CITS3401 Data Exploration and Mining Project 2

Wine Classification

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Introduction

The project specified that we are to develop several classifiers for wines using different classification methods to compare how machine learning performs compared to experts when rating different wines. The initial data is split into two groups, red wine and white wine, available from the UCI Machine Learning Repository[1]. Data analysis was done using Weka[2], data mining software created by Machine Learning Group at the University of Waikato, New Zealand. Specifically, we are using the classification and clustering tools in Weka Explorer for analysis.

Data Preprocessing

The initial data provided was in two files, winequality-red.csv and winequality-white.csv, that where converted to Weka's ARFF file format using an online conversion tool[3]. This tool was used to output two datasets, dataset 1 (ds1-red.arff and ds1-white.arff) and dataset 2 (ds2-red.arff and ds2-white.arff). Dataset 1 contains all the information that was in the original data, and is used to create the classifier. The fields in this dataset are numeric, apart from the quality which is nominal, making it is possible to group wines that receive the same rankings in Weka. Dataset 2 is contains all the numerical information from the original data and does not contain any information about the rankings from the wine tasters. The aim is to cluster these so that the wines fall into groups similar to the quality attribute of dataset 1.

Clustering

Dataset 2 requires clustering before it can be classified as the quality attribute of each data point has been removed. Simple K means clustering was used and after experimenting, fixed acidity, volatile acidity, citric acid, and density were ignored for the red wine data. This gave a roughly similar distrubution to the qualities found in the red wine dataset. In the white wine dataset, ignoring the different attributes has very little effect on the clustering. After ignoring different combinations of attributes, it does not seem possible to cluster the data points into a similar distrubution to the initial data. Comparing the clustering with the red wines quality information shows a 35% accuracy in the groupings, while for the clustered white wines there is only 25% accuracy.

```
Red Wine
```

```
Scheme:weka.clusterers.SimpleKMeans -N 6 -A
"weka.core.EuclideanDistance -R first-last" -I 500 -S 10
Clustered Instances
0
        632 (40%)
1
        128 (
               8%)
2
        196 (12%)
3
         32 ( 2%)
4
        339 (21%)
5
        272 (17%)
```

White Wine

```
Scheme:weka.clusterers.SimpleKMeans -N 7 -A
"weka.core.EuclideanDistance -R first-last" -I 500 -S 10
Clustered Instances
0
       1093 ( 22%)
1
        536 (11%)
2
        832 (17%)
3
        569 (12%)
4
        885 (18%)
5
        658 (13%)
6
        325 (7%)
```

Classification

Naive Bayesian

Naive Bayes classifiers are simple to implement, fast, and are used in real world situations such as spam filters. They work by looking at the traits of an object, and using each individual trait to determine how likely it is that the object falls into a specific classification. Their downside is that they assume the presence or absence of particular traits has no affect on the classification. We used cross-validation for Naive Bayes with 10 folds in Weka to calculate classify all the wines, and compared the classifications with the quality fields. Experimentation showed that increasing the number of folds for the red wine only had minimal effect on the accuracy of the classification.

Dataset 1

Red Wine

```
Scheme: weka.classifiers.bayes.NaiveBayes
Relation:
              ds1red
Instances:
              1599
Attributes:
              12
Test mode: 10-fold cross-validation
Correctly Classified Instances
                                        880
                                                           55.0344 %
Incorrectly Classified Instances
                                        719
                                                           44.9656 %
Kappa statistic
                                          0.311
Mean absolute error
                                          0.1763
                                          0.3198
Root mean squared error
Relative absolute error
                                         82.1845 %
Root relative squared error
                                         97.7154 %
Total Number of Instances
                                       1599
```

White Wine

Scheme: weka.classifiers.bayes.NaiveBayes

Relation: ds1white Instances: 4898 Attributes: 12

Test mode: 10-fold cross-validation

Correctly Classified Instances 2168 44.263 % Incorrectly Classified Instances 2730 55.737 %

Kappa statistic 0.2169
Mean absolute error 0.1721
Root mean squared error 0.3221
Relative absolute error 89.1485 %
Root relative squared error 103.6855 %

Total Number of Instances 4898

Dataset 2

Red Wine

Scheme:weka.classifiers.bayes.NaiveBayes

Relation: ds2red_clustered-weka.filters.unsupervised.attribute.Remove-R1

Instances: 1599
Attributes: 12

Test mode: 10-fold cross-validation

Correctly Classified Instances 1424 89.0557 % Incorrectly Classified Instances 175 10.9443 %

Kappa statistic 0.8548
Mean absolute error 0.0534
Root mean squared error 0.1698
Relative absolute error 21.401 %
Root relative squared error 48.0797 %

