

# proj-Breast Cancer Detection

2023-04-04

Loading the dataset

```
path <- file.choose()
```

```
df <- read.csv(path)
df <- df[,-33]
head(df)
```

```
##      id diagnosis radius_mean texture_mean perimeter_mean area_mean
## 1  842302         M      17.99      10.38         122.80      1001.0
## 2  842517         M      20.57      17.77         132.90      1326.0
## 3 84300903         M      19.69      21.25         130.00      1203.0
## 4 84348301         M      11.42      20.38          77.58       386.1
## 5 84358402         M      20.29      14.34         135.10      1297.0
## 6  843786         M      12.45      15.70          82.57       477.1
## smoothness_mean compactness_mean concavity_mean concave.points_mean
## 1      0.11840      0.27760      0.3001      0.14710
## 2      0.08474      0.07864      0.0869      0.07017
## 3      0.10960      0.15990      0.1974      0.12790
## 4      0.14250      0.28390      0.2414      0.10520
## 5      0.10030      0.13280      0.1980      0.10430
## 6      0.12780      0.17000      0.1578      0.08089
## symmetry_mean fractal_dimension_mean radius_se texture_se perimeter_se
## 1      0.2419      0.07871      1.0950      0.9053      8.589
## 2      0.1812      0.05667      0.5435      0.7339      3.398
## 3      0.2069      0.05999      0.7456      0.7869      4.585
## 4      0.2597      0.09744      0.4956      1.1560      3.445
## 5      0.1809      0.05883      0.7572      0.7813      5.438
## 6      0.2087      0.07613      0.3345      0.8902      2.217
## area_se smoothness_se compactness_se concavity_se concave.points_se
## 1 153.40      0.006399      0.04904      0.05373      0.01587
## 2  74.08      0.005225      0.01308      0.01860      0.01340
## 3  94.03      0.006150      0.04006      0.03832      0.02058
## 4  27.23      0.009110      0.07458      0.05661      0.01867
## 5  94.44      0.011490      0.02461      0.05688      0.01885
## 6  27.19      0.007510      0.03345      0.03672      0.01137
## symmetry_se fractal_dimension_se radius_worst texture_worst perimeter_worst
## 1  0.03003      0.006193      25.38      17.33      184.60
## 2  0.01389      0.003532      24.99      23.41      158.80
## 3  0.02250      0.004571      23.57      25.53      152.50
## 4  0.05963      0.009208      14.91      26.50      98.87
## 5  0.01756      0.005115      22.54      16.67      152.20
## 6  0.02165      0.005082      15.47      23.75      103.40
```

```
##   area_worst smoothness_worst compactness_worst concavity_worst
## 1    2019.0         0.1622         0.6656         0.7119
## 2    1956.0         0.1238         0.1866         0.2416
## 3    1709.0         0.1444         0.4245         0.4504
## 4     567.7         0.2098         0.8663         0.6869
## 5    1575.0         0.1374         0.2050         0.4000
## 6     741.6         0.1791         0.5249         0.5355
##   concave.points_worst symmetry_worst fractal_dimension_worst
## 1             0.2654             0.4601             0.11890
## 2             0.1860             0.2750             0.08902
## 3             0.2430             0.3613             0.08758
## 4             0.2575             0.6638             0.17300
## 5             0.1625             0.2364             0.07678
## 6             0.1741             0.3985             0.12440
```

Handling the NA Values with mean of the feature records and also omitting remaining NA value's records

```
df$radius_mean <- ifelse(is.na(df$radius_mean),
  ave(df$radius_mean, FUN = function(x)mean(x, na.rm = TRUE)),
  df$radius_mean)
df$area_mean <- ifelse(is.na(df$area_mean),
  ave(df$area_mean, FUN = function(x)mean(x, na.rm = TRUE)),
  df$area_mean)
df$concave.points_worst <- ifelse(is.na(df$concave.points_worst),
  ave(df$concave.points_worst, FUN = function(x)mean(x, na.rm = TRUE)),
  df$concave.points_worst)
df$area_worst = ifelse(is.na(df$area_worst),
  ave(df$area_worst, FUN = function(x)mean(x, na.rm = TRUE)),
  df$area_worst)
df$concave.points_mean = ifelse(is.na(df$concave.points_mean),
  ave(df$concave.points_mean, FUN = function(x)mean(x, na.rm = TRUE)),
  df$concave.points_mean)
df$area_se = ifelse(is.na(df$area_se),
  ave(df$area_se, FUN = function(x)mean(x, na.rm = TRUE)),
  df$area_se)
df$concavity_se = ifelse(is.na(df$concavity_se),
  ave(df$concavity_se, FUN = function(x)mean(x, na.rm = TRUE)),
  df$concavity_se)
```

