## **KNN**

Record % correctly classified and confusion matrix (ALL positive class). Derive the confusion matrix when AML is the positive class. Calculate the class-dependent TP and FP rates. Show your work and compare to Weka.

	Predicted class	
True Class	Yes	No
Yes	TP: True Positive	FN: False Negative
No	FP: False Positive	TN: True Negative

$$TP \ rate = \frac{TP}{TP + FN}$$

$$FP \ rate = \frac{FP}{FP + TN}$$

Correctly Classified Instances

68

94.4444 %

TP Rate FP Rate Class 1.000 0.143 ALL 0.857 0.000 AML

=== Confusion Matrix ===

ALL as the positive class (Weka):

$$TP \ rate = \frac{44}{44+0} = 1.000$$

$$FP \ rate = \frac{4}{4 + 24} = 0.143$$

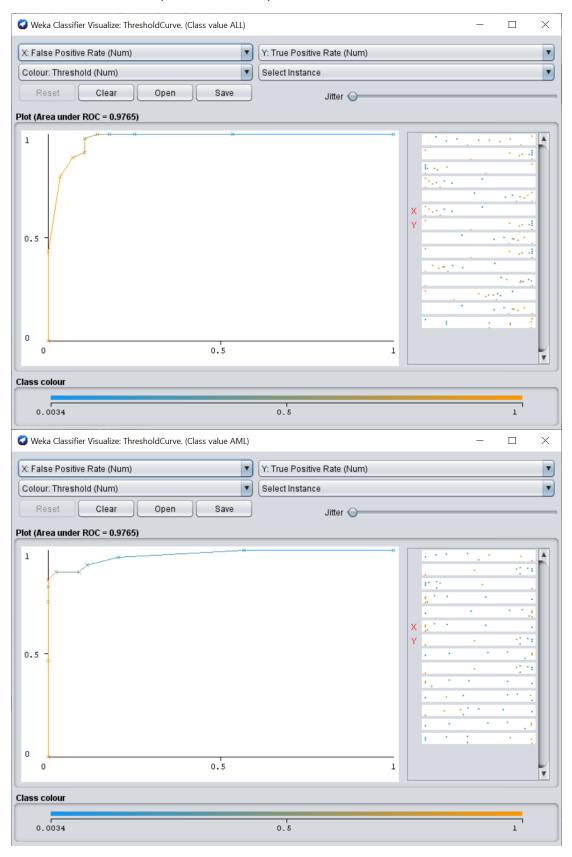
AML as the positive class (Derived):

$$TP \ rate = \frac{24}{24+4} = 0.857$$

$$FP\ rate = \frac{0}{0+44} = 0.000$$

My calculated values match the Weka results.

KNN: Capture the ROC curves for ALL positive and AML positive.



## **ZeroR**

Record % correctly classified and confusion matrix (ALL positive class). Derive the confusion matrix when AML is the positive class. Calculate the class-dependent TP and FP rates. Show your work and compare to Weka.

	Predicted class	
True Class	Yes	No
Yes	TP: True Positive	FN: False Negative
No	FP: False Positive	TN: True Negative

$$TP \ rate = \frac{TP}{TP + FN}$$

$$FP \ rate = \frac{FP}{FP + TN}$$

Correctly Classified Instances

44

61.1111 %

TP Rate FP Rate Class 1.000 1.000 ALL 0.000 0.000 AML

=== Confusion Matrix ===

ALL as the positive class (Weka):

$$TP \ rate = \frac{44}{44+0} = 1.000$$

$$FP \ rate = \frac{28}{28+0} = 1.000$$

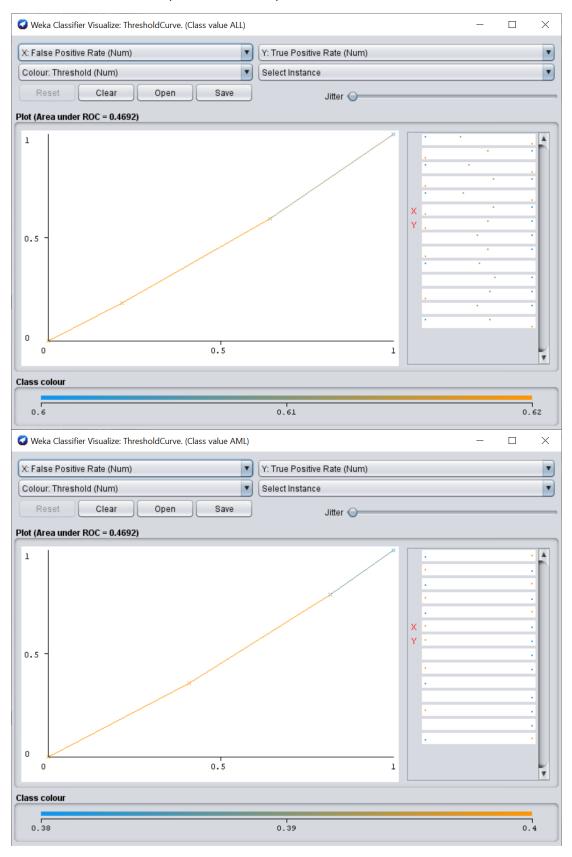
AML as the positive class (Derived):