Total Number of Instances 1599

White Wine

Scheme:weka.classifiers.bayes.NaiveBayes

Relation: ds2white_clustered-weka.filters.unsupervised.attribute.Remove-R1

Instances: 4898
Attributes: 12

Test mode: 10-fold cross-validation

Correctly Classified Instances 4267 87.1172 % Incorrectly Classified Instances 631 12.8828 %

Kappa statistic 0.8467
Mean absolute error 0.0594
Root mean squared error 0.1667
Relative absolute error 24.7449 %
Root relative squared error 48.0952 %

Total Number of Instances 4898

Support Vector Machine

Support Vector Machine's are learning models and algorithms that analyze data and find patterns, then use the patterns for classification of data. Unlike the Naive Bayesian classifier, the support vector machine is a non-probabilistic classifier, meaning that it will not provide uncertainty for the results. This means that each different category is separated by as large a gap as possible.

Dataset 1

Red Wine

```
Scheme: weka.classifiers.functions.SMO -C 1.0 -L 0.001 -P 1.0E-12 -N 0 -V -1
-W 1 -K "weka.classifiers.functions.supportVector.PolyKernel -C 250007
-E 1.0"
Relation:
              ds1red
Instances:
              1599
Attributes:
Test mode: 10-fold cross-validation
                                                          58.349 %
Correctly Classified Instances
                                        933
Incorrectly Classified Instances
                                        666
                                                          41.651 %
Kappa statistic
                                          0.2905
Mean absolute error
                                          0.2349
Root mean squared error
                                          0.3301
Relative absolute error
                                        109.5032 %
Root relative squared error
                                        100.851 %
Total Number of Instances
                                       1599
```

White Wine

```
Scheme:weka.classifiers.functions.SMO -C 1.0 -L 0.001 -P 1.0E-12 -N 0 -V -1
-W 1 -K "weka.classifiers.functions.supportVector.PolyKernel -C 250007
-E 1.0"
Relation:
              ds1white
Instances:
              4898
Attributes:
              12
Test mode: 10-fold cross-validation
Correctly Classified Instances
                                                          52.0621 %
                                       2550
Incorrectly Classified Instances
                                       2348
                                                          47.9379 %
Kappa statistic
                                          0.1905
Mean absolute error
                                          0.2137
Root mean squared error
                                          0.3168
Relative absolute error
                                        110.7083 %
Root relative squared error
                                        101.9859 %
Total Number of Instances
                                       4898
```

Dataset 2

Red Wine

```
Scheme: weka.classifiers.functions.SMO -C 1.0 -L 0.001 -P 1.0E-12 -N 0 -V -1
-W 1 -K "weka.classifiers.functions.supportVector.PolyKernel -C 250007
-E 1.0"
Relation:
              ds2red_clustered-weka.filters.unsupervised.attribute.Remove-R1
              1599
Instances:
Attributes:
              12
Test mode: 10-fold cross-validation
                                                          94.1839 %
Correctly Classified Instances
                                       1506
                                                           5.8161 %
Incorrectly Classified Instances
                                         93
Kappa statistic
                                          0.9215
                                          0.2238
Mean absolute error
Root mean squared error
                                          0.3128
                                         89.7115 %
Relative absolute error
Root relative squared error
                                         88.5878 %
Total Number of Instances
                                       1599
```

White Wine

```
Scheme: weka.classifiers.functions.SMO -C 1.0 -L 0.001 -P 1.0E-12 -N 0 -V -1
-W 1 -K "weka.classifiers.functions.supportVector.PolyKernel -C 250007
-E 1.0"
Relation:
            ds2white_clustered-weka.filters.unsupervised.attribute.Remove-R1
Instances: 4898
Attributes: 12
Test mode: 10-fold cross-validation
                                                          96.1821 %
Correctly Classified Instances
                                      4711
                                                           3.8179 %
Incorrectly Classified Instances
                                       187
Kappa statistic
                                         0.9545
Mean absolute error
                                         0.2047
Root mean squared error
                                         0.3021
Relative absolute error
                                        85.1906 %
                                        87.1569 %
Root relative squared error
Total Number of Instances
                                      4898
```

Neural Network

Neural Networks are based off of animal's central nervous systems, by using several input sensors that transform the data before handing it on to another neuron. The neurons are connected together in a network and work simultaneously , rather then sequentially, to process the data. Real world applications for neural networks include speech and handwriting recognition.

