P2-DiabetesData

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I have written a python script to read the text data and convert it to csv. The feature names have been simplified.

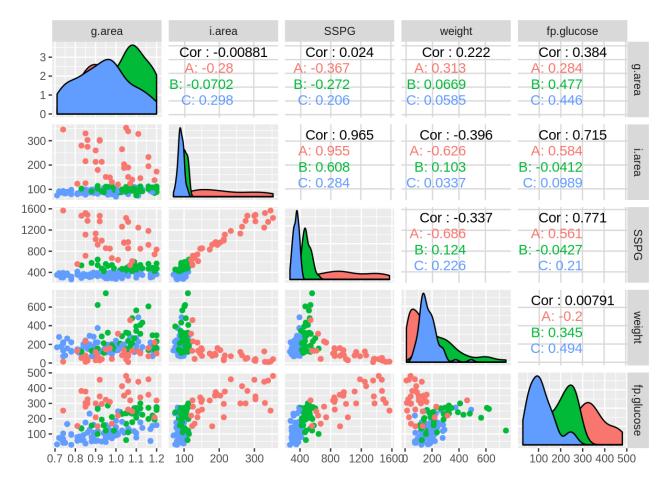
```
##
                                         SSPG
                                                         weight
        g.area
                          i.area
##
   Min.
           :0.7100
                     Min.
                            : 70
                                    Min.
                                           : 269.0
                                                     Min.
                                                             : 10.0
   1st Qu.:0.8800
                     1st Qu.: 90
                                    1st Qu.: 352.0
                                                     1st Qu.:118.0
##
   Median :0.9800
                     Median: 97
                                    Median : 413.0
                                                     Median :156.0
##
           :0.9773
   Mean
                     Mean
                           :122
                                    Mean
                                         : 543.6
                                                     Mean
                                                             :186.1
##
   3rd Qu.:1.0800
                     3rd Qu.:112
                                    3rd Qu.: 558.0
                                                     3rd Qu.:221.0
##
##
   Max.
          :1.2000
                     Max.
                            :353
                                    Max.
                                           :1568.0
                                                     Max.
                                                            :748.0
      fp.glucose
##
                        class
         : 29.0
## Min.
                    Min.
                           :1.000
   1st Qu.:100.0
                    1st Qu.:2.000
##
## Median :159.0
                    Median :3.000
          :184.2
                          :2.297
## Mean
                    Mean
   3rd Qu.:257.0
                    3rd Qu.:3.000
## Max.
           :480.0
                    Max.
                           :3.000
```

```
## [1] "g.area" "i.area" "SSPG" "weight" "fp.glucose" ## [6] "class"
```

Here are the prior probabilities of each class from class 1, class 2, followed by class 3, resp.

```
## [1] 0.2275862 0.2482759 0.5241379
```

Here is the pariwise scatter-plot for each of the five variables



Classes have difference covariance matrix The variances of feature distribution for classes are different as can be seen in the plot above. Consider, i.area, the variance for class 1 is larger than that for classes 2, and 3.

The classes are not multi-variate normal LDA assumes that the density of features for a given class is normall distributed. It can be seen from the image that the feature densities do not have an ideal normal distribution but are very close to normal. For i.area and SSPG, the distribution of class A(1), may not look normal, but it can be assumed to be normal with a large variance.

Test-train division

The data is split into 70% training and 25% test set. In the training data, the data points belonging to class 1, class 2, and class 3 are shown below.

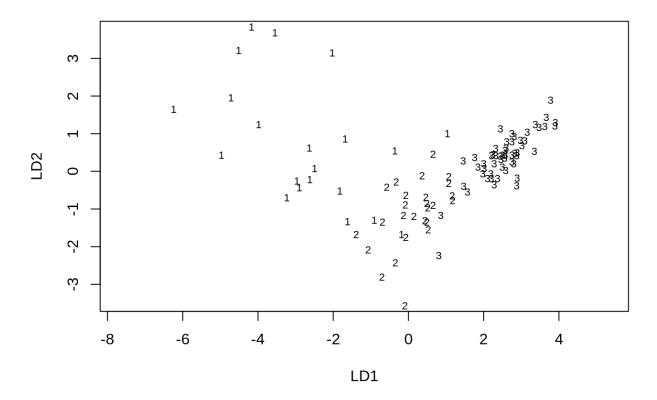
[1] 21 26 54

The the test data, the count of data points belonging to class 1, class 2, and class 3 are shown below.

