.8377 0.3604
##
       567
leveneTest(compactness mean~ diagnosis, data=cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 39.892 5.428e-10 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(concavity mean~ diagnosis, data=cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 70.484 3.723e-16 ***
##
```

```
leveneTest(points_mean ~ diagnosis, data=cancer)
```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

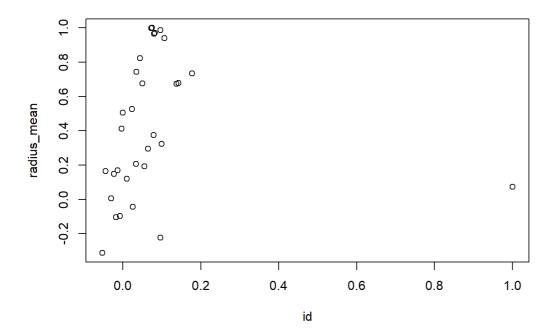
```
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 94.906 < 2.2e-16 ***
## 567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
leveneTest(symmetry_mean ~ diagnosis, data=cancer)
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 2.036 0.1542
      567
##
leveneTest(dimension_mean ~ diagnosis, data=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 Analysis
#PCA
dim(cancer)
## [1] 569 32
attach (cancer)
head(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
                               16.83
## 3 905520
               В
                     11.04
                                           70.92
                     11.28
                               13.39
## 4 868871
               В
                                          73.00
                               13.21
## 5 9012568
               В
                     15.19
                                          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
## 4
         0.11640
                                0.04635
                 0.06934
0.07722
## 5
         0.07963
                                 0.03393
                                          0.02657
        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
                                   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
## smoothness_se compactness_se concavity_se points_se symmetry_se
## 1 0.008045 0.011800 0.01683 0.012410 0.01924
      0.007470
                                             0.03504
## 2
                  0.035810
                            0.03354 0.013650
                 0.009355
## 3
     0.005158
                                             0.01718
                            0.01056 0.007483
      0.011270
## 4
                  0.034980
                            0.02187 0.019650
                                             0.01580
                  0.014850
       0.005012
                             0.01551 0.009155
## 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
## 2
     0.003318
                  11.88
                                        78.28
                                                 424.8
                                                 471.4
## 3
    0.002198
                 12.41
                            26.44
                                        79.93
## 4 0.003442
                 11.92
                            15.77
                                        76.53
## 5 0.001767
                            15.73
                 16.20
                                       104.50
                                                 819.1
## 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
## 5
          0.1126
                        0.1737
                                   0.25600
                                   0.13620
                                              0.08178
                      0.1937
## 6
          0.1249
                                             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
```

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

```
id radius_mean texture_mean perimeter_mean
## id 1.00000000 0.07462647 0.09976989 0.07315941
## radius_mean 0.07462647 1.00000000 0.32378189 0.99785528
## texture_mean 0.09976989 0.32378189 1.00000000 0.32953306
## perimeter_mean 0.07315941 0.99785528 0.32953306
                                                                             1.00000000
## area_mean 0.09689282 0.98735717 0.32108570 0.98650680
## smoothness mean -0.01296820 0.17058119 -0.02338852 0.20727816
## area mean smoothness mean compactness mean concavity mean
## id
                       0.09689282 -0.01296820 0.0000957011 0.05007995
## radius_mean 0.98735717 0.17058119 0.5061235775 0.67676355
## texture_mean 0.32108570 -0.02338852 0.2367022221 0.30241783
## perimeter_mean 0.98650680 0.20727816 0.5569362109
## area_mean 1.00000000 0.17702838 0.4985016822
## smoothness_mean 0.17702838 1.00000000 0.6591232152
                                                                                       0.71613565
                                                                                       0.68598283
0.52198377
       points_mean symmetry_mean dimension_mean radius se
##
## id 0.0441581 -0.02211406 -0.05251145 0.1430476
## radius_mean 0.8225285 0.14774124 -0.31163083 0.6790904
## texture_mean 0.2934641 0.07140098 -0.07643718 0.2758687
## area mean 0.8232689 0.15129308 -0.28310981 0.7325622
## smoothness mean 0.5536952 0.55777479 0.58479200 0.3014671
## texture_se perimeter_se area_se smoothness_se
## id -0.00752619 0.1373311 0.1777419 0.096780574
                       -0.00752619 0.1373311 0.1777419 0.096780574
## id
## radius_mean -0.09731744 0.6741716 0.7358637 -0.222600125
## texture_mean 0.38635762 0.2816731 0.2598450 0.006613777
## perimeter_mean -0.08676108 0.6931349 0.7449827 -0.202694026
## area_mean -0.06628021
                                            0.7266283 0.8000859 -0.166776667
## smoothness_mean 0.06840645 0.2960919 0.2465524 0.332375443
## compactness_se concavity_se points_se symmetry_se ## id 0.03396097 0.05523932 0.07876807 -0.017306295
## id 0.03396097 0.05523932 0.07876807 -0.017306295  
## radius_mean 0.20599998 0.19420362 0.37616896 -0.104320881  
## texture_mean 0.19197461 0.14329308 0.16385103 0.009127168  
## perimeter_mean 0.25074368 0.22808235 0.40721692 -0.081629327  
## area_mean 0.21258255 0.20766006 0.37232028 -0.072496588  
## smoothness_mean 0.31894330 0.24839568 0.38067569 0.20774376
## dimension_se radius_worst texture_worst perimeter_worst ## id 0.025725324 0.08240534 0.06471955 0.07998587
                       0.025725324 0.08240534 0.06471955 0.07998587
0.96513651
                                                                                  0.35803958
                         0.054457520 0.35257295 0.91204459
## perimeter_mean -0.005523391 0.96947636 0.30303837
## area_mean -0.019886963 0.96274609 0.28748863
                                                                                    0.97038689
## area_mean -0.019886963
                                                                                    0.95911957
## area_mean -0.019886963 0.96274609 0.28748863 0.95911957  
## smoothness_mean 0.283606699 0.21312014 0.03607180 0.23885263
     area_worst smoothness_worst compactness_worst
##
                        0.1071865 0.01033803 -0.0029681
## id
## radius_mean 0.9410825
                                             0.11961614
                                                                       0.4134628
## texture mean 0.3435459 0.07750336
                                                                      0.2778296
## perimeter_mean 0.9415498
                                             0.15054940
                                                                       0.4557742
## area_mean 0.9592133 0.12352294 0.3904103
## smoothness_mean 0.2067184 0.80532420 0.4724684
## concavity_worst points_worst symmetry_worst
## id 0.02320274 0.03517358 -0.04422425
## radius_mean 0.52691146 0.74421420 0.16395333
## texture_mean 0.30102522 0.29531584 0.10500791
## perimeter_mean 0.56387926 0.77124079 0.18911504
## area_mean 0.51260592 0.72201663 0.14356991
## smoothness_mean 0.43492571 0.50305335 0.39430948
## dimension_worst
## id -0.029865636
## radius_mean 0.007065886
## texture_mean 0.119205351
## perimeter_mean 0.051018530
## area_mean 0.003737597
## smoothness mean 0.499316369
```

