

Problem Statement

Routine breast cancer screening allows the disease to be diagnosed and treated prior to it causing noticeable symptoms. The process of early detection involves examining the breast tissue for abnormal lumps or masses. If a lump is found, a fine-needle aspiration biopsy is performed, which uses a hollow needle to extract a small sample of cells from the mass. A clinician then examines the cells under a microscope to determine whether the mass is likely to be malignant or benign.

If machine learning could automate the identification of cancerous cells, it would provide considerable benefit to the health system. Automated processes are likely to improve the efficiency of the detection process, allowing physicians to spend less time diagnosing and more time treating the disease. An automated screening system might also provide greater detection accuracy by removing the inherently subjective human component from the process.

Apply the k-NN algorithm to perform diagnosis Benign or Malignant

The Dataset

The dataset for the above problem was imported from https://github.com/stedy/Machine-Learning-with-R-datasets/blob/master/wisc_bc_data.csv

s

```
data = read.csv("wisc_bc_data.csv") str(data)
```

```
## 'data.frame':      569 obs. of  32 variables:
## $ id   : int 842302 842517 84300903 84348301 84358402 843786 844359 84458202 8449 ## $ diagnosis      : Factor w/ 2 levels "B","M": 2 2 2 2 2 2 2 2 2 2 ...
## $ radius_mean      : num 18 20.6 19.7 11.4 20.3 ...
## $ texture_mean     : num 10.4 17.8 21.2 20.4 14.3 ...
## $ perimeter_mean   : num 122.8 132.9 130 77.6 135.1 ...
## $ area_mean        : num 1001 1326 1203 386 1297 ...
## $ smoothness_mean  : num 0.1184 0.0847 0.1096 0.1425 0.1003 ... ## $ compactness_mean : num 0.2776 0.0786 0.1599 0.2839 0.1328 ... ## $ concavity_mean   : num 0.3001 0.0869 0.1974 0.2414 0.198 ...
## $ concave.points_mean : num 0.1471 0.0702 0.1279 0.1052 0.1043 ...
## $ symmetry_mean     : num 0.242 0.181 0.207 0.26 0.181 ...
## $ fractal_dimension_mean : num 0.0787 0.0567 0.06 0.0974 0.0588 ...
## $ radius_se        : num 1.095 0.543 0.746 0.496 0.757 ... ## $ texture_se       : num 0.905 0.734 0.787 1.156 0.781 ...
## $ perimeter_se     : num 8.59 3.4 4.58 3.44 5.44 ... ## $ area_se         : num 153.4 74.1 94 27.2 94.4 ...
## $ smoothness_se    : num 0.0064 0.00522 0.00615 0.00911 0.01149 ...
## $ compactness_se   : num 0.049 0.0131 0.0401 0.0746 0.0246 ...
## $ concavity_se     : num 0.0537 0.0186 0.0383 0.0566 0.0569 ...
## $ concave.points_se : num 0.0159 0.0134 0.0206 0.0187 0.0188 ...
## $ symmetry_se      : num 0.03 0.0139 0.0225 0.0596 0.0176 ...
## $ fractal_dimension_se : num 0.00619 0.00353 0.00457 0.00921 0.00511 ...
## $ radius_worst     : num 25.4 25 23.6 14.9 22.5 ...
## $ texture_worst    : num 17.3 23.4 25.5 26.5 16.7 ...
## $ perimeter_worst  : num 184.6 158.8 152.5 98.9 152.2 ...
## $ area_worst       : num 2019 1956 1709 568 1575 ...
## $ smoothness_worst : num 0.162 0.124 0.144 0.21 0.137 ...
```

```
## $ compactness_worst      : num 0.666 0.187 0.424 0.866 0.205 ...
## $ concavity_worst        : num 0.712 0.242 0.45 0.687 0.4 ...
## $ concave.points_worst   : num 0.265 0.186 0.243 0.258 0.163 ...
## $ symmetry_worst         : num 0.46 0.275 0.361 0.664 0.236 ...
## $ fractal_dimension_worst: num 0.1189 0.089 0.0876 0.173 0.0768 ...
```

```
data = data[-1] #Removal of patient indices
```

Including Libraries

The following libraries were needed to be installed and imported for the assignment.

```
library(pca3d) library(class)
library(gmodels) library(ggbiplot)
```

```
## Loading required package: ggplot2
```

