

# Wine Quality Classification Using Machine Learning

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## Introduction

While there are many definitions of machine learning, in a nutshell, machine learning is a tool for turning data into knowledge. People use machine learning to learn the rules governing the data and use them to make predictions or decisions without explicitly programming to execute the task. In this Case Study, we will analyze and dissect a data set with the goal of correctly classifying wines into their correct quality classification.

## Wine Data Set:

Our group would like to use this powerful tool to make our fair share of predictions. The dataset used in the project is quality of red wine from UC Irvine's Wine Quality Dataset. With 1599 instances and 11 input variables, we believe that the dataset is sufficient for our use. The goal of the project is to determine the quality of the wine using input variables and whether these variables are relevant in determining the quality of the wine in the first place.

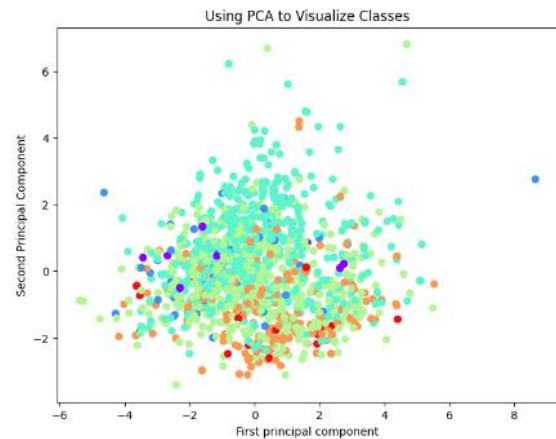


Figure 1 Shows the data visualized.

Our group used four machine learning methods: Random Forest Classifier, Support Vector Machine, Logistic Regression, and Artificial Neural Network. We provide a brief background on each method below.

## Algorithms Used:

**Random Forest Classifier:** is an ensemble learning method that consists of multiple decision trees. The decision trees are trained on a slightly different set of observations and the nodes in each tree are split considering a limited number of features. The final predictions of the random forest classifier are made by averaging the predictions of each individual tree.

**Support Vector Machines:** are a popular type of supervised learning algorithm used

in machine learning for classification and regression tasks. SVM works by finding the best possible hyperplane that separates different classes of data. A hyperplane is a decision boundary that helps to classify the input data into different classes. In SVM, the hyperplane that best separates the two classes is the one that maximizes the margin between the classes. The margin is the distance between the hyperplane and the closest data points from both classes. SVM can also use a kernel trick to transform the input data into a higher-dimensional space, where it may become easier to separate the data into classes. SVM has many applications in the fields of image recognition, bioinformatics, text classification, and more.

**Artificial neural network** is a computing system vaguely inspired by the biological neural networks of animal brains. It uses multiple layers of perceptrons which can be categorized into input layer, hidden layers, and output layer. Input layer nodes transmit input values to the hidden layer nodes without performing any computations. Each hidden layer node then computes the weighted sum of its inputs to form a scalar net activation. Then they output the nonlinear function of the net activation. After a single or multiple hidden layers, the output node computes the net activation based on hidden node outputs. Each output node emits the nonlinear function of the net activation.

**Multi-Class Logistic Regression:** is a popular statistical method that is used to analyze and model relationships between multiple independent variables and a categorical dependent variable. It is a technique used in machine learning, data science, and predictive analytics to build

models that classify data into multiple classes. While we did use this model, we did not write an analysis on it as it seemed redundant.

Using these methods and tuning the hyperparameters of each model, we were able to predict the quality of wine with reasonable accuracy.

## Methods

All the methods used will be expressed in detail as well as the steps taken to classify the data.

### Exploratory Data Analysis:

Before starting to manipulate the data, it's often helpful to graph the data to see if there are strong between certain classes in the data, this can be seen by the correlation matrix in Figure 2.

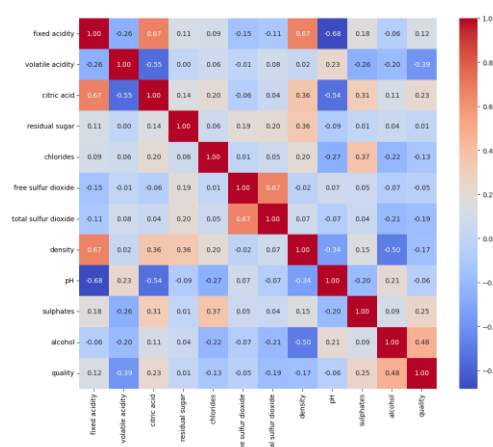


Figure 2 Shows the correlation matrix of all classes.

We also performed Principal Component Analysis, or PCA, to reduce the dimensionality of the dataset and visualize them in a plot for better understanding of how our data is structured. Figure 3 shows the data visualization using PCA.

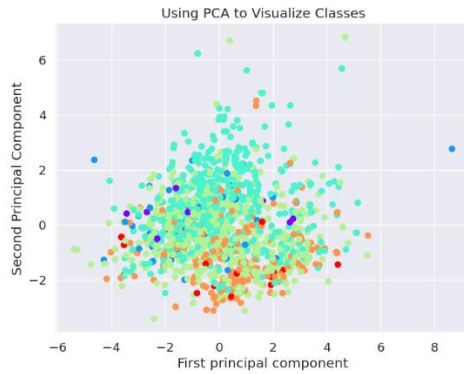


Figure 3 Visualization using PCA.

### Data Pre-Processing:

Before the data set can be used, it must be inspected and deemed fit to use with the desired machine learning algorithms.

Data Pre-Processing is often very useful in increasing the accuracy and reliability of machine learning models. The first steps we took in cleaning our data were to check for and remove any null values, separate the input variables from the output variable, and rescale the data to more optimal ranges when training machine learning models. We also considered the significance of features with the least predictive capabilities.

An effective way to trim down the number of features we are working with, which can be beneficial for both computation times and the generalization of models like the ANN and SVM, is to select the k best features. There are many ways to score the k best features, but a popular method is comparing the ratio of explained variance to unexplained variance, or, in other words, the ANOVA F-value. We used this technique through sklearn to remove the four features with the lowest scores from our dataset: density, pH, and chlorides.

	fixed acidity	volatile acidity	citric acid
0	7.4	0.700	0.00
1	7.8	0.880	0.00
2	7.8	0.760	0.04
3	11.2	0.280	0.56
4	7.4	0.700	0.00

Figure 4 Shows the first three features of the data set.