```
c <- (cor(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
cancer_pca <- prcomp(cancer[,-2],scale=TRUE)
head(cancer_pca)</pre>

```
## $sdev
## [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
##
                        PC1
                                   PC2
                                               PC3
## id
                  ## radius mean
## texture_mean
                  ## 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
                 -0.25828025 -0.060541625 0.001758736 0.019497124
## concavity mean
                 -0.26073811 0.034167392 -0.027579607
## points mean
## symmetry_mean
                 -0.13797774 -0.190684979 -0.040962032 0.067502543
                 -0.06414779 -0.366531055 -0.020817875 0.047957856
## dimension_mean
                 -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
                  -0.20307642 0.152771289 0.215790250 0.108568705
## area se
## smoothness se
                 -0.01467821 -0.203189876 0.311787845 0.044368664
                 -0.17028840 -0.232503362 0.154557465 -0.026425360
## compactness se
                 -0.15354367 -0.196846081 0.176560052 0.002248291
## concavity_se
## points_se
                 -0.18340675 -0.129965181 0.223850479 0.075252232
## symmetry_se
                 -0.04241552 -0.183558627 0.285265066 0.046936126
                  -0.10249607 -0.279584139 0.211893354
## dimension se
## radius_worst
                  ## texture_worst
                  -0.23663734 0.199295985 -0.050431945 0.014068572
## perimeter worst
                  ## area_worst
## 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 PC8
## id 0.011327587 -0.316733438 0.9071156324 -0.096362415
## radius_mean -0.038129861 0.029588521 -0.0422987777 -0.116427419
## texture_mean 0.049091450 -0.031394323 0.0149935618 0.001875482
             -0.037715592 0.028394008 -0.0435888242 -0.106272097
## perimeter mean
## compactness_mean -0.011786637 -0.004903894 -0.0453031106 0.043145968
## dimension mean 0.044767906 -0.112784169 -0.0410588768 0.306499872
## smoothness_se
             0.232745603 -0.330867850 -0.0678348099 -0.260833914
## compactness_se -0.280298048 0.066788120 0.02222220211 0.021001944
## concavity_se
             -0.354164595 0.049699104 0.0336810725 -0.219193299
## points se
             -0.195758558 -0.023197526 -0.0378517870 -0.370217167
## symmetry_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
            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
## compactness mean 0.099214048 -0.19518602 0.005172841 0.307036205
## concavity_mean 0.075750464 0.03395563 0.134664686 -0.124553100
## points_mean
             0.116569072 -0.14261678 0.006124860 0.071564686
## compactness_se -0.148593714 -0.11518343 -0.038615749 0.206959272
## compactness_worst -0.111218083 -0.06445290 0.175401648 0.196805544
## concavity_worst -0.035467377 0.19661986 0.295581609 -0.184959562
## dimension worst -0.087222442 -0.11291399 -0.007071634 -0.118625115
##
                  PC13 PC14 PC15 PC16
## id
            -0.004841084 -0.006500099 0.006885943 -0.002753492
0.065047550 -0.067879244 -0.010152279 -0.014636050
## area_mean
## smoothness mean 0.315872261 -0.046461624 -0.444044654 0.117493291
```

```
## 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
## dimension_mean
               0.236120382 -0.106390748 0.377436108 -0.518333769
## compactness_se -0.263398426 -0.010320713 -0.491903153 -0.167886977
## 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 FC17
-0.007779983 -0.019707372 0.005442248 0.020454908
##
## id
               0.150008977 0.209908003 -0.156773206 0.211821385
## radius_mean
## 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
## symmetry_mean
              0.165691481 -0.194702559 -0.169168737 -0.058503329
## dimension mean 0.039119713 0.046298986 -0.086779501 -0.062879947
## 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
##
                 PC21 PC22 PC23 PC24
             PC21 PC22 PC23 PC24
0.009870917 0.006195707 0.003190337 -0.010289027
## id
## 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
```

```
"" arca_bc
                  0.2/00/3010 0.20/310111 0.000/10/30
## smoothness_se -0.095370809 0.066602974 -0.051852247 -0.057154068
## 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 -0.183664583 0.01859418 0.128713229 0.131697326
## texture_mean 0.099441545 -0.08442059 0.024821224 0.017622634
##
                 PC25 PC26 PC27 PC28
## perimeter_mean -0.117262178 -0.02743488 0.124670225 0.115650274
## area_mean 0.070557041 0.21057100 -0.361014547 -0.467489167
## compactness_mean -0.102198309 -0.39651346 -0.262695425 -0.098624638
## points_mean 0.082349955 0.18630114 -0.389316679 0.453345398
## symmetry_mean 0.018841491 0.02451053 0.015910368 0.015157593
## dimension_mean -0.134601525 0.20670502 0.096796804 0.101343150
## smoothness se 0.016193934 -0.03100551 0.008066566 0.009312365
## compactness_se -0.122457873 -0.17364984 0.049404535 -0.046651501
## concavity_se
                 0.186159613 -0.01600952 -0.091931364 0.083824645
## points_se
## symmetry_se
## dimension_se
## radius_worst
                 -0.107166573 0.12999049 0.018674110 0.011675700
                 0.002613811 0.01936313 0.016991197 0.019891112
0.076177800 0.08458109 -0.035156906 0.012141785
                 -0.158114412 -0.07144112 0.195812320 0.178796461
## texture worst -0.118609952 0.11802219 -0.036347107 -0.021473842
## 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
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
## radius_se -1.922290e-01 -0.1186768422 8.679321e-03
## texture_se
                -5.624974e-03 0.0086942153 1.063104e-03
## 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
                 -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
## radius_worst
                 4.126296e-01 0.6356796555 1.356846e-01
## texture worst
                 -3.896988e-04 -0.0172219636 -1.020237e-03
```

```
## 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
##
## $center
##
               id
                       radius_mean
                                      texture_mean
                                                    perimeter_mean
##
      3.037183e+07
                      1.412729e+01
                                     1.928965e+01
                                                     9.196903e+01
##
        area mean smoothness mean compactness mean concavity mean
      6.548891e+02
                     9.636028e-02 1.043410e-01
                                                     8.879932e-02
##
      points mean symmetry mean dimension mean
                                                       radius se
##
      4.891915e-02
                     1.811619e-01 6.279761e-02
                                                     4.051721e-01
##
       texture se
                     perimeter se
                                       area se smoothness se