$$TP \ rate = \frac{0}{0+28} = 0.000$$

$$FP\ rate = \frac{0}{0+44} = 0.000$$

My calculated values match the Weka results.

**ZeroR:** Capture the ROC curves for ALL positive and AML positive.



## **KNN**

```
=== Run information ===
Scheme:
              weka.classifiers.lazy.IBk -K 5 -W 0 -A "weka.core.neighboursearch.LinearNNSearch -A
\"weka.core.EuclideanDistance -R first-last\""
Relation:
              leukemia gene expression data names changed
Instances:
              72
Attributes:
             151
              [list of attributes omitted]
Test mode:
              5-fold cross-validation
=== Classifier model (full training set) ===
IB1 instance-based classifier
using 5 nearest neighbour(s) for classification
Time taken to build model: 0 seconds
=== Stratified cross-validation ===
=== Summary ===
Correctly Classified Instances
                                        68
                                                         94.4444 %
Incorrectly Classified Instances
                                                          5.5556 %
                                         4
Kappa statistic
                                         0.88
                                         0.0917
Mean absolute error
Root mean squared error
                                        0.2235
Relative absolute error
                                       19.2509 %
Root relative squared error
                                        45.7979 %
Total Number of Instances
                                        72
=== Detailed Accuracy By Class ===
                 TP Rate FP Rate Precision Recall
                                                       F-Measure MCC
                                                                           ROC Area PRC Area
                                                                                               Class
                 1.000
                                   0.917
                                              1.000
                                                       0.957
                                                                  0.886
                                                                           0.976
                                                                                     0.978
                                                                                               ALL
                          0.143
                 0.857
                          0.000
                                   1.000
                                              0.857
                                                       0.923
                                                                  0.886
                                                                           0.976
                                                                                     0.967
                                                                                               AML
Weighted Avg.
                0.944
                          0.087
                                   0.949
                                              0.944
                                                       0.944
                                                                  0.886
                                                                           0.976
                                                                                     0.973
=== Confusion Matrix ===
 a b
         <-- classified as
 44 0 | a = ALL
 4 24 | b = AML
```

## **ZeroR**

```
=== Run information ===
Scheme:
             weka.classifiers.rules.ZeroR
Relation:
             leukemia gene expression data names changed
Instances:
             72
Attributes: 151
             [list of attributes omitted]
Test mode:
             5-fold cross-validation
=== Classifier model (full training set) ===
ZeroR predicts class value: ALL
Time taken to build model: 0 seconds
=== Stratified cross-validation ===
=== Summary ===
                                       44
                                                        61.1111 %
Correctly Classified Instances
Incorrectly Classified Instances
                                       28
                                                        38.8889 %
Kappa statistic
                                        0
Mean absolute error
                                        0.4765
Root mean squared error
                                        0.4879
Relative absolute error
                                               %
                                      100
Root relative squared error
                                      100
                                               %
Total Number of Instances
                                      72
=== Detailed Accuracy By Class ===
                TP Rate FP Rate Precision Recall
                                                      F-Measure MCC
                                                                          ROC Area PRC Area Class
                1.000
                         1.000
                                  0.611
                                             1.000
                                                      0.759
                                                                 ?
                                                                          0.469
                                                                                   0.596
                                                                                             ALL
                0.000
                         0.000
                                  ?
                                             0.000
                                                      ?
                                                                 ?
                                                                          0.469
                                                                                   0.373
                                                                                             AML
Weighted Avg.
                0.611
                         0.611
                                  ?
                                             0.611
                                                      ?
                                                                 ?
                                                                          0.469
                                                                                   0.509
=== Confusion Matrix ===
 a b <-- classified as
44 0 | a = ALL
28 0 | b = AML
```