Removing the ID Column as this doesn't affect the result

```
df <- df[, -1]
```

Encoding the Categorical data for Diagnosis where 1 represents M (Malignant) and 2 represents B (Benign)

```
df$diagnosis <- factor(df$diagnosis, levels = c('M', 'B'), labels = c(1, 2))
head(df)
```

```
##   diagnosis radius_mean texture_mean perimeter_mean area_mean smoothness_mean
## 1         1      17.99       10.38         122.80      1001.0         0.11840
## 2         1      20.57       17.77         132.90      1326.0         0.08474
## 3         1      19.69       21.25         130.00      1203.0         0.10960
```

```

## 4      1      11.42      20.38      77.58      386.1      0.14250
## 5      1      20.29      14.34      135.10     1297.0      0.10030
## 6      1      12.45      15.70      82.57      477.1      0.12780
## compactness_mean concavity_mean concave.points_mean symmetry_mean
## 1      0.27760      0.3001      0.14710      0.2419
## 2      0.07864      0.0869      0.07017      0.1812
## 3      0.15990      0.1974      0.12790      0.2069
## 4      0.28390      0.2414      0.10520      0.2597
## 5      0.13280      0.1980      0.10430      0.1809
## 6      0.17000      0.1578      0.08089      0.2087
## fractal_dimension_mean radius_se texture_se perimeter_se area_se
## 1      0.07871      1.0950      0.9053      8.589 153.40
## 2      0.05667      0.5435      0.7339      3.398 74.08
## 3      0.05999      0.7456      0.7869      4.585 94.03
## 4      0.09744      0.4956      1.1560      3.445 27.23
## 5      0.05883      0.7572      0.7813      5.438 94.44
## 6      0.07613      0.3345      0.8902      2.217 27.19
## smoothness_se compactness_se concavity_se concave.points_se symmetry_se
## 1      0.006399      0.04904      0.05373      0.01587 0.03003
## 2      0.005225      0.01308      0.01860      0.01340 0.01389
## 3      0.006150      0.04006      0.03832      0.02058 0.02250
## 4      0.009110      0.07458      0.05661      0.01867 0.05963
## 5      0.011490      0.02461      0.05688      0.01885 0.01756
## 6      0.007510      0.03345      0.03672      0.01137 0.02165
## fractal_dimension_se radius_worst texture_worst perimeter_worst area_worst
## 1      0.006193      25.38      17.33      184.60 2019.0
## 2      0.003532      24.99      23.41      158.80 1956.0
## 3      0.004571      23.57      25.53      152.50 1709.0
## 4      0.009208      14.91      26.50      98.87 567.7
## 5      0.005115      22.54      16.67      152.20 1575.0
## 6      0.005082      15.47      23.75      103.40 741.6
## smoothness_worst compactness_worst concavity_worst concave.points_worst
## 1      0.1622      0.6656      0.7119      0.2654
## 2      0.1238      0.1866      0.2416      0.1860
## 3      0.1444      0.4245      0.4504      0.2430
## 4      0.2098      0.8663      0.6869      0.2575
## 5      0.1374      0.2050      0.4000      0.1625
## 6      0.1791      0.5249      0.5355      0.1741
## symmetry_worst fractal_dimension_worst
## 1      0.4601      0.11890
## 2      0.2750      0.08902
## 3      0.3613      0.08758
## 4      0.6638      0.17300
## 5      0.2364      0.07678
## 6      0.3985      0.12440