                                     4.033708e+01
                                                     7.040979e-03
##
      1.216853e+00
                     2.866059e+00
                                                       symmetry_se
##
                     concavity se
                                       points se
    compactness se
##
      2.547814e-02
                      3.189372e-02
                                      1.179614e-02
                                                      2.054230e-02
                                   texture_worst perimeter_worst
##
      dimension_se
                      radius_worst
##
      3.794904e-03
                      1.626919e+01
                                      2.567722e+01
                                                     1.072612e+02
\#\,\#
       area_worst smoothness_worst compactness_worst
                                                    concavity_worst
                                                     2.721885e-01
##
      8.805831e+02
                   1.323686e-01 2.542650e-01
                   symmetry worst dimension worst
##
      points worst
##
      1.146062e-01
                     2.900756e-01
                                     8.394582e-02
##
## $scale
               id
##
                      radius mean
                                     texture mean
                                                   perimeter mean
##
      1.250206e+08
                      3.524049e+00
                                     4.301036e+00
                                                     2.429898e+01
        area_mean smoothness_mean compactness_mean
##
                                                   concavity_mean
                   1.406413e-02
                                                   7.971981e-02
                                   5.281276e-02
##
      3.519141e+02
##
                                                         radius se
       points mean
                     symmetry_mean
                                    dimension mean
                    2.741428e-02
                                   7.060363e-03
                                                     2.773127e-01
##
       3.880284e-02
                     perimeter_se
##
       texture_se
                                         area_se
                                                     smoothness_se
                                                     3.002518e-03
##
      5.516484e-01
                      2.021855e+00
                                      4.549101e+01
                     concavity_se
##
    compactness_se
                                       points_se
                                                      symmetry_se
                     3.018606e-02
                                     6.170285e-03
                                                     8.266372e-03
##
     1.790818e-02
##
     dimension se
                     radius worst
                                     texture worst perimeter worst
##
     2.646071e-03 4.833242e+00 6.146258e+00 3.360254e+01
##
       area_worst smoothness_worst compactness_worst concavity_worst
\# \#
      5.693570e+02 2.283243e-02 1.573365e-01
                                                     2.086243e-01
##
     points worst symmetry worst dimension worst
                      6.186747e-02 1.806127e-02
##
      6.573234e-02
##
## $x
##
                             PC2
                                         PC3
                                                    PC4
##
    [1,]
         2.50194550 -0.096948052 -0.4489597008 2.334117555 0.697715477
          1.46743888 -1.686300588 1.1542038774 0.336210935 0.459625382
\# \#
    [2,]
          2.92902790 -0.383199241 -0.8955890691 -0.116482776 0.984413772
##
    [3,1
          1.99534236 -1.330465924 1.1172875551 2.050276131 0.253038461
##
    [4,]
         2.50025164 2.010350968 -0.7584034865 1.986216853 -1.135370958
##
    [5,1
##
   [6,] 2.01830827 -0.782420945 0.1125197163 -0.653227961 0.018415769
   [7,] 1.09348444 -1.293936715 1.4189069050 -3.356740242 -0.951965047
##
   [8,] -4.54183732 -3.099372405 -2.2612175075 -2.200950296 3.459973177
##
   [9,] 2.69412302 -1.421092569 0.9516155379 0.007476733 1.250335936
##
   [10,]
         2.08173487 -1.812218651 -1.1720406658 1.137014305 0.414757427
##
   [11,] -5.08416948 2.000137924 -0.7512306390 -0.076186113 -0.385543991
   [12,]
##
         2.42030088 -0.010337119 -0.8066693631 -0.070644793 0.075814497
##
   [13,]
          0.42109190 - 0.121776139 - 0.5410059634 - 0.365041743 - 1.511092195
          3.72127373
                     1.784296564 1.2485358731 0.129289545 0.520260616
##
   [14.]
         -2.50683608 2.519115720 0.7679362671 -0.814023090 0.426551810
##
   [15,]
         ##
   [16,]
         0.71352261 1.556166034 -0.7062636839 1.691564007 -0.340941385
##
   [17.]
## [18,]
         ## [19,] -7.18617681 -0.044244030 2.0258549210 0.087485348 0.664035557
## [20,] 1.88203558 -1.429919420 -1.0832586631 1.705943528 0.884175937
## [21,] 2.83789903 0.393325503 -0.3614479874 0.812880484 0.040063252
## [22,] -0.33608363 -3.535859467 0.0806908813 -1.552997026 1.956881414
## [23,] 2.34862913 -0.683708564 1.1197182845 -0.295906284 0.539770064
   [24,] -1.40123647 -1.312422336 0.6099807955 1.894731466 0.034197398
##
##
   [25,] 1.35378261 -3.553367632 1.7257338713 -0.038167893 0.343071810
   [26,]
         0.36757240 -0.131511726 -2.0517854155 1.050460940 -0.752865683
```

```
[27,]
          1.96773313 0.957849703 -0.1042456721 -0.088392018 0.547006886
##
##
   [28,]
         -3.99611352 -0.554685751 -2.7691318844 -1.904122982 0.524809460
          1.94377603 -2.549019049 -1.0010501923 0.023504095 -1.204979082
##
   [29.]
          3.14007439
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## [229,] 0.150104370 -3.349342e-01 -1.184740e-02 -2.692893e-02
## [230,] -0.261499163 -1.385273e-01 7.101842e-02 -4.465156e-02
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## [438,] 0.064842228 -1.767356e-01 1.073602e-01 -3.916015e-02
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## [440,] 0.211675368 2.179147e-02 -1.482077e-01 2.622333e-01
## [441,] -0.063261860 1.408576e-01 1.293660e-01 1.780952e-01
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## [446,] -0.209186142 2.148137e-01 3.565643e-02 5.023195e-02
## [447,] 0.250877405 5.162165e-02 -1.712514e-01 4.138032e-02
## [448,] -0.221909881 -5.656071e-02 9.059298e-02 6.376507e-03
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## [453,] 0.214432386 6.462198e-01 4.572245e-01 -1.856783e-01
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## [455,] -0.305309372 -4.839912e-02 -1.447690e-02 -2.087116e-01
## [456,] 0.062926245 -3.128287e-02 1.289289e-02 1.772915e-02
## [457,] 0.591158273 -5.824376e-02 -1.919702e-01
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## [458,] -1.238236947 2.562185e-01 -7.671496e-01