### Class Imbalance:

Class imbalance refers to when some classes in the data set have much lower representation, as seen in Figure 2, classes 3, 4, 7, & 8 have much lower instances than those of 5 & 6. This will cause the algorithms to be very biased towards classes 5 & 6 and ignore the other classes for the most part. It's also important to note that the data set gives the wine a quality score between 0 and 10 but the data set has no representation for qualities 0, 1, 2, 9 & 10, which is problematic.

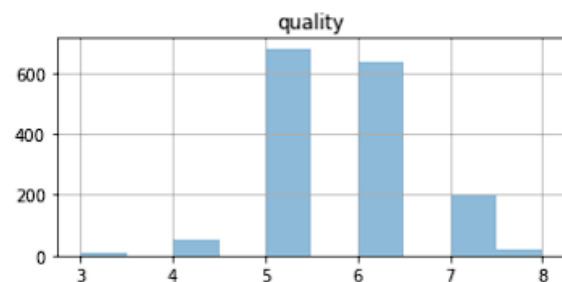


Figure 5 Shows how unbalanced the data is.

### Solutions To Class Imbalance:

There are many ways to deal with Class Imbalance as it's a common problem in Machine Learning and data processing. Several strategies were explored below.

### Oversampling Minority Class:

By copying the small amount of data, we have for the minority class and duplicating them, they are more represented, but the downside is if too many copies are made, the model may be overfit to those minority instances.

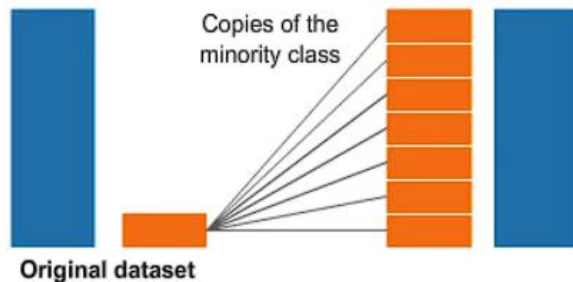


Figure 6 Diagram showing how oversampling works.

### Under Sampling Majority Class:

By leaving out some of the instances of the major classes, the classes become more balanced, but the downfall is useful data is lost.

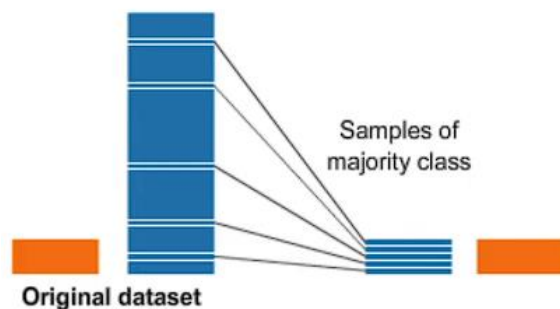


Figure 7 Diagram showing how under sampling works.

### Synthetic Minority Oversampling (SMOTE):

Instead of just duplicating the minority class to increase the number of instances, synthetic instances can be used by using the nearest neighbors approach. The downside is that it is assumed that the correlation between the different features is relatively linear.

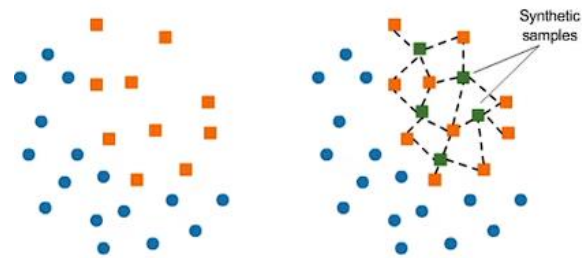


Figure 8 Graphic showing synthetic instances.

### Data Imputation:

Although it is a common practice to identify missing values in a dataset and replace them with a numeric value, what we are trying is creating new data for classes with no other instances available. Using the mean value of each input variable, we created one instance for each of the missing classes.

total sulfur dioxide	sulphates	alcohol	quality
46.467792	0.658149	10.422983	0
46.467792	0.658149	10.422983	1
46.467792	0.658149	10.422983	2
46.467792	0.658149	10.422983	9
46.467792	0.658149	10.422983	10

Figure 9 Table shows the instances added for classes 0,1,2,9, & 10

### Hyperparameter Tuning:

With all the algorithms used in this case study, several hyperparameters need to be chosen for the algorithm to run on. Instead of choosing the parameters through trial and error, grid search was used for an exhaustive sampling of the hyperparameter space to find the optimal results. We usually use accuracy as a measure of a model's predictive score.

## Random Forest Classifier (RFC):

Hyper-parameter tuning for the random forest classifier algorithm was done in tandem with training the random forest classifier with the following datasets: the trimmed version, the original dataset, and the SMOTE dataset. While we could have relied on sklearn's built in oop-scoring, which generates out-of-bag samples to test with, for this machine learning model we chose to create a validation set before creating three versions of the hyper-parameter tuning set: two modified hyper-parameter training sets and the original (with all 11 features). Get picture of one of the decision trees:

This allowed for more consistent testing methods, as we were able to validate the scores of the trained models using the same validation set. We manually coded in that the final model would use the original data set after seeing the results of the testing on the validation set, which used the entire training data set, including the validation set to get as many minority classifications as possible in the training set. Here are some graphs displaying the trend of hyper-parameters around the local maxima for score of the random forest classifier trained on the original training set.

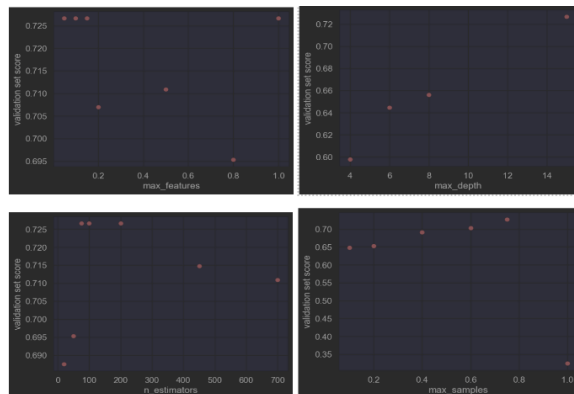


Figure 10 Shows the optimal parameters for RFC.