```

Splitting the data into train and test data using dplyr package

```
library(dplyr)
```

```
## Warning: package 'dplyr' was built under R version 4.2.3
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##   filter, lag
```

```
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
df$id <- 1:nrow(df)
```

```
# Splitting into 80-training and 20-test dataset
trn <- df%>dplyr::sample_frac(0.80)
tst <- dplyr::anti_join(df,trn,by='id')
```

```
trn <- trn[,-32]
tst <- trn[,-32]
df <- df[,-32]
```

```
head(trn)
```

```
##   diagnosis radius_mean texture_mean perimeter_mean area_mean smoothness_mean
## 1         2      11.890       18.35         77.32      432.2         0.09363
## 2         2       8.734       16.84         55.27      234.3         0.10390
## 3         2      12.890       15.70         84.08      516.6         0.07818
## 4         1      17.990       10.38        122.80     1001.0         0.11840
## 5         2      11.800       17.26         75.26      431.9         0.09087
## 6         1      15.340       14.26        102.50      704.4         0.10730
##   compactness_mean concavity_mean concave.points_mean symmetry_mean
## 1      0.11540      0.06636      0.03142      0.1967
## 2      0.07428      0.00000      0.00000      0.1985
## 3      0.09580      0.11150      0.03390      0.1432
## 4      0.27760      0.30010      0.14710      0.2419
## 5      0.06232      0.02853      0.01638      0.1847
## 6      0.21350      0.20770      0.09756      0.2521
##   fractal_dimension_mean radius_se texture_se perimeter_se area_se
## 1      0.06314      0.2963      1.5630      2.087      21.46
## 2      0.07098      0.5169      2.0790      3.167      28.85
## 3      0.05935      0.2913      1.3890      2.347      23.29
## 4      0.07871      1.0950      0.9053      8.589     153.40
## 5      0.06019      0.3438      1.1400      2.225      25.06
## 6      0.07032      0.4388      0.7096      3.384      44.91
##   smoothness_se compactness_se concavity_se concave.points_se symmetry_se
## 1      0.008872      0.04192      0.05946      0.017850      0.02793
## 2      0.015820      0.01966      0.00000      0.000000      0.01865
## 3      0.006418      0.03961      0.07927      0.017740      0.01878
## 4      0.006399      0.04904      0.05373      0.015870      0.03003
## 5      0.005463      0.01964      0.02079      0.005398      0.01477
## 6      0.006789      0.05328      0.06446      0.022520      0.03672
##   fractal_dimension_se radius_worst texture_worst perimeter_worst area_worst
## 1      0.004775      13.25      27.10      86.20      531.2
## 2      0.006736      10.17      22.80      64.01      317.0
## 3      0.003696      13.90      19.69      92.12      595.6
## 4      0.006193      25.38      17.33     184.60     2019.0
```

```