## [459,] 0.101701285 -8.905947e-03 -1.834538e-01 1.857571e-01
## [460,] -0.027289191 -1.938181e-01 1.048415e-01 3.829356e-02
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## [462,] -0.193424611 1.479751e-01 1.220193e-01 -1.296267e-01
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## [465,] 0.284431074 -5.373005e-01 -2.197313e-01 9.126616e-02
## [466,] -0.387817362 3.272425e-01 3.568318e-02 -2.214587e-01
## [467,] -0.002403087 2.411783e-01 2.350863e-01 -5.933900e-02
## [468,] -0.007050347 2.245577e-01 -2.302748e-02 -1.105619e-01
## [469,] 0.135831371 -1.488714e-01 -7.360296e-02 -1.879613e-04
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## [475,] 0.490390249 4.206388e-02 1.444518e-03 3.732401e-01
## [476,] 0.194558951 1.065368e-01 -2.703042e-01 -1.832076e-01
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## [478,] -0.131892905 -1.728015e-01 -3.374861e-01 -2.453675e-02
## [479,] 0.237160597 -7.589834e-02 1.284184e-01 -3.049761e-01
## [480,] 0.278073334 7.642517e-02 7.834827e-02 -4.114467e-01
## [481,] 0.035489310 7.554027e-02 1.064340e-01 2.369870e-02
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## [483,]
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          0.207366238 -5.127782e-01 -3.057416e-01 -5.080697e-02
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## [486,]
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## [17,] -6.569926e-02 -1.493094e-01 1.716667e-02 0.0597777408
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## [568,] 3.274830e-04 0.1276164883 -1.587248e-02 -6.099950e-02
## [569,] -1.465535e-01 -0.0395651763 2.851123e-02 -1.446121e-01
##
                  PC28
                               PC29
                                            PC30
    [1,] 0.0535613058 0.0151848822 1.598541e-02 1.396101e-03
##
    [2,] -0.1869655269  0.0270113107 -8.033300e-04  8.096490e-03
##
    [3,] -0.0765306661 -0.0146403879 1.030789e-02 9.074601e-03
##
    [4,] -0.0171166487 -0.0478284944 2.386299e-02 2.650750e-04
##
    [5,] 0.0383536434 0.0324507997 -2.312178e-03 -2.563269e-03
##
    [6,] 0.0095543400 -0.0044034309 3.869919e-03 -2.931194e-03
##
    [7,] 0.0378787886 0.0203768196 4.010932e-04 -1.269166e-02
##
    [8,] 0.1590301296 0.0211851026 2.063600e-02 -1.136547e-03
    [9,] -0.0586366278 -0.0157101706 1.578442e-02 8.784686e-03
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   [10,] 0.0208900863 -0.0229256417 -3.656928e-02 3.436465e-03
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   [11,] -0.1836996047 -0.0702399725 -1.960853e-02 1.499158e-02
##
##
   [12,] 0.0313667590 0.0214758148 -5.815388e-03 2.101407e-03
   [13,] -0.0115189160 -0.0236487859 -1.336927e-02 -9.541018e-04
   [15,] 0.0526092277 -0.0106431857 1.382817e-02 -1.138219e-02
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   [18,] 0.0091836992 0.0184535080 3.004816e-03 -8.284949e-03
##
  [19,] 0.2263597091 0.0319045180 -5.109774e-02 1.888498e-02
## [20,] 0.0030067774 -0.0211027308 -1.938230e-02 9.852988e-04
   [21,] -0.0198932675 -0.0136803305 -8.551012e-03 -1.325546e-03
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   [26,] 0.0315139751 -0.0401102951 4.734871e-02 9.919702e-03
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  [34,] -0.0590833361  0.0130416480  7.452318e-03  9.309105e-03
  [35,] -0.0252578940 0.0047644298 1.129211e-02 -7.572504e-04
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  [37,] 0.0145404352 -0.0403402638 4.594615e-02 4.646818e-05
   [38,] 0.0620363491 0.0259186233 -1.038252e-02 1.782104e-02
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   [39,] 0.0551020192 0.0091668531 1.530145e-02 1.309894e-02
##
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   [41,] -0.0507089905 -0.0501071660 -7.727756e-03 -1.182598e-02
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   ##
   [44,]
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        0.0328784540 -0.0146923196 -7.281896e-03 -5.593681e-03
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  [48,] -0.2137482100 -0.0381373067 -2.763514e-02 4.146805e-03
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   [50,] 0.0129741769 0.0087123575 -7.947914e-03 -1.263665e-02
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   [51,] 0.0815819893 0.0152921924 -6.954635e-03 -2.622044e-03
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   [53,] -0.0402841172  0.0107722059 -1.137519e-02  3.710160e-03
   [54,] 0.0411842454 -0.0065017458 2.731969e-03 -7.122045e-03
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        0.0025384196 -0.0130582355 -1.408453e-02 3.403444e-03
   [98,]
  [99,] -0.0327142984 -0.0275644972 1.513022e-03 -7.353861e-04
## [100,] -0.2069422358 -0.0652489816 2.144969e-02 -6.508857e-03
## [101,] -0.1105311481 -0.0013877012 1.828318e-02 -9.785484e-03
## [102,] 0.0483677669 -0.0459407868 1.111969e-02 3.880046e-03
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## [105,] 0.0467342886 0.0108345237 -3.142075e-02 1.271146e-03
## [106,] -0.0413517840 -0.0235905790 -2.795827e-02 3.569201e-03
## [107,] 0.0038392754 0.0547225235 -1.913600e-03 -2.788389e-03
## [108,] -0.0547154320 -0.0165025183 1.460897e-02 2.125402e-03
## [109,] -0.0339964747 0.0000718563 2.155622e-02 4.147422e-03
## [110,] -0.0774286268 -0.0086110360 -2.290418e-04 1.782624e-03
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## [125,]
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## [127,] -0.0494097662  0.0115558742  1.951383e-02 -2.997311e-03
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## [141,] -0.0740378147 -0.0175467891 -2.181434e-02 8.579655e-03
## [142,] 0.2529700286 -0.0338815826 -4.553172e-02 -4.712428e-02
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## [149,] 0.1056232897 -0.0076881080 5.237851e-03 -4.690007e-03
## [150,] 0.0155594355 0.0230259878 -2.954818e-02 -2.359282e-03
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## [155,] 0.0423785082 -0.1189758996 -1.719611e-02 1.501597e-02
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## [157,] 0.0150952486 -0.0038538689 2.906355e-02 -3.245762e-03
## [158,] 0.0037113226 -0.0488170641 -2.437811e-03 -2.296581e-03
## [159,] 0.0547616829 0.0188567292 4.834882e-02 4.374965e-03
## [160,] -0.0326339555 -0.0084983195 1.205413e-02 2.884325e-03
## [161,] -0.0211644709 0.0026083217 6.400260e-02 1.161854e-02
## [162,] 0.1029920697 -0.1320359748 2.703355e-02 1.442639e-02
## [163,] 0.0324474322 0.0041253801 -1.291301e-02 -2.752354e-03
## [164,] 0.0700441492 -0.0250198232 3.430259e-03 8.358832e-03
## [165,] -0.2638728581 -0.0789245094 2.625865e-02 -7.628811e-03
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## [180,] -0.0341987777 0.0002996804 -1.240101e-02 3.102445e-04