## Support Vector Machines (SVM):

It's important to note that for each data set, the hyperparameters are not necessarily the same, and that for the majority of data set types, c is at 1000 meaning that the algorithm will choose a smaller-margin hyperplane if that will cause more of the training instances to be classified correctly. This can clearly be seen in Figure 10.

```
Regular Data : {'C': 1, 'kernel': 'rbf'}  
best score: 0.6067493872549019  
Oversampled Data : {'C': 1000, 'kernel': 'rbf'}  
best score: 0.6845814977973569  
Undersampled Data : {'C': 1000, 'kernel': 'poly'}  
best score: 0.6875210694565533  
SMOTE Data : {'C': 1000, 'kernel': 'rbf'}  
best score: 0.683311836194797  
Imputed Data : {'C': 1000, 'kernel': 'rbf'}  
best score: 0.5710942367601245
```

Figure 11 Shows the optimal parameters for SVM.

## Artificial Neural Networks (ANN):

ANNs were implemented using the scikit-learn Package in python. While the Sklearn Package computes the mathematics behind the ANN algorithm, the user must still choose the hyperparameters for the algorithm, these hyperparameters are the activation function to be used, the solver, the learning rate, and the maximum number of iterations. As can be seen in Figure 11, as the data was transformed from the original data set, the hidden layer sizes tripled and the max\_iter also significantly grew. This shows that changing the dataset muddled the relationship a bit to the point that now it takes significantly more resources for the algorithm to run but the accuracy was over all better.

```
Regular Data : {'hidden_layer_sizes': (100,), 'learning_rate_init': 0.01, 'max_iter': 100}  
best score: 0.6106556372549019  
Oversampled Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.01, 'max_iter': 500}  
best score: 0.6867841409691631  
Undersampled Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.001, 'max_iter': 500}  
best score: 0.702615518744551  
SMOTE Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.01, 'max_iter': 500}  
best score: 0.7214946446687596  
Imputed Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.001, 'max_iter': 300}  
best score: 0.5691958722741433
```

Figure 12 Shows the optimal parameters for ANN.



## Results & Analysis:

After running the methods explained above, the results are analyzed, and conclusions are made about the data.

### Random Forest Classifier (RFC):

I found that the scores of the model trained on the original training set scored better on the validation set overall. This did not surprise me because of the nature of the hyper-parameter tuning.

The random forest classifier models allow for tuning of the power/generalization capabilities, and I thought that the grid-search would find the most optimal amount of generalization on its own. Also, with random forest classifier you can tune the depth of the trees, which would practically have the same effect as removing the features with the lowest predictive power.

It makes sense that the random forest classifier trained on the original data set is able to achieve the highest accuracy score out of the three, because the red wine data set is far from linearly separable. We managed to get an accuracy score of .7 on the final test set, but this could be an anomaly, and an accuracy of at least .65 is more reliable for our trained model. See the appendix for more details on the score.

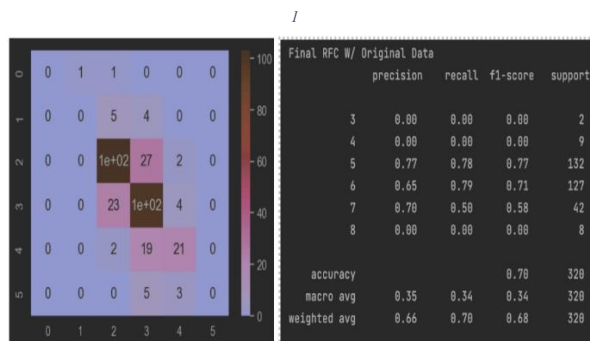


Figure 13 Final RFC: Final RFC Results

```
Regular Data : 0.65625
Oversampled Data : 0.475
Undersampled Data : 0.2625
SMOTE Data : 0.359375
Imputed Data : 0.359375
```

Figure 14 RFC baseline results tested on regular data.

```
Regular Data : 0.6375
Oversampled Data : 0.748898678414097
Undersampled Data : 0.7419354838709677
SMOTE Data : 0.7481662591687042
Imputed Data : 0.40923076923076923
```

Figure 15 RFC results after hyperparameter tuning.

### Support Vector Machines (SVM):

We managed to achieve a respectable accuracy of around .6 with the support vector machines model. The kernel “rbf” was by far the best kernel method when testing on the validation set. This is likely because it is more powerful than the other methods, and our data seems to require a powerful model.

SVM tested best on the validation set with the oversampled data, but this was very close, and overall, it seems like the less we touched the data set, the better it did on the validation set. This is likely because of how the GridSearchCV function evaluates its training scores, which leads to each modified training set overfitting their own data set, rather than being able to generalize on to the original validation set.

```
Regular Data : 0.6
Oversampled Data : 0.00625
Undersampled Data : 0.396875
SMOTE Data : 0.00625
Imputed Data : 0.4125
```

Figure 16 SVM baseline results.

```
Regular Data : 0.6
Oversampled Data : 0.7268722466960352
Undersampled Data : 0.7311827956989247
SMOTE Data : 0.8092909535452323
Imputed Data : 0.40615384615384614
```

*Figure 17 SVM results after hyperparameter tuning.*

### Artificial Neural Networks (ANN):

The ANN algorithm was run with both the original data set as well as the processed datasets to determine what gives the best results. Using the optimal hyperparameters calculated in the above section, the regular data performed the best as can be seen in Figure 19.

```
Regular Data : 0.6
Oversampled Data : 0.303125
Undersampled Data : 0.23125
SMOTE Data : 0.234375
Imputed Data : 0.296875
```

*Figure 18 ANN baseline results.*

If, however, the test data was processed in the same way that the training data was, then it can be seen as shown in Figure 20, that there is very significant model performance improvement compared to the regular data. The SMOTE data performed the best with an accuracy of about 84%, which is very good.

```
Regular Data : 0.578125
Oversampled Data : 0.7290748898678414
Undersampled Data : 0.7150537634408602
SMOTE Data : 0.8398533007334963
Imputed Data : 0.12923076923076923
```

*Figure 19 ANN results after hyperparameter tuning.*

### General Analysis:

In general, it seems that the best model was the random forest classifier trained on the original data set.

