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 required clustering before it could 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 different combinations of attributes had very little effect on the clustering and so it did 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. These clusters where then outputted to ds1-red-clustered.arff and ds1-white-clustered.arff for use in classification. Red Wine

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

Both dataset 1 and dataset 2 where processed by three different classifiers, naive bayes, neural networks and support vector machines. The aim was to see how correct each classifier was and how efficiently it performed on different size inputs. To do this, all variables, suck as test mode were kept constant.

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

Root mean squared error 0.3198

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 $\tt ds2white_clustered_weka.filters.unsupervised.attribute.Remove-R1$ Instances: Attributes: 12 Test mode: 10-fold cross-validation 87.1172 % Correctly Classified Instances 4267 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 needs to be 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:
              12
Test mode: 10-fold cross-validation
Correctly Classified Instances
                                                          58.349 %
                                       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
```

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 52.0621 % Correctly Classified Instances 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 -W 1 -K "weka.classifiers.function -E 1.0"			·1		
Relation: ds2red_clustered-weka.filters.unsupervised.attribute.Remove-R1					
Instances: 1599					
Attributes: 12					
Test mode:10-fold cross-validation					
Correctly Classified Instances	1506	94.1839 %			
Incorrectly Classified Instances	93	5.8161 %			
Kappa statistic	0.9215				
Mean absolute error	0.2238				
Root mean squared error	0.3128				
Relative absolute error	89.7115 %				
Root relative squared error	88.5878 %				
more relative refugion error					

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" $\tt ds2white_clustered_weka.filters.unsupervised.attribute.Remove-R1$ Relation: Instances: 4898 Attributes: 12 Test mode: 10-fold cross-validation Correctly Classified Instances 4711 96.1821 % Incorrectly Classified Instances 3.8179 % 187 Kappa statistic 0.9545 Mean absolute error 0.2047 Root mean squared error 0.3021 Relative absolute error 85.1906 % Root relative squared error 87.1569 % 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.MV 0 -S 0 -E 20 -H a	ultilayerPerceptro	on -L 0.3 -M	0.2 -N 500
Relation: ds1red			
Instances: 1599			
Attributes: 12			
Test mode:10-fold cross-validation			
Correctly Classified Instances	967	60.4753	%
Incorrectly Classified Instances	632	39.5247	%
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		

 $Scheme: we ka. classifiers. functions. \verb|Multilayer| Perceptron -L 0.3 -M 0.2 -N 500|$ -V 0 -S 0 -E 20 -H a Relation: ds1white Instances: 4898 Attributes: 12 Test mode:10-fold cross-validation 55.247 % Correctly Classified Instances 2706 Incorrectly Classified Instances 44.753 % 2192 Kappa statistic 0.2839 Mean absolute error 0.1601 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					
Relation: ds2red_clustered-weka.filters.unsupervised.attribute.Remove-R1					
Instances: 1599					
Attributes: 12					
Test mode:10-fold cross-validation					
Correctly Classified Instances	1531	95.7473 %			
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 %				
Root relative squared error	29.6501 %				
Total Number of Instances	1599				

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: 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 Relative absolute error 8.0972 % 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. Dataset 2's results were ignored when analysing the performance of the different classifiers.

For dataset 1, 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 there is very little vairance in the data. 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.

The nueral network classifier gives the best classification rate for dataset 1, and a similar classification as the support vector machine classifier for dataset 2. It takes a significantly for time than the support vector machine however, so for larger datasets both could be seen as valid options. The naive Bayesian classifier gave poor results for dataset 1 as expected from it's relatively simplistic algorithms. Across large data sets, the support vector machine is the most efficient method for classifying the data due to it's higher accuracy and low running time.

Figures

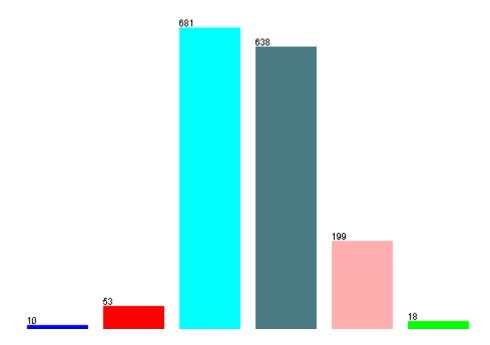


Figure 1: Dataset 1 Red Wine Distribution

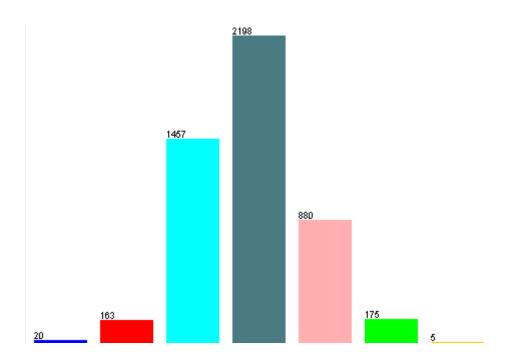


Figure 2: Dataset 1 White Wine Distribution

Bibliography

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