## [181,] 0.0409088669 0.0165848011 2.757407e-02 6.055236e-03
## [182,] 0.0957711588 0.0256512152 9.719448e-04 5.484105e-04
## [183,] 0.0034861498 0.0042715325 -2.109341e-02 -2.514913e-03
## [184,] 0.0182635901 -0.0014884541 -1.693941e-02 1.576631e-02
## [185,] -0.0153716350 -0.0019545732 -1.360106e-02 9.613744e-03
## [186,] 0.0262076696 -0.0114572595 2.606248e-02 1.052111e-03
## [187,] -0.0682213414 -0.0050720262 -3.422509e-02 -1.011332e-04
## [188,] 0.0390749887 0.0175751718 -1.622265e-02 1.862621e-03
## [189,] -0.0593689344 -0.0186617141 2.093184e-02 -5.483462e-03
## [191,] 0.0937831254 0.0057336425 5.356388e-03 5.899756e-03
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## [198,] 0.0116182552 0.0025851687 -9.083080e-03 8.212111e-03
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## [203,] -0.0607609902 0.0155759941 -2.345218e-02 9.257839e-03
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## [209,] -0.0144120095 0.0083469672 2.166718e-02 -2.414526e-03
## [210,] -0.0110996987 0.1370702906 -1.603930e-02 4.133387e-03
## [211,] -0.0381426451 0.0171359015 6.290327e-03 1.075495e-02
## [214,] 0.0652718210 -0.0111549889 -1.407231e-02 -8.327849e-03
## [215,] -0.0408947713 -0.0166146907 7.978440e-03 -4.868288e-04
## [217,] -0.0157885793 -0.0016117964 -7.704393e-04 -8.937990e-04
## [218,] -0.0705155682 -0.0476980872 7.135127e-03 9.321918e-04
## [219,] 0.0845005649 0.0007302365 1.973105e-02 3.460458e-03
## [220,] -0.3021173359 -0.0233926389 8.616636e-03 -1.816951e-03
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## [227,] 0.3014725649 0.0096765848 2.010773e-02 -4.149831e-02
## [228,] 0.0565387128 -0.0057568519 -5.965070e-03 3.186364e-03
## [229,] -0.0016450638 -0.0574026217 9.502027e-03 -6.027354e-04
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## [241,] -0.0573744623 -0.0291055063 8.163172e-03 6.137371e-03
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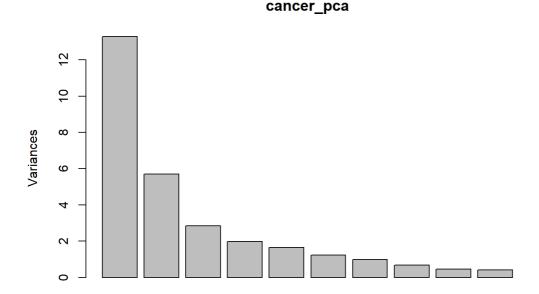
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## [484,] -0.0090227346  0.0248997990  3.734145e-02  1.790719e-03
## [485,] -0.0125412778  0.0127037430  1.009832e-02  4.256908e-03
## [486,] 0.0636391030 -0.0323824607 -2.607947e-02 -4.951705e-03
## [487,] -0.1397578741 0.0088616024 -6.041273e-04 1.252251e-02
## [488,] -0.0747773145 -0.0420757978 -5.595981e-03 -1.960175e-02
## [489,] 0.0802388777 -0.0138229184 -2.376301e-02 -2.479727e-03
## [491,] 0.0989654491 -0.0140864165 4.367477e-03 2.171240e-03
         0.0282618964 -0.0680236124 8.422818e-02 -2.374215e-02
## [492,]
## [493,] 0.2172752231 0.0082839891 1.938583e-02 1.442880e-03
## [494,] -0.0123190204 -0.0095134802 -9.186819e-03 3.113970e-03
## [495,] -0.2199572217 -0.0294567698 1.576101e-02 -5.252938e-03
## [496,] 0.0634010466 -0.0082540868 1.922416e-02 -1.087247e-02
## [497,] -0.0533707600 -0.0089322195 3.592632e-03 -3.646989e-03
## [498,] 0.0480566608 0.0142425925 7.108301e-02 1.083214e-02
## [499,] 0.1442753476 -0.0422424796 -1.782818e-02 4.751815e-02
## [500,] 0.0165308656 -0.0083255759 -2.727051e-02 5.957476e-03
## [501,] -0.0100329601 -0.0372407659 2.363354e-02 -9.646078e-03
## [502,] 0.0171915088 -0.0178647571 1.050801e-03 -8.568039e-03
## [503,] 0.0756534229 0.0607336581 9.157892e-03 1.261060e-02
## [504,] 0.0297769033 0.0973985283 3.350919e-02 -2.700494e-02
## [505,] 0.0350448873 0.0275088283 1.063655e-02 2.760851e-05
## [506,] -0.0285875298 -0.0016575587 -1.182673e-03 -4.053176e-03
## [507,] 0.0189829021 0.0207446755 -7.102198e-03 -1.406710e-02
## [508,] -0.0172826029 0.0125580405 -9.707108e-03 -9.199495e-03
## [509,] -0.1441724544 -0.0394471024 -6.987832e-02 -7.615465e-03
## [510,] 0.0875496770 -0.0391355476 8.573809e-03 8.449619e-03
## [511,] 0.0981611536 -0.0150404803 1.875760e-02 1.130911e-02
## [512,] 0.0260553682 -0.0198184704 2.964767e-02 -1.229652e-02
## [513,] 0.2306612233 -0.1768569002 7.487201e-02 1.145739e-03
## [514,] 0.0443660379 -0.0127577223 1.385273e-02 5.862815e-04
## [515,] 0.0244686942 0.0136171599 -3.015268e-02 3.004115e-02
## [516,] -0.0464406428 -0.0461814081 7.006575e-03 -2.118240e-02
## [517,] -0.1804905504 0.0326011719 5.896892e-03 -1.843209e-03
## [519,] 0.0216068656 -0.0476515075 -3.416270e-02 -1.304970e-02
## [520,] 0.1523810055 -0.0451309386 -3.033061e-02 1.749013e-02
## [521,] -0.0591849335 -0.0255713138 1.596715e-02 -1.397577e-04
## [522,] -0.0395616001 0.0091315682 3.740507e-02 1.008395e-02
## [523,] 0.0513025394 0.0081396144 -1.686257e-03 -1.284561e-03
## [524,] -0.1277410469 -0.0095049270 6.099755e-02 -1.943908e-02
## [525,] -0.1521848402 -0.0408818910 4.394310e-02 1.438141e-03
## [527,] -0.0502031636  0.0248580682 -2.886409e-02  1.869812e-02
## [528,] -0.0157216742 0.0417373451 6.411718e-03 -3.945196e-02
## [529,] 0.0226840232 -0.0011725978 1.220802e-02 2.561845e-04
```

```
## [531,] 0.0142770079 0.0250853629 -3.624568e-03 -3.363374e-03
## [532,] -0.0065668831 0.0016770508 5.785196e-02 -3.674048e-02
         0.0776102509 0.0225164413 -5.199929e-03 -1.656424e-03
         0.0328960130 -0.0287557957 -4.530040e-02 -6.688662e-03
## [536,] 0.1253022047 0.0006555599 1.231382e-03 4.937076e-03
## [537,] -0.0282099570 0.0141624831 -2.623194e-03 1.917073e-03
## [538,] 0.0033372312 -0.0091753971 -4.953264e-03 6.203430e-04
## [539,] 0.0828029895 -0.0089573807 -1.291559e-02 -4.762266e-03
## [540,] 0.1346756425 0.0092818155 -4.752541e-03 3.466008e-03
## [541,] 0.0686944450 -0.0043570810 1.903166e-02 -1.182699e-02
## [542,] -0.0958839897 -0.0266441447 3.100822e-02 5.635812e-03
## [543,] 0.0391931001 0.0061086959 -2.330714e-03 -6.398906e-06
## [544,] -0.0884505800 0.0230539154 -5.910469e-02 -6.112608e-03
## [545,] 0.0461121057 0.0512587241 -3.534171e-02 7.330315e-02
## [546,] -0.0181344013 -0.0079308291 -5.151052e-02 -9.360687e-03
## [547,] -0.1284012027 0.0731570237 2.171499e-02 3.565962e-03
## [548,] -0.0652554774  0.0267267995 -1.453407e-02  1.372139e-03
## [549,] 0.0144803135 0.0290173096 -1.418995e-03 -4.295967e-03
## [550,] 0.0020030080 0.0038173956 1.280350e-02 8.471746e-04
## [551,] -0.0460309869 -0.0698233075 2.414560e-04 1.845563e-03
## [553,] 0.0013530758 0.1356038629 -3.808346e-02 3.787346e-03
## [554,] -0.0092761156 -0.0162335677 -6.313714e-03 8.823042e-04
## [555,] 0.0583525074 0.0104867651 -7.650307e-03 -7.688319e-03
## [556,] 0.0365396314 0.1285384513 -1.926274e-02 -3.420722e-03
## [557,] -0.0825366557 0.0120057039 -9.118959e-04 2.230323e-03
## [558,] 0.0596863336 0.0432872298 -3.428184e-03 -7.399596e-03
## [559,] 0.2118838404 -0.0181142975 3.600294e-02 2.217638e-04
## [561,] 0.0612930701 -0.1122543797 -6.714921e-02 -1.368662e-02
## [562,] 0.0536497985 -0.0177312339 -1.023821e-02 -3.139075e-03
## [563,] 0.0572798895 0.0130302469 -1.209907e-02 -5.552985e-03
## [564,] -0.0318972163 -0.0229019521 -3.253642e-02 -9.340876e-04
## [565,] 0.0048045501 0.0281111851 4.728344e-03 -5.899725e-04
## [566,] 0.0068911201 -0.0175118860 -2.580188e-04 5.826148e-03
## [567,] 0.0471706437 0.0507946253 1.250840e-03 3.425887e-03
## [568,] 0.1477176175 0.0013794388 -1.939084e-03 -1.131610e-02
## [569,] -0.0426995400 -0.0728970801 3.245528e-02 -6.532395e-03
```