The first likely cause is that the score used to tune the hyper-parameters was much more representative of the actual final testing data. Not because the data set is the least altered, but because with the manual grid search, which we implemented only for random forest classifier, we were able to test and score each set of hyperparameters on a validation set much more representative of the test set than the scores generated by the GridSearchCV function.

A second, less interesting cause, is that the random forest classifier model is just better with these types of data sets. As we saw, the red wine data set is far from linearly separable, and its minority classes are very small. A random forest classifier model could have an advantage if its training set represented the test set well (which it does) when there are minority classes as small as these. Its training algorithm does not rely on the gradient search method, which could be why it is better at handling minority cases.

### Conclusion:

The red wine data set is very difficult to accurately classify, but with powerful machine learning models, we were able to get some respectable results. While the random forest classifier model had the best training results, that is only if the actual population follows the data set. If, for example, our data set is misrepresentative of the actual population, then perhaps the models trained on some of our modified data sets, like SMOTE, would be better at predicting results from outside our data set. But, with the information and data we have, modifying the data set did not significantly benefit the accuracy of our results.

## Contributions:

### **Seif Elkhatab:**

- Programmed Neural Networks.
- Helped Program SMOTE.
- Wrote ANN in report.
- Wrote Background on Algorithms.
- Wrote Class imbalance section.

### **Ben Ko:**

- Programmed Hyper tuning.
- Programmed Data Preprocessing.
- Programmed MLR and SVM.
- Programmed Data Transformations.
- Wrote the introduction.

### **Ben Watkins:**

- Worked on Manual Grid search for RFC and RFC coding.
- Helped with editing and writing reports, especially RFC related sections.

## Appendix:



```
In [ ]: import matplotlib.pyplot as plt
import pandas as pd
from sklearn.decomposition import PCA
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import confusion_matrix, classification_report
from sklearn.model_selection import GridSearchCV
from sklearn.neural_network import MLPClassifier
from sklearn.feature_selection import SelectKBest, chi2
from collections import Counter
import seaborn as sns
from imblearn.over_sampling import RandomOverSampler, SMOTE
from imblearn.under_sampling import RandomUnderSampler
import warnings
warnings.filterwarnings("ignore")
```

## 1. Data Preprocessing

### Importing dataset & Checking for missing data

```
In [ ]: headerList = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density', 'pH', 'sulphates',
                        'residual dry matter']

wineData = pd.read_csv('winequality-red.csv', header = 0, names = headerList, sep=";")
print(wineData.head(10))
print(wineData)

#Summarative functions
wineData.dtypes
wineData.describe()
wineData.info()

#Plotting histogram of each variable
wineData.hist(alpha=0.5, figsize=(15, 10))
plt.tight_layout()
plt.show()

for h in headerList:
    wineData[h] = pd.to_numeric(wineData[h], errors='coerce')

print("\nChecking for null values: \n")
wineData.isna().sum()
wineData = wineData.fillna(0)
print("\nChecking for null values after using fillna(): \n")
wineData.isna().sum()
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
0	7.4	0.70	0.00	1.9	0.076
1	7.8	0.88	0.00	2.6	0.098
2	7.8	0.76	0.04	2.3	0.092
3	11.2	0.28	0.56	1.9	0.075
4	7.4	0.70	0.00	1.9	0.076
5	7.4	0.66	0.00	1.8	0.075
6	7.9	0.60	0.06	1.6	0.069
7	7.3	0.65	0.00	1.2	0.065
8	7.8	0.58	0.02	2.0	0.073
9	7.5	0.50	0.36	6.1	0.071

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates \
0	11.0	34.0	0.9978	3.51	0.56
1	25.0	67.0	0.9968	3.20	0.68
2	15.0	54.0	0.9970	3.26	0.65
3	17.0	60.0	0.9980	3.16	0.58
4	11.0	34.0	0.9978	3.51	0.56
5	13.0	40.0	0.9978	3.51	0.56
6	15.0	59.0	0.9964	3.30	0.46
7	15.0	21.0	0.9946	3.39	0.47
8	9.0	18.0	0.9968	3.36	0.57
9	17.0	102.0	0.9978	3.35	0.80

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5
5	9.4	5
6	9.4	5
7	10.0	7
8	9.5	7
9	10.5	5

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
0	7.4	0.700	0.00	1.9	0.076
1	7.8	0.880	0.00	2.6	0.098
2	7.8	0.760	0.04	2.3	0.092
3	11.2	0.280	0.56	1.9	0.075
4	7.4	0.700	0.00	1.9	0.076
...	...	...	...	...	...
1594	6.2	0.600	0.08	2.0	0.090
1595	5.9	0.550	0.10	2.2	0.062
1596	6.3	0.510	0.13	2.3	0.076
1597	5.9	0.645	0.12	2.0	0.075
1598	6.0	0.310	0.47	3.6	0.067

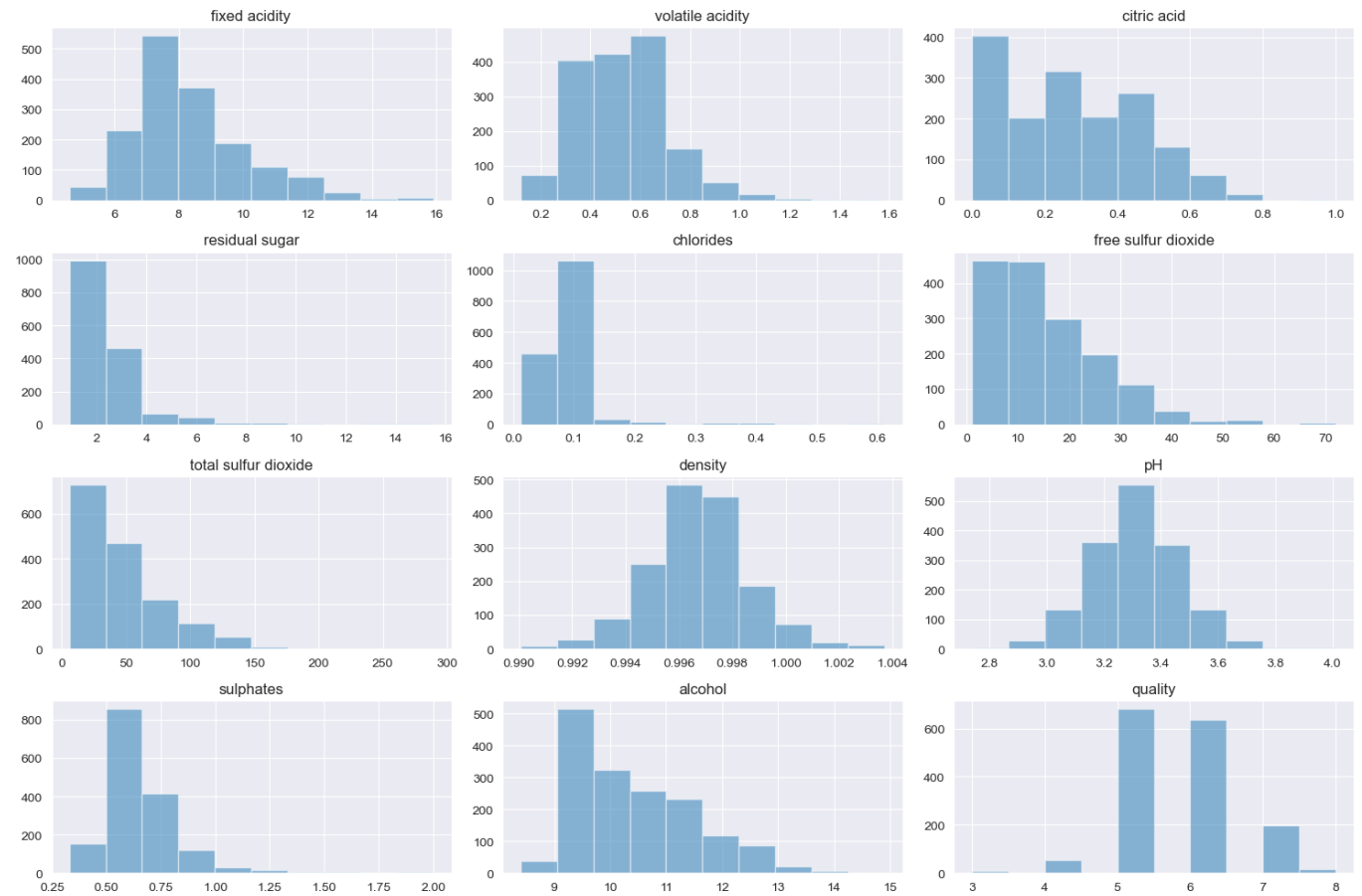
	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates \
0	11.0	34.0	0.99780	3.51	0.56
1	25.0	67.0	0.99680	3.20	0.68
2	15.0	54.0	0.99700	3.26	0.65
3	17.0	60.0	0.99800	3.16	0.58
4	11.0	34.0	0.99780	3.51	0.56
...	...	...	...	...	...
1594	32.0	44.0	0.99490	3.45	0.58
1595	39.0	51.0	0.99512	3.52	0.76
1596	29.0	40.0	0.99574	3.42	0.75
1597	32.0	44.0	0.99547	3.57	0.71
1598	18.0	42.0	0.99549	3.39	0.66

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5
...	...	...
1594	10.5	5
1595	11.2	6
1596	11.0	6
1597	10.2	5
1598	11.0	6