## 5          0.003071      13.45      24.49      86.00      562.0
## 6          0.004394      18.07      19.08     125.10     980.9
## smoothness_worst compactness_worst concavity_worst concave.points_worst
## 1          0.14050      0.3046      0.2806      0.11380
## 2          0.14600      0.1310      0.0000      0.00000
## 3          0.09926      0.2317      0.3344      0.10170
## 4          0.16220      0.6656      0.7119      0.26540
## 5          0.12440      0.1726      0.1449      0.05356
## 6          0.13900      0.5954      0.6305      0.23930
## symmetry_worst fractal_dimension_worst
## 1          0.3397      0.08365
## 2          0.2445      0.08865
## 3          0.1999      0.07127
## 4          0.4601      0.11890
## 5          0.2779      0.08121
## 6          0.4667      0.09946
```

```
head(tst)
```

```
## diagnosis radius_mean texture_mean perimeter_mean area_mean smoothness_mean
## 1          2      11.890      18.35      77.32      432.2      0.09363
## 2          2       8.734      16.84      55.27      234.3      0.10390
## 3          2      12.890      15.70      84.08      516.6      0.07818
## 4          1      17.990      10.38      122.80     1001.0      0.11840
## 5          2      11.800      17.26      75.26      431.9      0.09087
## 6          1      15.340      14.26     102.50      704.4      0.10730
## compactness_mean concavity_mean concave.points_mean symmetry_mean
## 1          0.11540      0.06636      0.03142      0.1967
## 2          0.07428      0.00000      0.00000      0.1985
## 3          0.09580      0.11150      0.03390      0.1432
## 4          0.27760      0.30010      0.14710      0.2419
## 5          0.06232      0.02853      0.01638      0.1847
## 6          0.21350      0.20770      0.09756      0.2521
## fractal_dimension_mean radius_se texture_se perimeter_se area_se
## 1          0.06314      0.2963      1.5630      2.087      21.46
## 2          0.07098      0.5169      2.0790      3.167      28.85
## 3          0.05935      0.2913      1.3890      2.347      23.29
## 4          0.07871      1.0950      0.9053      8.589     153.40
## 5          0.06019      0.3438      1.1400      2.225      25.06
## 6          0.07032      0.4388      0.7096      3.384      44.91
## smoothness_se compactness_se concavity_se concave.points_se symmetry_se
## 1          0.008872      0.04192      0.05946      0.017850      0.02793
## 2          0.015820      0.01966      0.00000      0.000000      0.01865
## 3          0.006418      0.03961      0.07927      0.017740      0.01878
## 4          0.006399      0.04904      0.05373      0.015870      0.03003
## 5          0.005463      0.01964      0.02079      0.005398      0.01477
## 6          0.006789      0.05328      0.06446      0.022520      0.03672
## fractal_dimension_se radius_worst texture_worst perimeter_worst area_worst
## 1          0.004775      13.25      27.10      86.20      531.2
## 2          0.006736      10.17      22.80      64.01      317.0
## 3          0.003696      13.90      19.69      92.12      595.6
## 4          0.006193      25.38      17.33     184.60     2019.0
## 5          0.003071      13.45      24.49      86.00      562.0
## 6          0.004394      18.07      19.08     125.10     980.9
```