plot(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
                     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 PC24
## 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(cancer_pca)

head(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
                           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
## [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
##
            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
# sample scores stored in cancer pca$x
# singular values (square roots of eigenvalues) stored in cancer pca$sdev
# loadings (eigenvectors) are stored in cancer_pca$rotation
# variable means stored in cancer pca$center
# variable standard deviations stored in sparrows_pca$scale
# A table containing eigenvalues and %'s accounted, follows
# Eigenvalues are sdev^2
(eigen cancer <- cancer pca$sdev^2) ## brackets for print
## [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_cancer) <- paste("PC",1:31,sep="")
eigen_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
                                             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
```

```
sumlambdas <- sum(eigen_cancer)
sumlambdas</pre>
```

```
## [1] 31
```

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

```
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
## 4.291620e-06
```

summary(eigen cancer)

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

summary(cancer_pca)

```
## Importance of components:
##
                          PC1
                                PC2
                                       PC3
                                                PC4
                      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
```

```
cumvar_cancer <- cumsum(propvar)
cumvar_cancer</pre>
```

```
## PC1 PC2 PC3 PC4 PC5 PC6 PC7
## 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_cancer,propvar,cumvar_cancer)
rownames(matlambdas) <- c("Eigenvalues","Prop. variance","Cum. prop. variance")
# Sample scores stored in cancer_pca$x
head(cancer_pca$x)</pre>
```

```
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
## [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),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
          В 2.929028 -0.38319924 -0.8955891 -0.1164828 0.98441377
## 3
## 4
          B 1.995342 -1.33046592 1.1172876 2.0502761 0.25303846
## 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
##
        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
```

head(round(matlambdas, 4))

```
##
                        PC1
                              PC2
                                    PC3
                                           PC4
                                                 PC5
                                                         PC6
                                                                PC7
                   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
## Eigenvalues
                    0.6718 0.4608 0.4031 0.3492 0.2939 0.2611 0.2413
                   0.0217 0.0149 0.0130 0.0113 0.0095 0.0084 0.0078
## Prop. variance
## Cum. prop. variance 0.9140 0.9289 0.9419 0.9532 0.9626 0.9711 0.9789
##
                      PC15 PC16 PC17 PC18 PC19 PC20 PC21
                    0.1570 0.0941 0.0798 0.0590 0.0526 0.0491 0.0311
## Eigenvalues
                   0.0051 0.0030 0.0026 0.0019 0.0017 0.0016 0.0010
## Prop. variance
## 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
##
                     0.0016 7e-04 1e-04
## Eigenvalues
                   0.0001 0e+00 0e+00
## Prop. variance
## Cum. prop. variance 1.0000 1e+00 1e+00
```

```
## 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
```

head(cancer_pca\$rotation)

```
PC1 PC2 PC3
               -0.02291216 0.03406849 0.09693844 -0.02659805
## id
             -0.21891302 0.23327140 -0.01139379 0.04218795
## radius_mean
## texture_mean -0.10384388 0.06004420 0.06689234 -0.60295431
## perimeter_mean -0.22753491 0.21458900 -0.01212479 0.04275280
## area mean -0.22104577 0.23066882 0.02629315 0.05411472
## smoothness mean -0.14241471 -0.18642221 -0.10318240 0.15809818
##
                     PC5 PC6 PC7 PC8
## id
               0.01132759 -0.316733438 0.90711563 -0.096362415
## radius_mean -0.03812986 0.029588521 -0.04229878 -0.116427419
## perimeter mean -0.03771559 0.028394008 -0.04358882 -0.106272097
## area_mean -0.01056223 0.006113155 -0.02892567 -0.047414568
## smoothness mean 0.36575006 -0.262508993 -0.14034036 -0.123541189
##
                PC9 PC10 PC11 PC12
                0.14911564 -0.16926751 0.05818900 -0.006721252
## id
## radius_mean -0.04627083 -0.22402704 -0.07946608 -0.042213788
## texture_mean -0.08872717 0.11945674 -0.25325809 0.304032359
## perimeter_mean -0.03623074 -0.22634517 -0.06986593 -0.017573055
## area mean -0.08064986 -0.18600385 -0.06279537 -0.110760120
## smoothness mean 0.27899640 -0.06133822 0.08466155 0.135321954
      PC13 PC14 PC15 PC16
##
## id
              -0.004841084 -0.006500099 0.006885943 -0.002753492
## perimeter_mean 0.038470392 -0.044684430 -0.048019221 0.039590476
                0.065047550 -0.067879244 -0.010152279 -0.014636050
## area mean
## smoothness_mean 0.315872261 -0.046461624 -0.444044654 0.117493291
##
                PC17 PC18 PC19 PC20
               -0.007779983 -0.01970737 0.005442248 0.02045491
## id
## perimeter_mean   0.113792993   0.20123366 -0.168413120   0.22707927
## area mean 0.130173978 0.25146046 -0.269145594 -0.04549963
## smoothness_mean 0.203117911 0.16817161 0.354463321 -0.16035826
      PC21 PC22 PC23 PC24
0.009870917 0.006195707 0.003190337 -0.010289027
##
## id
               0.009870917 0.006195707 0.003190337 -0.010289027
## 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
\# \#
                PC25
                           PC26
                                      PC27 PC28
## id
               -0.004233388 -0.00132610 -0.002571324 -0.001623875
## radius mean -0.183664583 0.01859418 0.128713229 0.131697326
## texture mean 0.099441545 -0.08442059 0.024821224 0.017622634
## 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
     PC29 PC30 PC31
-1.891724e-05 -0.0006852263 -7 1225816-05
##
              -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
```

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