Dataset 1

Red Wine

Scheme:weka.classifiers.functions.MultilayerPerceptron -L 0.3 -M 0.2 -N 500 -V 0 -S 0 -E 20 -H a Relation: ds1red 1599 Instances: Attributes: 12 Test mode: 10-fold cross-validation Correctly Classified Instances 967 60.4753 % Incorrectly Classified Instances 39.5247 % 632 Kappa statistic 0.3585 Mean absolute error 0.1657 Root mean squared error 0.3021 Relative absolute error 77.2334 % Root relative squared error 92.3027 % Total Number of Instances 1599

White Wine

Scheme:weka.classifiers.functions.MultilayerPerceptron -L 0.3 -M 0.2 -N 500 -V 0 -S 0 -E 20 -H a Relation: ds1white Instances: 4898 Attributes: 12 Test mode: 10-fold cross-validation Correctly Classified Instances 2706 55.247 % 44.753 % Incorrectly Classified Instances 2192 Kappa statistic 0.2839 0.1601 Mean absolute error Root mean squared error 0.289 Relative absolute error 82.9327 % Root relative squared error 93.0254 % Total Number of Instances 4898

Dataset 2

Red Wine

Scheme: weka.classifiers.functions.MultilayerPerceptron -L 0.3 -M 0.2 -N 500 -V 0 -S 0 -E 20 -H a ds2red_clustered-weka.filters.unsupervised.attribute.Remove-R1 Relation: Instances: Attributes: 12 Test mode: 10-fold cross-validation Correctly Classified Instances 95.7473 % 1531 Incorrectly Classified Instances 68 4.2527 % Kappa statistic 0.9432 Mean absolute error 0.0183 Root mean squared error 0.1047 Relative absolute error 7.3361 % 29.6501 % Root relative squared error Total Number of Instances 1599

White Wine

Scheme: weka.classifiers.functions.MultilayerPerceptron -L 0.3 -M 0.2 -N 500 -V 0 -S 0 -E 20 -H a ds2white_clustered-weka.filters.unsupervised.attribute.Remove-R1 Relation: Instances: 4898 Attributes: 12 Test mode: 10-fold cross-validation 95.0796 % Correctly Classified Instances 4657 Incorrectly Classified Instances 241 4.9204 % Kappa statistic 0.9415 Mean absolute error 0.0195 Root mean squared error 0.1053 8.0972 % Relative absolute error 30.3967 % Root relative squared error Total Number of Instances 4898

Results

The project stated that dataset 1, the unmodified dataset, and dataset 2, the clustered dataset, were to be compared through the use of classifiers. Dataset 2 had it's classification clusters generated algorithmically from scientific data, while dataset 1's classifications where assigned depending each wines sensory information. For this reason we could assume that the classifications in dataset two will be easier to classify, as they are based off of objective data rather than subjective.

The white wine data falls into a bell curve shape, while the majority of red wine classifications fall into two distinct bins. This makes classification hard as the majority of our data is This difference can be seen in Figure 1 and Figure 2.

The information supplied about each wine is incomplete, as it is missing attributes such as grape type, brand and cost that can effect a testers perception[7]. Wines that sell

at a higher price may be percieved as better wines, even though they contain the same physiochemical properties as a cheaper wine. The range of wines are limited to varients of the Portuguese "Vinho Verde" wine and the classification of the wines is subjective. This means that for someone tasting a wine from a different region may see the same physiochemical properties, but classify the wine completely differently.

Figures

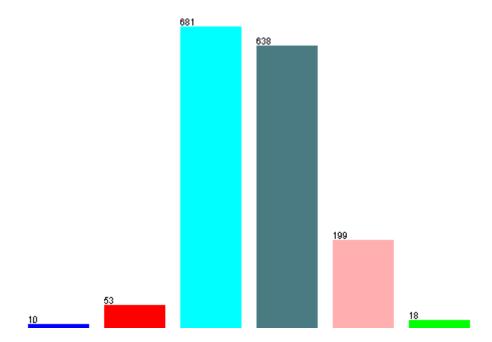


Figure 1: Red Wine Distribution

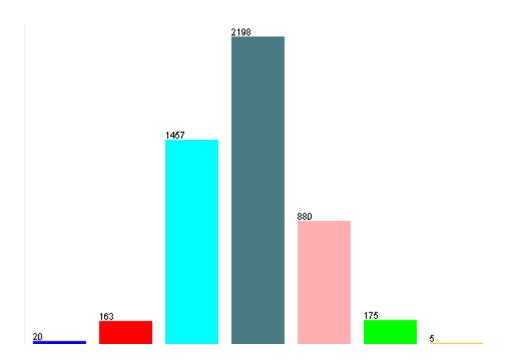


Figure 2: White Wine Distribution

Bibliography

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