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#### **CPTS 434**

## Neural Network Design & Applications

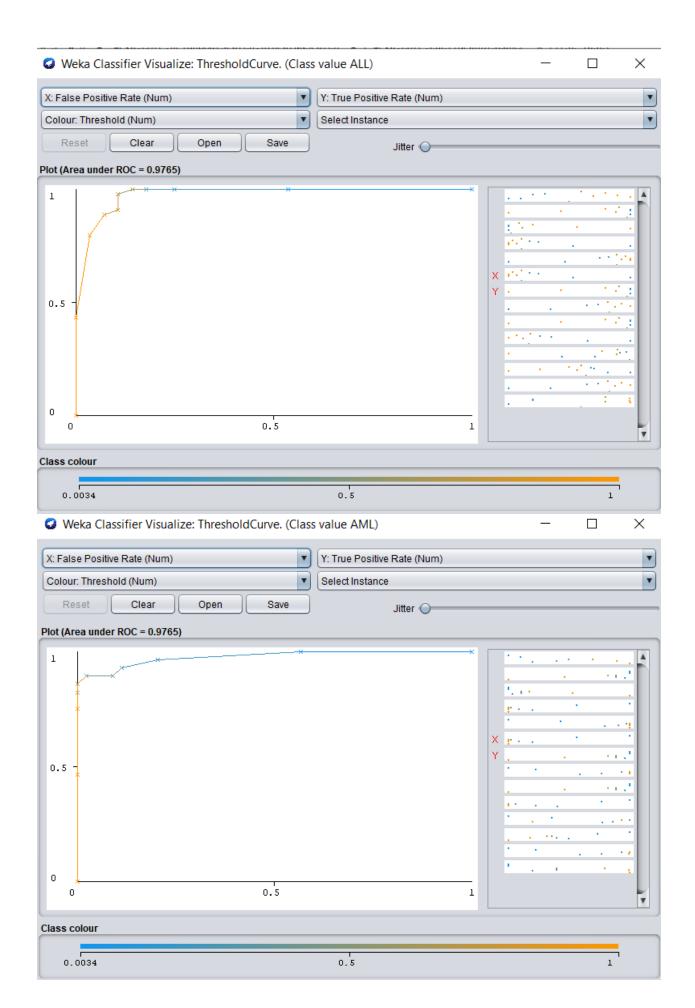
09-24-2019

#### Homework #2

#### **KNN Classifier**

```
--- Run information ---
             weka.classifiers.lazy.IBk -K 5 -W 0 -A "weka.core.neighboursearch.LinearNNSearch -A \"weka.core.EuclideanDistance -R first-last\""
Scheme:
             leukemia gene expression data names changed
Instances:
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
                                                     94.4444 %
Incorrectly Classified Instances
                                                      5.5556 %
                                      0.88
Kappa statistic
                                      0.0917
Mean absolute error
Root mean squared error
Relative absolute error
                                     19.2509 %
Root relative squared error
                                     45.7979 %
Total Number of Instances
=== Detailed Accuracy By Class ===
                TP Rate FP Rate Precision Recall F-Measure MCC
                                                                      ROC Area PRC Area Class
                1.000 0.143 0.917
0.857 0.000 1.000
                                        1.000
                                                             0.886
                                                                     0.976
                                                                              0.978
                                                   0.957
                                                                                         ALL
                                           0.857
                                                   0.923
                                                             0.886
                                                                     0.976
                                                                                0.967
                                                                                         AML
Weighted Avg.
                0.944 0.087
=== Confusion Matrix ===
  a b <-- classified as
4 24 | b = AML
=== Confusion Matrix =
 a b <-- classified as
24 \ 4 \mid b = AML
 0.44 \mid a = ALL
TP rate = 24/28 = 0.857\%
FP rate = 0/44 = 0.000\%
```

This matches the calculated accuracy provided by WEKA.



### ZeroR Classifier

```
=== Run information ===
Scheme:
            weka.classifiers.rules.ZeroR
Relation:
           leukemia gene expression data names changed
Instances:
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 ===
Correctly Classified Instances 44
                                                61.1111 %
                                  28
Incorrectly Classified Instances
                                                 38.8889 %
Kappa statistic
                                  0.4765
Mean absolute error
                                  0.4879
Root mean squared error
Relative absolute error
                                100 %
                                100
Root relative squared error
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 ?
0.611 ?
                                                        ?
                                                                0.469 0.373 AML
Weighted Avg. 0.611 ₹ 0.611 ?
                                                        ?
                                                                0.469 0.509
=== Confusion Matrix ===
 a b <-- classified as
 44 0 | a = ALL
↑28 0 | b = AML
=== Confusion Matrix ===
a b <-- classified as
44 0 \mid b = AML
28 0 \mid a = ALL
TP rate = 0/28 = 0.000\%
FP rate = 0/44 = 0.000\%
```

This matches the calculated accuracy provided by WEKA.