```
## diagnosis PC1 PC2 PC3 PC4 PC5
## 1 B 2.204253 -0.3436398 0.2160542 0.1384470 -0.09800974
        M -3.711879 0.5786765 -0.3638272 -0.2331395 0.16504470
## 2
        PC6 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
        PC11 PC12 PC13 PC14 PC15
## 2 0.0013284090 -0.010133446 -0.005565970 0.06292076 0.04131016
##
    PC16 PC17 PC18 PC19 PC20
## 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
##
    PC26 PC27
                     PC28 PC29
## 1 -0.000879209 -0.003967725 -0.001909259 -0.0003418423 -0.0009838392
## 2 0.001480555 0.006681499 0.003215121 0.0005756496 0.0016567480
```

tabmeansPC <- tabmeansPC[rev(order(tabmeansPC\$diagnosis)),]
tabmeansPC</pre>

```
## 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
                                              PC10
## 2 -0.007364189 -0.02848926 -0.08261104 -0.05686895 0.01688516
## 1 0.004373132 0.01691799 0.04905754 0.03377092 -0.01002704
        PC11 PC12 PC13 PC14 PC15
## 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 PC20
## 2 0.04238026 -0.0001874111 -0.010278530 0.02202108 0.016391505
## PC21 PC22 PC23 PC24 PC25
## 2 0.014077577 0.010482771 0.005093974 -0.004394179 -0.013157604
## 1 -0.008359794 -0.006225063 -0.003024993 0.002609428 0.007813479
    PC26 PC27 PC28 PC29 PC30
##
## 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
        0.0066814986 -0.0039677247
## PC27
## 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
       0.5786764540 -0.3436397990
## PC2
## 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
```

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

```
##
                 В
## 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC1 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC2 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC3 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC4 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC5 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC6 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC7 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC8 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC9 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC10 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC11 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC12 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC13 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC14 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC15 by 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~cancer\$diagnosis,data=diag_pca)

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

```
##
## Welch Two Sample t-test
##
## data: PC17 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC18 by cancer$diagnosis
## t = 0.67787, df = 288.75, p-value = 0.4984
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03118389  0.06394850
## sample estimates:
## mean in group B mean in group M
## 0.006103777  -0.010278530
```

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

```
##
## Welch Two Sample t-test
##
## data: PC19 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC20 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC21 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC22 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC23 by 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~cancer\$diagnosis,data=diag_pca)

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

```
##
## Welch Two Sample t-test
##
## data: PC25 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC26 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC27 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC28 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC29 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC30 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## Welch Two Sample t-test
##
## data: PC31 by 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~cancer$diagnosis,data=diag_pca)
```

```
##
## F test to compare two variances
##
## data: PC1 by 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
## 0.2931115</pre>
```

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

```
##
## F test to compare two variances
##
## data: PC2 by 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
## 0.5845061
```

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

```
##
## F test to compare two variances
##
## data: PC3 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC4 by 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
## ratio of variances
## 0.9955172
```

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

```
##
## F test to compare two variances
##
## data: PC5 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC6 by 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
## 0.5078212
```

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

```
##
## F test to compare two variances
##
## data: PC7 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC8 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC9 by 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
## ratio of variances
```

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

```
##
## F test to compare two variances
##
## data: PC10 by 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
## 0.6503607
```

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

```
##
## F test to compare two variances
##
## data: PC11 by 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
## 1.110734
```

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

```
##
## F test to compare two variances
##
## data: PC12 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC13 by 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
## 1.008108
```

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

```
##
## F test to compare two variances
##
## data: PC14 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC15 by 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
## 0.8268605
```

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

```
##
## F test to compare two variances
##
## data: PC16 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC17 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC18 by 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
## 0.3050177</pre>
```

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

```
##
## F test to compare two variances
##
## data: PC19 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC20 by 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
## ratio of variances
## 0.3614279</pre>
```

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

```
##
## F test to compare two variances
##
## data: PC21 by 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
## ratio of variances
```

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

```
##
## F test to compare two variances
##
## data: PC22 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC23 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC24 by 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~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC25 by 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
## ratio of variances
## 0.4411905
```

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

```
##
## F test to compare two variances
##
## data: PC26 by 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
## ratio of variances
## 0.3779774
```

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

```
##
## F test to compare two variances
##
## data: PC27 by 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
## 0.3938058
```

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

```
##
## F test to compare two variances
##
## data: PC28 by 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~cancer\$diagnosis,data=diag pca)

```
##
## F test to compare two variances
##
## data: PC29 by 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
## 0.2129389</pre>
```

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

```
##
## F test to compare two variances
##
## data: PC30 by 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</pre>
```

var.test(PC31~cancer\$diagnosis,data=diag_pca)

```
##
## F test to compare two variances
##
## data: PC31 by 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</pre>
```

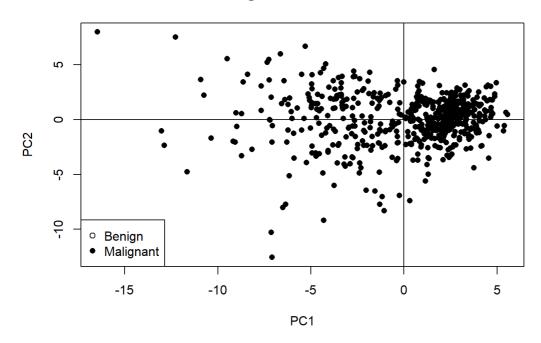
```
# Levene's tests (one-sided)
library(car)
(LTPC1 <- leveneTest(PC1~cancer$diagnosis,data=diag_pca))</pre>
```

```
## 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 '**' 0.05 '.' 0.1 ' ' 1
```

```
(p_PC1_1sided <- LTPC1[[3]][1]/2)</pre>
## [1] 8.268824e-15
(LTPC2 <- leveneTest(PC2~cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 18.786 1.73e-05 ***
        567
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(p_PC2_1sided=LTPC2[[3]][1]/2)
## [1] 8.651441e-06
(LTPC3 <- leveneTest(PC3~cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 27.65 2.063e-07 ***
##
## ---
## 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~cancer$diagnosis,data=diag pca))
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 2e-04 0.989
       567
(p_PC4_1sided <- LTPC4[[3]][1]/2)</pre>
## [1] 0.4944984
(LTPC5 <- leveneTest(PC5~cancer$diagnosis,data=diag_pca))
## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 6.8535 0.009083 **
## 567
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(p_PC5_1sided <- LTPC5[[3]][1]/2)</pre>
## [1] 0.004541533
```

