

# 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 were 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 distribution 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 distribution 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 were 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

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

Both dataset 1 and dataset 2 were 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, such 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
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 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      933           58.349 %
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      2550           52.0621 %
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
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 %
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

Correctly Classified Instances      4711                96.1821 %
Incorrectly Classified Instances    187                   3.8179 %
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 than 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
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
```

#### 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 %
Incorrectly Classified Instances    2192           44.753 %
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
```

### White Wine

```
Scheme:weka.classifiers.functions.MultilayerPerceptron -L 0.3 -M 0.2 -N 500
-V 0 -S 0 -E 20 -H a
Relation:      ds2white_clustered-weka.filters.unsupervised.attribute.Remove-R1
Instances:     4898
Attributes:    12
Test mode:10-fold cross-validation

Correctly Classified Instances      4657                95.0796 %
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 %
Root relative squared error          30.3967 %
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 its classification clusters generated by an algorithmic based on scientific data, while **dataset 1**'s classifications were assigned depending on 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 analyzing 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 variance 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 perceived as better wines, even though they contain the same physiochemical properties as a cheaper wine. The range of wines are limited to variants 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 neural network classifier gives the best classification rate for **dataset 1**, and a similar classification as the support vector machine classifier for **dataset 2**. Both could be seen as valid options, however the neural network takes significantly more time so may not be favorable for larger datasets . The naive Bayesian classifier gave poor results for **dataset 1** as expected from its relatively simplistic algorithms. Across large data sets, the support vector machine is the most efficient method for classifying the data due to its higher accuracy and low running time.

## Figures

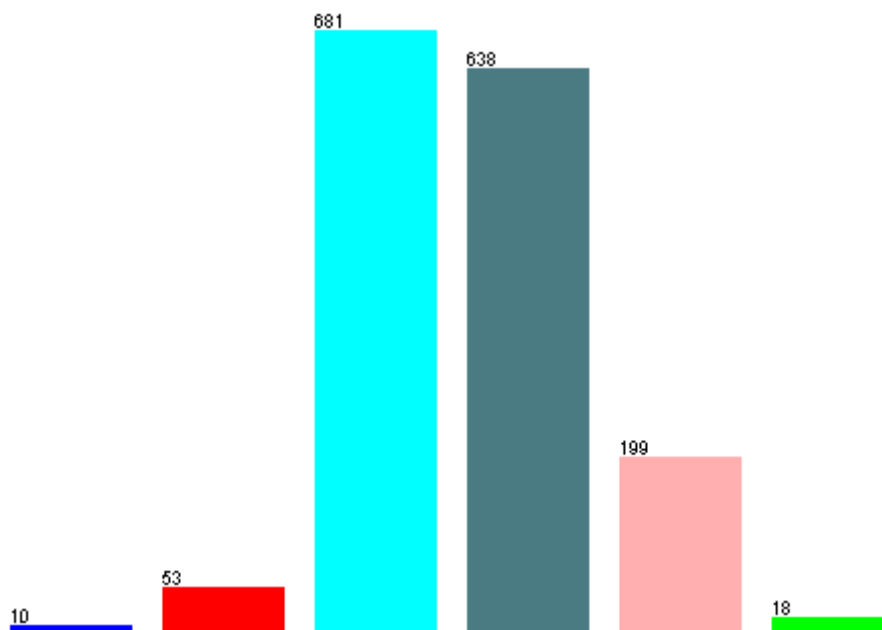


Figure 1: Dataset 1 Red Wine Distribution

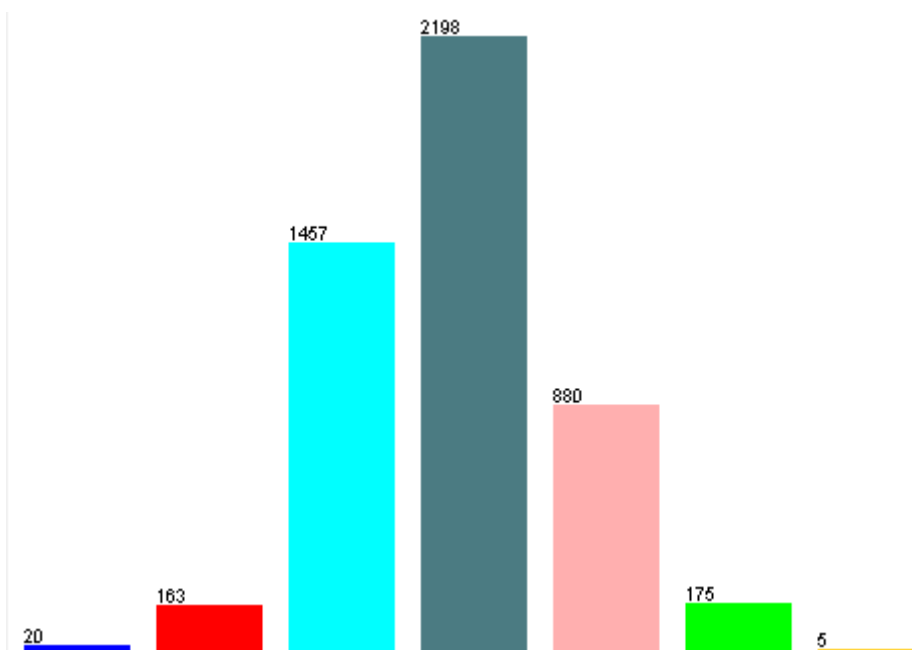


Figure 2: Dataset 1 White Wine Distribution

# Bibliography

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