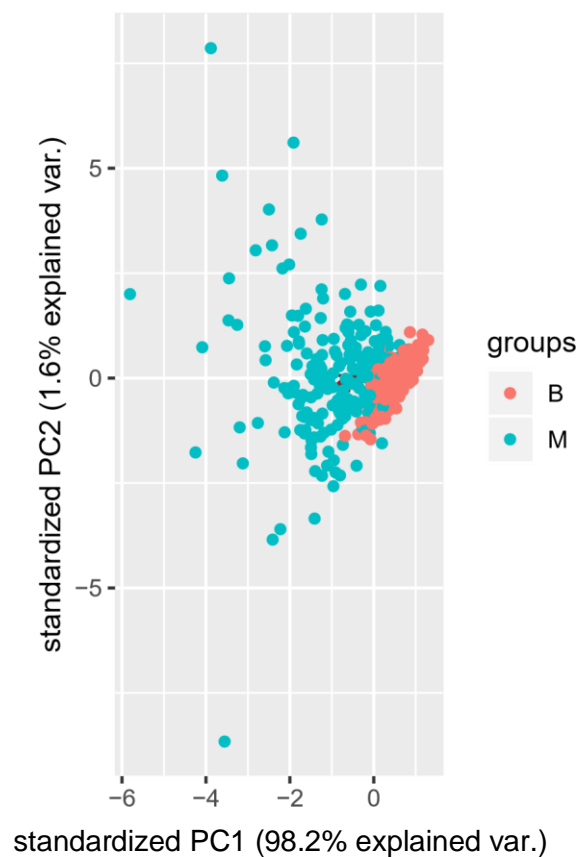
```
## Loading required package: plyr
```

```
## Loading required package: scales
```

```
## Loading required package: grid
```

Principal Component Analysis

```
#pca3d(princomp(data[,c(2:31)]), group = data$diagnosis)
ggbiplot(princomp(data[,c(2:31)]), groups = data$diagnosis, varname.size = 0)
```



The graph above shows the existence of spacial separation in the euclidian space between the two tumour catrgories.