```
[1599 rows x 12 columns]
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1599 entries, 0 to 1598
Data columns (total 12 columns):
```

#	Column	Non-Null Count	Dtype
0	fixed acidity	1599 non-null	float64
1	volatile acidity	1599 non-null	float64
2	citric acid	1599 non-null	float64
3	residual sugar	1599 non-null	float64
4	chlorides	1599 non-null	float64
5	free sulfur dioxide	1599 non-null	float64
6	total sulfur dioxide	1599 non-null	float64
7	density	1599 non-null	float64
8	pH	1599 non-null	float64
9	sulphates	1599 non-null	float64
10	alcohol	1599 non-null	float64
11	quality	1599 non-null	int64

```
dtypes: float64(11), int64(1)
memory usage: 150.0 KB
```



Checking for null values:

Checking for null values after using fillna():

```
Out[ ]:
fixed acidity      0
volatile acidity   0
citric acid        0
residual sugar     0
chlorides          0
free sulfur dioxide 0
total sulfur dioxide 0
density            0
pH                0
sulphates         0
alcohol           0
quality           0
dtype: int64
```

Our data is imbalanced as seen from the histogram. We will adapt multiple strategies to address the issue.

## 2. Exploratory Data Analysis

### 1. Principal Component Analysis (PCA)

```
In [ ]:
#PCA
df_pca = wineData.copy()
X_pca = df_pca.loc[:, 'fixed acidity':'alcohol']
y_pca = df_pca['quality']

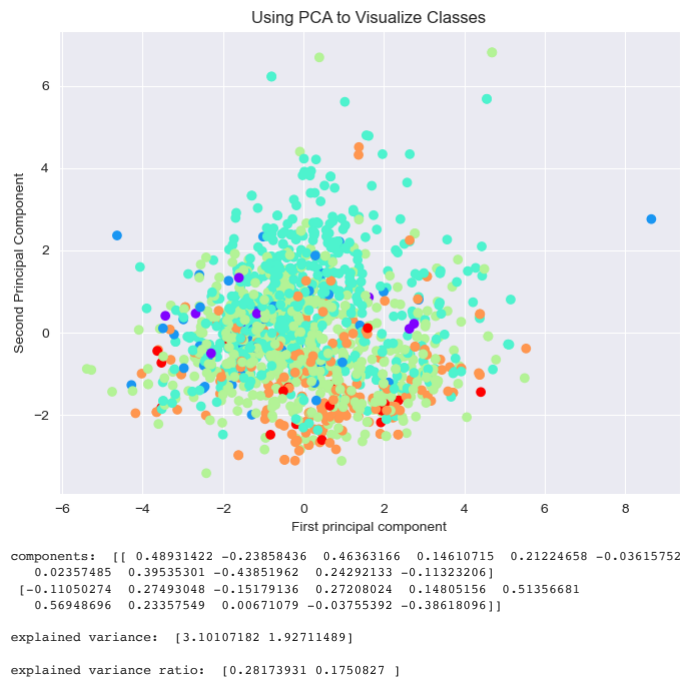
X_pca.tail()
X_pca = StandardScaler().fit_transform(X_pca)

#Fit PCA
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X_pca)

