

Naive Bayes

Breast Cancer. Use the 'wisc_bc_data.csv' dataset and build a Naïve Bayes model to predict 'diagnosis' cases M: malignant or B: benign depending on all the numerical predictors. Use set.seed(1234) and 455 cases to train the model.

Predict cases of malignant and benign in test dataset. Calculate the confusion matrix

```
> wbcd = read.csv(file.choose(), header =TRUE)
> str(wbcd)
'data.frame':
                569 obs. of 32 variables:
 $ id
                          : int 842302 842517 84300903 84348301 84358402 843786 844359 84458
202 844981 84501001 ...
 $ diagnosis
                          : chr "M" "M" "M" "M" ...
 $ 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 ...
                          : num 1001 1326 1203 386 1297 ...
 $ area_mean
                          : num 0.1184 0.0847 0.1096 0.1425 0.1003 ...
 $ smoothness_mean
 $ compactness_mean
                          : num 0.2776 0.0786 0.1599 0.2839 0.1328 ...
                          : num 0.3001 0.0869 0.1974 0.2414 0.198 ...
 $ concavity_mean
 $ 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 ...
                          : num 17.3 23.4 25.5 26.5 16.7 ...
 $ texture_worst
 $ perimeter_worst
                          : num 184.6 158.8 152.5 98.9 152.2 ...
                          : num 2019 1956 1709 568 1575 ...
 $ area_worst
                          : num 0.162 0.124 0.144 0.21 0.137 ...
 $ smoothness_worst
 $ 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 ...
> wbcd = wbcd[,-1]
> set.seed(1234)
> s = sample(569, 455)
> train = wbcd[s,]
> test = wbcd[-s,]
> train$diagnosis=as.factor(train$diagnosis)
> test$diagnosis=as.factor(test$diagnosis)
> fit = naiveBayes(diagnosis~.,data=train)
```

Naive Bayes

Naive Bayes 2