Pre-Processing : Normalization

```
normalize <- function(x) {  
  return ((x - min(x)) / (max(x) - min(x)))  
}
```

```
data_n = as.data.frame(lapply(data[2:31], normalize)) summary(data_n)
```

```
##      radius_mean      texture_mean      perimeter_mean      area_mean  
## Min.      :0.0000    Min.      :0.0000    Min.      :0.0000    Min.      :0.0000  
## 1st Qu.:0.2233      1st Qu.:0.2185      1st Qu.:0.2168      1st Qu.:0.1174  
## Median :0.3024      Median :0.3088      Median :0.2933      Median :0.1729  
## Mean      :0.3382    Mean      :0.3240    Mean      :0.3329    Mean      :0.2169  
## 3rd Qu.:0.4164      3rd Qu.:0.4089      3rd Qu.:0.4168      3rd Qu.:0.2711  
## Max.      :1.0000    Max.      :1.0000    Max.      :1.0000    Max.      :1.0000  
## smoothness_mean compactness_mean concavity_mean      concave.points_mean  
## Min.      :0.0000    Min.      :0.0000    Min.      :0.00000    Min.      :0.0000  
## 1st Qu.:0.3046      1st Qu.:0.1397      1st Qu.:0.06926      1st Qu.:0.1009  
## Median :0.3904      Median :0.2247      Median :0.14419      Median :0.1665  
## Mean      :0.3948    Mean      :0.2606    Mean      :0.20806    Mean      :0.2431  
## 3rd Qu.:0.4755      3rd Qu.:0.3405      3rd Qu.:0.30623      3rd Qu.:0.3678  
## Max.      :1.0000    Max.      :1.0000    Max.      :1.00000    Max.      :1.0000  
## symmetry_mean      fractal_dimension_mean      radius_se  
## Min. :0.0000 Min. :0.0000 Min. :0.00000 ## 1st Qu.:0.2823 1st Qu.:0.1630  
1st Qu.:0.04378 ## Median :0.3697 Median :0.2439 Median :0.07702 ##  
Mean :0.3796 Mean :0.2704 Mean :0.10635 ## 3rd Qu.:0.4530 3rd  
Qu.:0.3404 3rd Qu.:0.13304 ## Max. :1.0000 Max. :1.0000 Max. :1.00000  
##      texture_se      perimeter_se      area_se      smoothness_se  
## Min. :0.0000 Min. :0.00000 Min. :0.00000 Min. :0.0000 ## 1st Qu.:0.1047 1st Qu.:0.04000  
1st Qu.:0.02064 1st Qu.:0.1175 ## Median :0.1653 Median :0.07209 Median :0.03311  
Median :0.1586 ## Mean :0.1893 Mean :0.09938 Mean :0.06264 Mean :0.1811 ## 3rd  
Qu.:0.2462 3rd Qu.:0.12251 3rd Qu.:0.07170 3rd Qu.:0.2187 ## Max. :1.0000 Max. :1.00000  
Max. :1.00000 Max. :1.0000  
## compactness_se      concavity_se      concave.points_se symmetry_se  
## Min. :0.00000 Min. :0.00000 Min. :0.0000 Min. :0.0000 ## 1st Qu.:0.08132 1st Qu.:0.03811  
1st Qu.:0.1447 1st Qu.:0.1024 ## Median :0.13667 Median :0.06538 Median :0.2070 Median  
:0.1526 ## Mean :0.17444 Mean :0.08054 Mean :0.2235 Mean :0.1781 ## 3rd Qu.:0.22680  
3rd Qu.:0.10619 3rd Qu.:0.2787 3rd Qu.:0.2195 ## Max. :1.00000 Max. :1.00000 Max. :1.0000  
Max. :1.0000  
## fractal_dimension_se radius_worst      texture_worst      perimeter_worst  
## Min. :0.00000 Min. :0.0000 Min. :0.0000 Min. :0.0000 ## 1st Qu.:0.04675 1st Qu.:0.1807 1st  
Qu.:0.2415 1st Qu.:0.1678 ## Median :0.07919 Median :0.2504 Median :0.3569 Median  
:0.2353 ## Mean :0.10019 Mean :0.2967 Mean :0.3640 Mean :0.2831 ## 3rd Qu.:0.12656 3rd  
Qu.:0.3863 3rd Qu.:0.4717 3rd Qu.:0.3735 ## Max. :1.00000 Max. :1.0000 Max. :1.0000 Max.  
:1.0000  
## area_worst smoothness_worst compactness_worst concavity_worst ## Min. :0.00000  
Min. :0.0000 Min. :0.0000 Min. :0.00000 ## 1st Qu.:0.08113 1st Qu.:0.3000 1st Qu.:0.1163 1st
```

```

Qu.:0.09145 ## Median :0.12321 Median :0.3971 Median :0.1791 Median :0.18107 ## Mean
:0.17091 Mean :0.4041 Mean :0.2202 Mean :0.21740 ## 3rd Qu.:0.22090 3rd Qu.:0.4942 3rd
Qu.:0.3025 3rd Qu.:0.30583 ## Max. :1.00000 Max. :1.0000 Max. :1.0000 Max. :1.00000
## concave.points_worst symmetry_worst fractal_dimension_worst
## Min. :0.0000 Min. :0.0000 Min. :0.0000 ## 1st Qu.:0.2231 1st
Qu.:0.1851 1st Qu.:0.1077 ## Median :0.3434 Median :0.2478 Median
:0.1640 ## Mean :0.3938 Mean :0.2633 Mean :0.1896 ## 3rd Qu.:0.5546
3rd Qu.:0.3182 3rd Qu.:0.2429
## Max. :1.0000 Max. :1.0000 Max. :1.0000

```

Dividing Training and Test Data

We take N_sample rows as test cases and use the rest to train the classifier.

```
N_sample = 100
```

```
s_index = sample(1:569, N_sample) data_Train = data_n[-s_index,] data_Test =
data_n[s_index,]
```

KNN

```
N_K = 25 data_knn = knn(train = data_Train, test = data_Test, cl = data[-s_index, 1], k = N_K)
```

Accuracy Analysis

The confusion matrix is shown below.

```
CrossTable(x = data[s_index,1], data_knn)
```

```

## ##
##      Cell Contents
## |-----|
## |      N | ## | Chi-square contribution
## |
## |      N / Row Total | ## |      N /
Col Total | ## |      N / Table Total |
## |-----|
## ##
## Total Observations in Table: 100
## ##
##      | data_knn
## data[s_index, 1] |      B |      M | Row Total |
## -----|-----|-----|-----|
##      B |      62 |      0 |      62 |
##      | 12.555 | 22.320 | ##      | 1.000 | 0.000 |
##      0.620 | ##      |      0.969 | 0.000 | | ##      |
##      0.620 | 0.000 | | ## -----|-----|-----|
##      M |      2 |      36 |      38 |
##      | 20.484 | 36.417 | ##      | 0.053 | 0.947 |
##      0.380 | ##      |      0.031 | 1.000 | | ##      |
##      0.020 | 0.360 | | ## -----|-----|-----|
##      Column Total |      64 |      36 |      100 |

```

| 0.640 | 0.360 | | ## -----|-----|-----|----
-----|
##