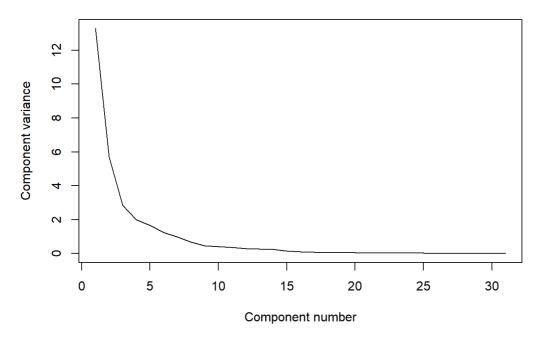
```
# Plotting the scores for the first and second components
plot(diag_pca$PC1, diag_pca$PC2,pch=ifelse(diag_pca$diagnosis == "S",1,16),xlab="PC1", ylab="PC2", main="569
entries against values for PC1 & PC2")
abline(h=0)
abline(v=0)
legend("bottomleft", legend=c("Benign","Malignant"), pch=c(1,16))
```

569 entries against values for PC1 & PC2



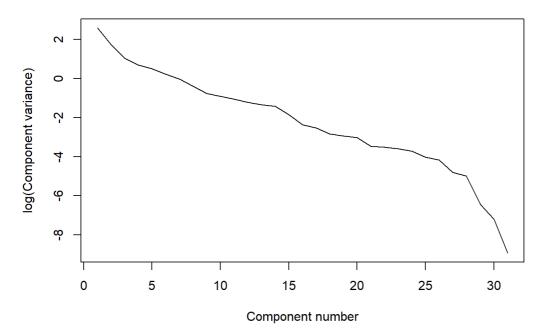
plot(eigen_cancer, xlab = "Component number", ylab = "Component variance", type = "l", main = "Scree diagram
")

Scree diagram



plot(log(eigen_cancer), xlab = "Component number", ylab = "log(Component variance)", type="l", main = "Log(eig
envalue) diagram")

Log(eigenvalue) diagram



```
print(summary(cancer_pca))
```

```
## Importance of components:
                            PC1
                                   PC2
                                          PC3
                                                  PC4
                                                          PC5
## 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
                                          PC9 PC10
##
                                 PC8
                                                         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
                         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
                                 PC20 PC21
##
                          PC19
                                               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
                        0.13431 0.1244 0.09040 0.08305 0.03987 0.02736
## Standard deviation
## 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(cancer_pca)
diag(cov(cancer_pca$x))
```

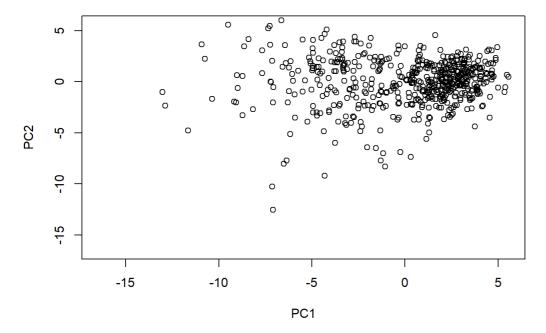
```
PC1
                 PC2
                      PC3 PC4
## 1.328806e+01 5.696805e+00 2.835395e+00 1.981357e+00 1.648815e+00
##
  PC6 PC7 PC8 PC9 PC10
## 1.234673e+00 9.782732e-01 6.717530e-01 4.607924e-01 4.031331e-01
       PC11 PC12 PC13 PC14 PC15
##
## 3.491550e-01 2.938904e-01 2.611469e-01 2.413302e-01 1.569736e-01
       PC16 PC17 PC18 PC19
## 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
```

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

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

head(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
           PC7
                     PC8
                              PC9
                                       PC10
                                                PC11
## [1,]
       0.5092015 \; -1.11423307 \quad 0.2840243 \quad 0.32463197 \; -0.3245353 \quad 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
                                                   PC17
## [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
## [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
## [5,] 0.03835364 0.032450800 -0.002312178 -0.002563269
## [6,] 0.00955434 -0.004403431 0.003869919 -0.002931194
```



```
###
#Factor Analysis
library (psych)
## Attaching package: 'psych'
## The following object is masked from 'package:car':
##
##
       logit
\#\# The following objects are masked from 'package:ggplot2':
##
##
       %+%, alpha
\verb|#install.packages("psych", lib="/Library/Frameworks/R.framework/Versions/3.5/Resources/library")|
library (psych)
fit.pc <- principal(cancer[-2], nfactors=4, rotate="varimax")</pre>
fit.pc
```

```
## Principal Components Analysis
## Call: principal(r = 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
## radius_mean 0.95 0.13 -0.14 0.10 0.951 0.049 1.1 ## texture_mean 0.25 0.06 0.05 0.91 0.897 0.103 1.2
## 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
## symmetry mean
                 0.14 0.58 0.33 -0.06 0.474 0.526 1.7
## 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)

```
## RC1 RC2 RC3 RC4

## [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

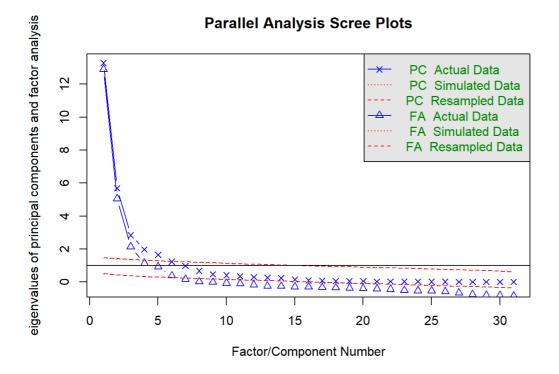
## [4,] -0.3869942 -0.38180634 0.79329588 -1.49911551

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

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

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

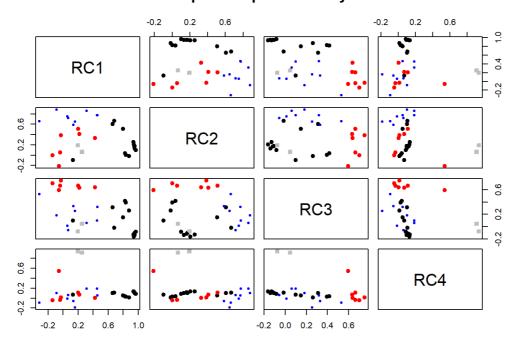
```
## Warning in fa.stats(r = r, f = f, phi = phi, n.obs = n.obs, np.obs
## = np.obs, : The estimated weights for the factor scores are probably
## incorrect. Try a different factor extraction method.
```



```
\#\# 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