[1] 12 10 22

LDA

```
## Call:
## lda(class ~ ., data = trainData)
## Prior probabilities of groups:
##
                       2
            1
## 0.2079208 0.2574257 0.5346535
## Group means:
         g.area
                    i.area
                                SSPG
                                      weight fp.glucose
## 1 0.9795238 204.57143 998.7619 128.7143
                                                  306.3810
## 2 1.0484615 100.76923 504.3846 277.4231
                                                  214.7308
## 3 0.9416667 91.42593 352.1852 171.0741
                                                  116.0926
##
## Coefficients of linear discriminants:
##
                         LD1
## g.area -0.979571205 -3.420970121
## i.area 0.039091321 0.033732203
## SSPG -0.013423147 -0.005573391
## weight
             -0.001082993 -0.005957180
## fp.glucose -0.005304303 0.000193961
##
## Proportion of trace:
      LD1
##
              LD2
## 0.8922 0.1078
```



The predictions done by LDA are as follows:

```
##
##
         1
            2
                3
            0
                0
##
      1 11
     2
            7
                1
##
         1
         0
            3 21
##
```

The miss-classification percentage for LDA on test data is:

```
## [1] 11.36364
```

LDAs performance on train data:

```
##
##
         1
            2
                3
##
     1 16
            0
                0
         4 23
                2
##
            3 52
##
     3
         1
```

Miss-classifications on train data:

```
## [1] 9.90099
```

QDA

```
## Call:
## qda(class ~ ., data = trainData)
## Prior probabilities of groups:
                     2
          1
## 0.2079208 0.2574257 0.5346535
##
## Group means:
                            SSPG weight fp.glucose
        g.area
                 i.area
## 1 0.9795238 204.57143 998.7619 128.7143
                                            306.3810
## 2 1.0484615 100.76923 504.3846 277.4231
                                            214.7308
## 3 0.9416667 91.42593 352.1852 171.0741
                                            116.0926
```

The predictions done by QDA on test data are as follows:

```
##
## 1 2 3
## 1 12 1 0
## 2 0 7 1
## 3 0 2 21
```

The miss-classification percentage of QDA on test data is:

```
## [1] 9.090909
```

Performance of QDA on train data:

```
##
## 1 2 3
## 1 12 1 0
## 2 0 7 1
## 3 0 2 21
```

The miss-classification percentage of QDA on train data is:

```
## [1] 3.960396
```

Performance of LDA and QDA

I did multiple tests by setting different seed values. In general QDA performed better than LDA. As seen above, for the current seed, QDA's test and train error are lesser than LDA's test and train error. The above observation might be due to the violation of the rule/assumption that all the features have the same variance. Violation of the above rule may make LDA a weaker candidate. QDA works when the variances are not the same.

Classification results for test sample

Following is the test sample:

```
## g.area i.area SSPG weight fp.glucose
## 1 0.98 122 544 186 184
```

LDA assigns this individual to:

```
## [1] 3
## Levels: 1 2 3
```

The posterior probabilities are shown below. It can be seen that LDA assigns the data point to class 3 with a probability which is only slightly greater than that for class 2.

```
## 1 2 3
## 1 0.002148242 0.475594 0.5222577
```

QDA assigns the individual to:

```
## [1] 2
## Levels: 1 2 3
```

The posterior probabilities are shown below. We can say that QDA classifies the data point very confidently to class 2.

```
## 1 0.4135972 0.5863852 1.756332e-05
```