	smoothness_worst	compactness_worst	concavity_worst	concave.points_worst
## 1	0.14050	0.3046	0.2806	0.11380
## 2	0.14600	0.1310	0.0000	0.00000
## 3	0.09926	0.2317	0.3344	0.10170
## 4	0.16220	0.6656	0.7119	0.26540
## 5	0.12440	0.1726	0.1449	0.05356
## 6	0.13900	0.5954	0.6305	0.23930

	symmetry_worst	fractal_dimension_worst
## 1	0.3397	0.08365
## 2	0.2445	0.08865
## 3	0.1999	0.07127
## 4	0.4601	0.11890
## 5	0.2779	0.08121
## 6	0.4667	0.09946

Implementing decision tree classifier

```
library(party)

## Warning: package 'party' was built under R version 4.2.3

## Loading required package: grid

## Loading required package: mvtnorm

## Loading required package: modeltools

## Loading required package: stats4

## Loading required package: strucchange

## Warning: package 'strucchange' was built under R version 4.2.3

## Loading required package: zoo

## Warning: package 'zoo' was built under R version 4.2.3

##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##   as.Date, as.Date.numeric

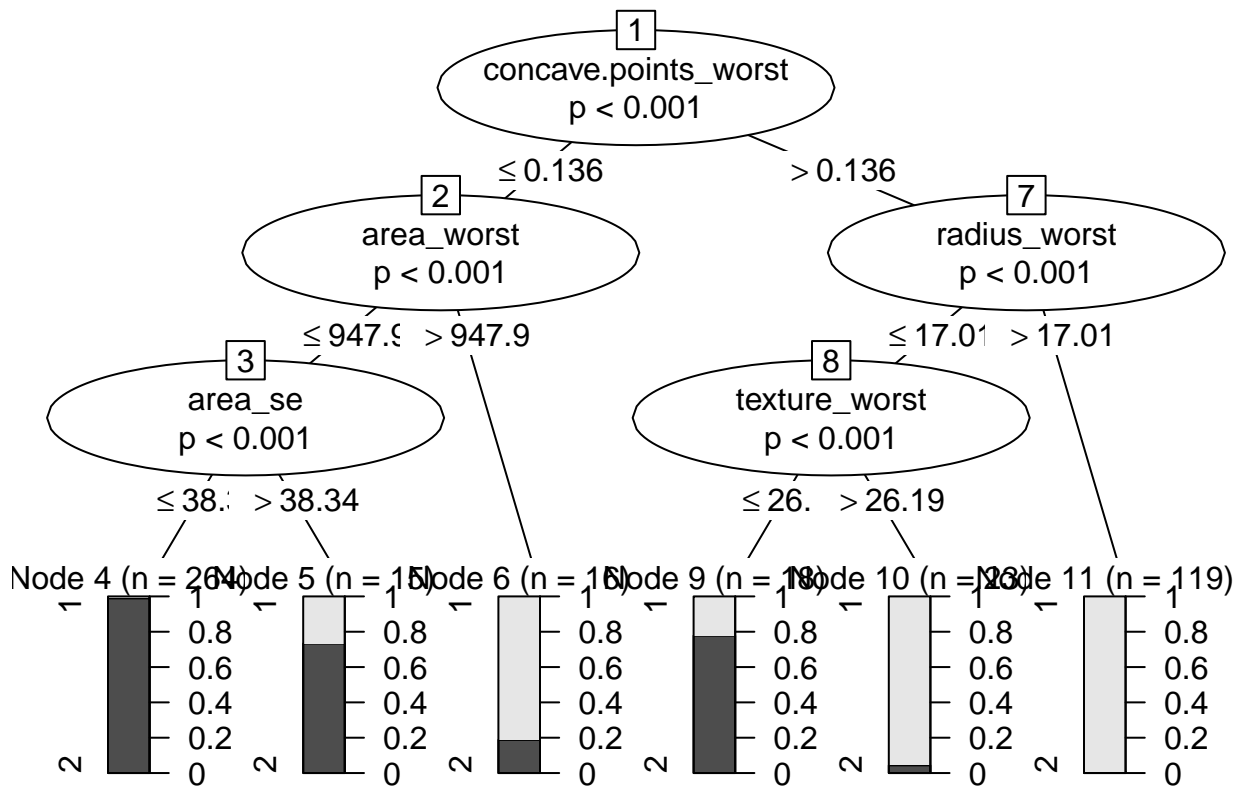
## Loading required package: sandwich

## Warning: package 'sandwich' was built under R version 4.2.3

##
## Attaching package: 'party'

## The following object is masked from 'package:dplyr':
##
##   where
```

```
dt <- ctree(diagnosis~., trn)
plot(dt)
```



```
dt
```

```
##
## Conditional inference tree with 6 terminal nodes
##
## Response: diagnosis
## Inputs: radius_mean, texture_mean, perimeter_mean, area_mean, smoothness_mean, compactness_mean, concave.points_worst
## Number of observations: 455
##
## 1) concave.points_worst <= 0.1357; criterion = 1, statistic = 286.707
## 2) area_worst <= 947.9; criterion = 1, statistic = 108.068
## 3) area_se <= 38.34; criterion = 1, statistic = 28.407
## 4)* weights = 264
## 3) area_se > 38.34
## 5)* weights = 15
## 2) area_worst > 947.9
## 6)* weights = 16
## 1) concave.points_worst > 0.1357
## 7) radius_worst <= 17.01; criterion = 1, statistic = 28.921
## 8) texture_worst <= 26.19; criterion = 1, statistic = 19.78
## 9)* weights = 18
## 8) texture_worst > 26.19
```

```
##      10)* weights = 23
##      7) radius_worst > 17.01
##      11)* weights = 119
```