X_pca.shape

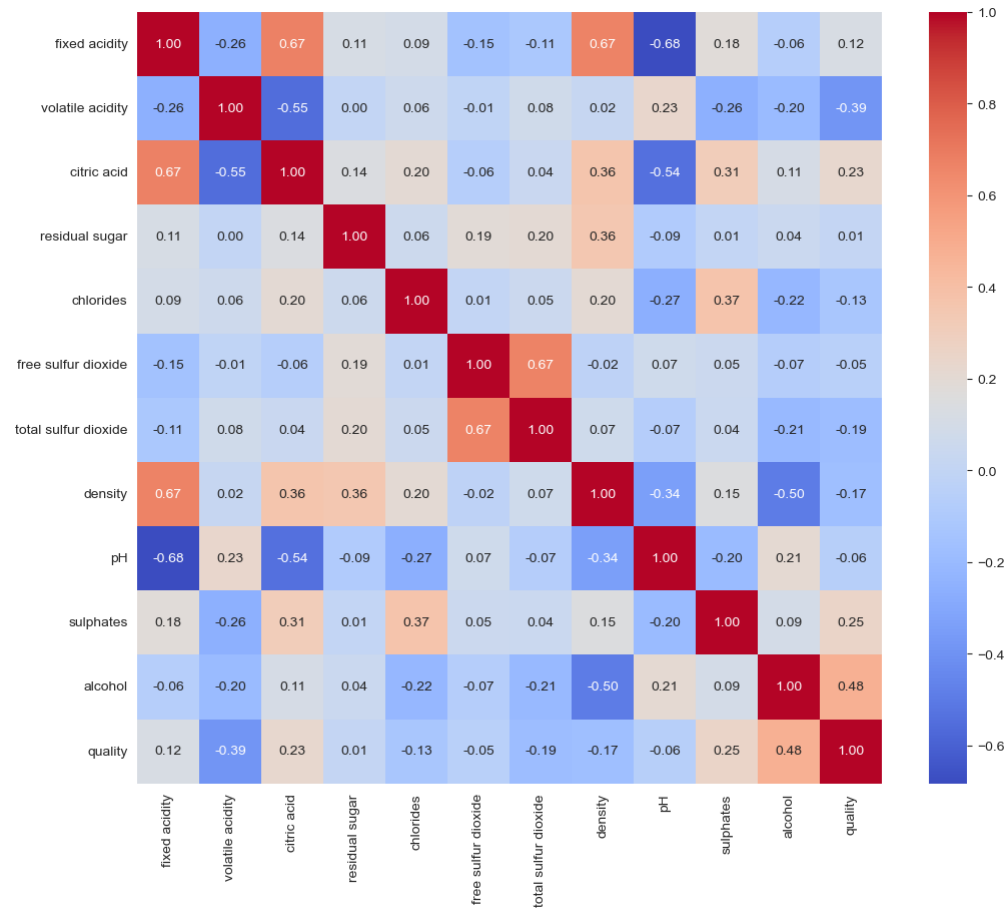
plt.figure(figsize=(8,6))
plt.scatter(X_pca[:,0],X_pca[:,1],c=y_pca,cmap='rainbow')
plt.xlabel('First principal component')
plt.ylabel('Second Principal Component')
plt.title("Using PCA to Visualize Classes")
plt.show()

print("components: ", pca.components_, "\n")
print("explained variance: ", pca.explained_variance_, "\n")
exp_var_rat = pca.explained_variance_ratio_
print("explained variance ratio: ", exp_var_rat)
```



## 2. Correlation Matrix

```
In [ ]: plt.figure(figsize=(12,10))
sns.heatmap(wineData.corr(),annot=True, cmap='coolwarm',fmt='.2f')
Out [ ]: <AxesSubplot:>
```



## 3. Univariate Selection

```
In [ ]: #Split data into training and test sets
x = wineData.loc[:, 'fixed acidity':'alcohol']
y = wineData['quality']

# apply SelectKBest class to extract best features
bestFeatures = SelectKBest(score_func=chi2, k=11)
bestFeaturesFit = bestFeatures.fit(X,y)
dfscores = pd.DataFrame(bestFeaturesFit.scores_)
```

```
dfcolumns = pd.DataFrame(X.columns)

# concatenate scores with predictor names
predScores = pd.concat([dfcolumns,dfscores],axis=1)
predScores.columns = ['Predictor','Score']
print(predScores.nlargest(11,'Score'))
```

	Predictor	Score
6	total sulfur dioxide	2755.557984
5	free sulfur dioxide	161.936036
10	alcohol	46.429892
1	volatile acidity	15.580289
2	citric acid	13.025665
0	fixed acidity	11.260652
9	sulphates	4.558488
3	residual sugar	4.123295
4	chlorides	0.752426
8	pH	0.154655
7	density	0.000230

## Dropping features from univariate selection

We are dropping bottom features as they have very low predictor scores and to save computation

```
In [ ]: #save the original data
XOriginal = wineData.loc[:, 'fixed acidity':'alcohol']
yOriginal = wineData['quality']

#Drop the bottom four features (smallest score)
wineData = wineData.drop(['density'], axis=1)
wineData = wineData.drop(['pH'], axis=1)
wineData = wineData.drop(['chlorides'], axis=1)
print(wineData)

X = wineData.loc[:, 'fixed acidity':'alcohol']
y = wineData['quality']
```

	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	7.4	0.700	0.00	1.9	
1	7.8	0.880	0.00	2.6	
2	7.8	0.760	0.04	2.3	
3	11.2	0.280	0.56	1.9	
4	7.4	0.700	0.00	1.9	
...	...	...	...	...	
1594	6.2	0.600	0.08	2.0	
1595	5.9	0.550	0.10	2.2	
1596	6.3	0.510	0.13	2.3	
1597	5.9	0.645	0.12	2.0	
1598	6.0	0.310	0.47	3.6	

	free sulfur dioxide	total sulfur dioxide	sulphates	alcohol	quality
0	11.0	34.0	0.56	9.4	5
1	25.0	67.0	0.68	9.8	5
2	15.0	54.0	0.65	9.8	5
3	17.0	60.0	0.58	9.8	6
4	11.0	34.0	0.56	9.4	5
...	...	...	...	...	...
1594	32.0	44.0	0.58	10.5	5
1595	39.0	51.0	0.76	11.2	6
1596	29.0	40.0	0.75	11.0	6
1597	32.0	44.0	0.71	10.2	5
1598	18.0	42.0	0.66	11.0	6

[1599 rows x 9 columns]

## Data for Random Forest Classifier