Predicted output of test data

```
dt_pred <- predict(dt,tst)
dt_pred
```

```
##      [1] 2 2 2 1 2 1 2 2 2 2 1 2 2 2 2 2 2 1 1 2 2 1 2 2 1 2 2 1 1 1 2 2 2 2 1 2 1
##      [38] 1 2 1 1 1 2 2 1 2 2 2 1 1 1 2 1 2 2 2 2 2 2 1 1 1 1 2 2 2 1 2 2 2 2 2 1 2
##      [75] 1 2 2 2 2 2 2 2 2 2 2 1 2 2 2 1 2 1 2 1 2 2 2 1 2 2 2 2 1 2 1 1 1 2 2 1 1
##     [112] 1 1 2 1 2 2 2 1 1 1 2 2 2 1 2 2 2 2 1 2 2 2 1 2 1 1 2 1 1 2 1 2 1 2 2 2 2
##     [149] 1 1 2 1 1 2 2 1 2 2 2 1 1 2 2 1 2 2 2 2 1 2 1 1 2 1 2 1 2 2 2 2 2 2 1 1 2
##     [186] 1 1 2 2 2 1 2 2 2 2 2 2 1 2 2 1 1 1 1 1 2 1 2 2 2 1 2 2 2 1 2 2 2 2 1 2 2
##     [223] 2 2 2 2 1 2 1 1 1 2 2 2 1 2 2 2 2 2 2 2 2 2 2 1 1 2 2 2 2 2 1 1 2 2 2 2
##     [260] 1 2 2 2 2 2 1 2 2 2 1 2 2 2 1 2 2 2 2 2 2 2 2 1 1 2 1 2 1 2 2 2 2 1 2 2 2
##     [297] 1 1 1 2 2 2 2 2 2 1 2 1 2 1 1 2 2 1 1 2 1 2 2 2 2 1 2 2 2 1 1 1 2 2 1 2 1
##     [334] 1 2 1 2 1 2 1 2 2 1 2 2 2 1 2 1 2 2 2 2 2 2 2 1 2 2 1 2 2 2 2 2 1 2 2 1 2 2
##     [371] 2 2 2 2 2 2 1 2 2 2 1 2 1 1 2 2 2 2 1 1 2 1 1 2 1 2 2 2 2 1 2 2 1 2 1 2 2
##     [408] 1 2 1 2 2 2 2 1 2 1 2 2 2 2 2 2 1 2 1 1 2 2 2 2 1 2 2 2 2 1 1 1 1 2 2 2
##     [445] 1 1 2 1 1 1 2 2 2 2
## Levels: 1 2
```

Confusion matrix for the decision tree model

```
library(caret)
```

```
## Warning: package 'caret' was built under R version 4.2.3
```

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

```
## Loading required package: lattice
```

```
confusionMatrix(dt_pred, tst$diagnosis)
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction   1    2
##              1 153   5
##              2   11 286
##
##              Accuracy : 0.9648
##              95% CI : (0.9435, 0.9798)
##              No Information Rate : 0.6396
##              P-Value [Acc > NIR] : <2e-16
##
##              Kappa : 0.9231
##
##              Mcnemar's Test P-Value : 0.2113
```



```
##
##          Sensitivity : 0.9329
##          Specificity : 0.9828
##          Pos Pred Value : 0.9684
##          Neg Pred Value : 0.9630
##          Prevalence : 0.3604
##          Detection Rate : 0.3363
##          Detection Prevalence : 0.3473
##          Balanced Accuracy : 0.9579
##
##          'Positive' Class : 1
##
```

Naive Bayes model

```
library(e1071)
```

```
## Warning: package 'e1071' was built under R version 4.2.3
```

```
nb <- naiveBayes(diagnosis ~ ., data = trn)
nb_pred <- predict(nb, newdata = tst)
confusionMatrix(nb_pred, tst$diagnosis)
```

```
## Confusion Matrix and Statistics
##
##          Reference
## Prediction  1   2
##          1 149  11
##          2  15 280
##
##          Accuracy : 0.9429
##          95% CI : (0.9174, 0.9623)
##          No Information Rate : 0.6396
##          P-Value [Acc > NIR] : <2e-16
##
##          Kappa : 0.8754
##
##          Mcnemar's Test P-Value : 0.5563
##
##          Sensitivity : 0.9085
##          Specificity : 0.9622
##          Pos Pred Value : 0.9313
##          Neg Pred Value : 0.9492
##          Prevalence : 0.3604
##          Detection Rate : 0.3275
##          Detection Prevalence : 0.3516
##          Balanced Accuracy : 0.9354
##
##          'Positive' Class : 1
##
```