```
In [ ]: #Create The Base Dataset
from collections import Counter
from imblearn.over_sampling import RandomOverSampler, SMOTE

# 1. Resized Data (less traits)
X_train_RFC, X_test_RFC, y_train_RFC, y_test_RFC = train_test_split(X, y, test_size = .2, random_state=10) #split the data
X_train_RFC.shape, y_train_RFC.shape, X_test_RFC.shape, y_test_RFC.shape
scaledData = StandardScaler()
X_train_RFC = scaledData.fit_transform(X_train_RFC)
X_test = scaledData.transform(X_test_RFC)
X_train_HP, X_test_HP, y_train_HP, y_test_HP = train_test_split(X_train_RFC, y_train_RFC, test_size = .2, random_state=11)

#Add in the Original Data set
# 2. Original Data (All traits) Using the same random state is essential to getting two identical matrices
X_train_Original, X_test_Original, y_train_Original, y_test_Original = train_test_split(XOriginal, yOriginal, test_size = .2, random_state=10) #split the data
X_train_Original.shape, y_train_Original.shape, X_test_Original.shape, y_test_Original.shape
scaledData = StandardScaler()
X_train_Original = scaledData.fit_transform(X_train_Original)
X_test_Original = scaledData.transform(X_test_Original)
```

## Addressing Imbalance in Class

### First Strategy: Oversampling minority class

```
In [ ]: oversample = RandomOverSampler(sampling_strategy='minority')
X_over, y_over = oversample.fit_resample(X, y)
print("Before RandomOverSampler : ", Counter(y))
print("After RandomOverSampler : ", Counter(y_over))

Before RandomOverSampler : Counter({5: 681, 6: 638, 7: 199, 4: 53, 8: 18, 3: 10})
After RandomOverSampler : Counter({5: 681, 3: 681, 6: 638, 7: 199, 4: 53, 8: 18})
```

### Second Strategy: Undersampling majority class

```
In [ ]: undersample = RandomUnderSampler(sampling_strategy='majority')
X_under, y_under = undersample.fit_resample(X, y)
print("Before RandomUnderSampler : ", Counter(y))
print("After RandomUnderSampler : ", Counter(y_under))

Before RandomUnderSampler : Counter({5: 681, 6: 638, 7: 199, 4: 53, 8: 18, 3: 10})
After RandomUnderSampler : Counter({6: 638, 7: 199, 4: 53, 8: 18, 3: 10, 5: 10})
```

### Third Strategy: SMOTE

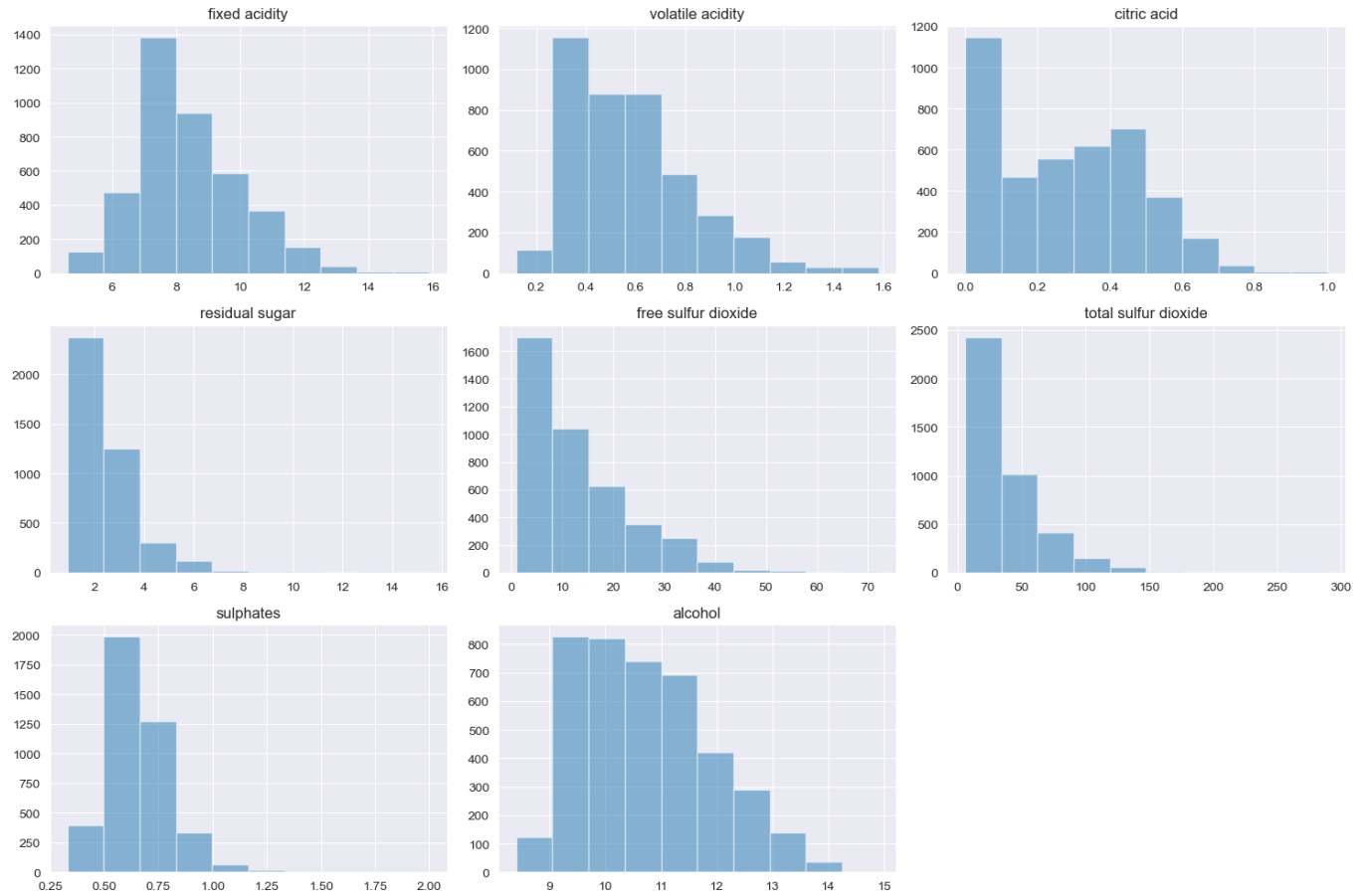
```
In [ ]: smoteOversample = SMOTE()
X_smote, y_smote = smoteOversample.fit_resample(X, y)