SVM model for different kernels

```
library(e1071)
# Linear kernel
svm <- svm(diagnosis ~ ., data = trn, kernel = "linear")
svm_pred <- predict(svm, newdata = tst)
confusionMatrix(svm_pred, as.factor(tst$diagnosis))
```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction  1    2
##           1 160    1
##           2    4 290
##
##           Accuracy : 0.989
##           95% CI : (0.9745, 0.9964)
##       No Information Rate : 0.6396
##       P-Value [Acc > NIR] : <2e-16
##
##           Kappa : 0.9761
##
##  McNemar's Test P-Value : 0.3711
##
##           Sensitivity : 0.9756
##           Specificity : 0.9966
##       Pos Pred Value : 0.9938
##       Neg Pred Value : 0.9864
##           Prevalence : 0.3604
##       Detection Rate : 0.3516
##       Detection Prevalence : 0.3538
##       Balanced Accuracy : 0.9861
##
##       'Positive' Class : 1
##
```

```
# Radial kernel
svm <- svm(diagnosis ~ ., data = trn, kernel = "radial")
svm_pred <- predict(svm, newdata = tst)
confusionMatrix(svm_pred, tst$diagnosis)
```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction  1    2
##           1 160    0
##           2    4 291
##
##           Accuracy : 0.9912
##           95% CI : (0.9776, 0.9976)
##       No Information Rate : 0.6396
##       P-Value [Acc > NIR] : <2e-16
##
##           Kappa : 0.9808
```

```
##
## McNemar's Test P-Value : 0.1336
##
##      Sensitivity : 0.9756
##      Specificity : 1.0000
##      Pos Pred Value : 1.0000
##      Neg Pred Value : 0.9864
##      Prevalence : 0.3604
##      Detection Rate : 0.3516
##      Detection Prevalence : 0.3516
##      Balanced Accuracy : 0.9878
##
##      'Positive' Class : 1
##
```

#### *# Sigmoid kernel*

```
svm <- svm(diagnosis ~ ., data = trn, kernel = "sigmoid")
svm_pred <- predict(svm, newdata = tst)
confusionMatrix(svm_pred, tst$diagnosis)
```

```
## Confusion Matrix and Statistics
##
##      Reference
## Prediction  1   2
##      1 149  10
##      2   15 281
##
##      Accuracy : 0.9451
##      95% CI : (0.92, 0.9641)
##      No Information Rate : 0.6396
##      P-Value [Acc > NIR] : <2e-16
##
##      Kappa : 0.88
##
## McNemar's Test P-Value : 0.4237
##
##      Sensitivity : 0.9085
##      Specificity : 0.9656
##      Pos Pred Value : 0.9371
##      Neg Pred Value : 0.9493
##      Prevalence : 0.3604
##      Detection Rate : 0.3275
##      Detection Prevalence : 0.3495
##      Balanced Accuracy : 0.9371
##
##      'Positive' Class : 1
##
```

#### *# Polynomial kernel*

```
svm <- svm(diagnosis ~ ., data = trn, kernel = "polynomial")
svm_pred <- predict(svm, newdata = tst)
confusionMatrix(svm_pred, tst$diagnosis)
```

```
## Confusion Matrix and Statistics
```

```

##
##           Reference
## Prediction   1   2
##           1 126   0
##           2  38 291
##
##           Accuracy : 0.9165
##           95% CI : (0.8872, 0.9402)
##           No Information Rate : 0.6396
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.8092
##
## Mcnemar's Test P-Value : 1.947e-09
##
##           Sensitivity : 0.7683
##           Specificity : 1.0000
##           Pos Pred Value : 1.0000
##           Neg Pred Value : 0.8845
##           Prevalence : 0.3604
##           Detection Rate : 0.2769
##           Detection Prevalence : 0.2769
##           Balanced Accuracy : 0.8841
##
##           'Positive' Class : 1
##

```