#Plotting histogram of each variable
X_smote.hist(alpha=0.5, figsize=(15, 10))

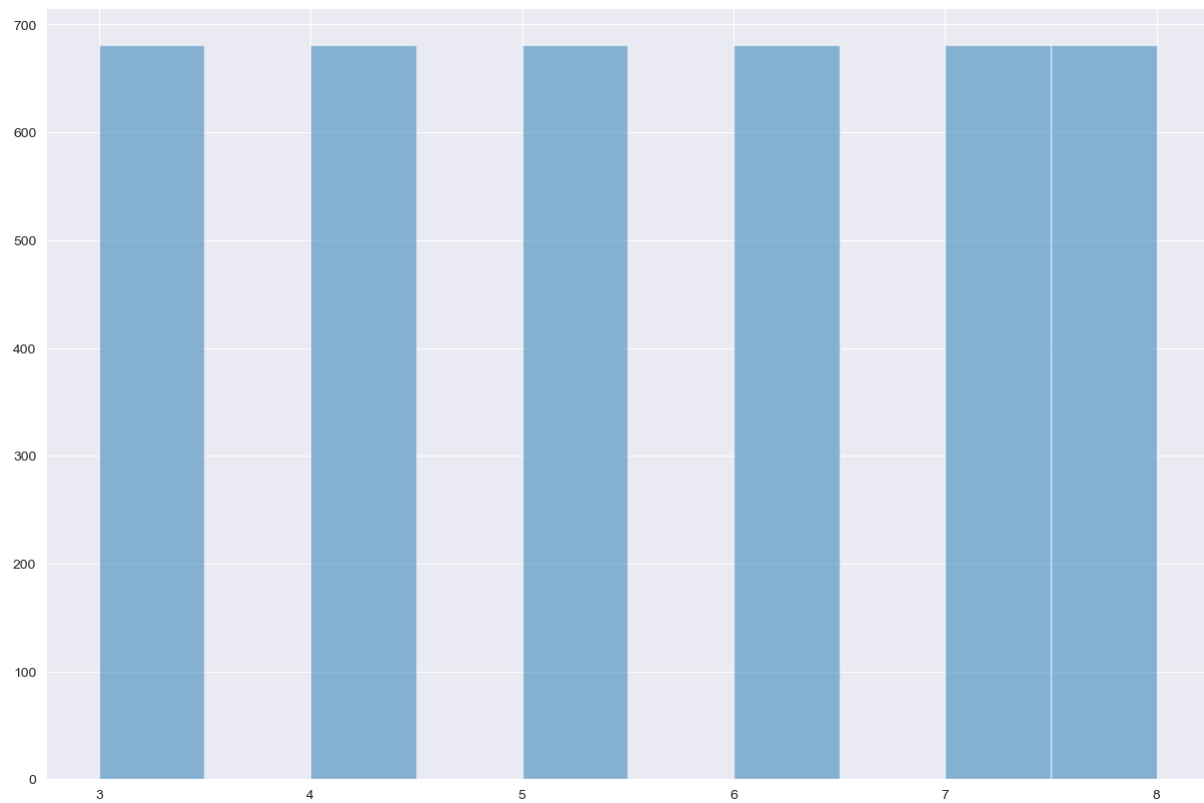
plt.tight_layout()
plt.show()

y_smote.hist(alpha=0.5, figsize=(15, 10))
plt.show()

from collections import Counter
print("Before SMOTE : ", Counter(y))
print("After SMOTE : ", Counter(y_smote))
```







## Fourth Strategy: Data Imputation

Filling in data from missing classes - 0, 1, 2, 9, & 10 with fraud data

```
In [ ]: avgX = X.mean(axis=0)
dfImpute = pd.DataFrame(
    [[avgX[0], avgX[1], avgX[2], avgX[3], avgX[4], avgX[5], avgX[6], avgX[7], 0],
     [avgX[0], avgX[1], avgX[2], avgX[3], avgX[4], avgX[5], avgX[6], avgX[7], 1],
     [avgX[0], avgX[1], avgX[2], avgX[3], avgX[4], avgX[5], avgX[6], avgX[7], 2],
     [avgX[0], avgX[1], avgX[2], avgX[3], avgX[4], avgX[5], avgX[6], avgX[7], 9],
     [avgX[0], avgX[1], avgX[2], avgX[3], avgX[4], avgX[5], avgX[6], avgX[7], 10]],
    columns=['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'free sulfur dioxide', 'total sulfur dioxide', 'sulphates', 'alcohol', '
')

dfImpute

X_add = dfImpute.loc[:, 'fixed acidity':'alcohol']
y_add = dfImpute['quality']
X_impute = pd.concat([X, X_add])
y_impute = pd.concat([y, y_add])
```

## 3. Comparing Machine Learning Models

### Modeling - Final data preparations

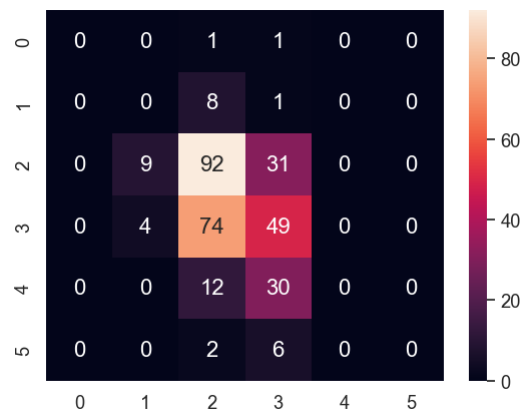
```
In [ ]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = .2, random_state=10) #split the data
X_train.shape, y_train.shape, X_test.shape, y_test.shape
scaledData = StandardScaler()
X_train = scaledData.fit_transform(X_train)
X_test = scaledData.transform(X_test)
```

```
In [ ]: def clas_report(X_train, y_train, x_test, y_test, model, title):
    model.fit(X_train, y_train)
    y_pred = model.predict(x_test)
    cm = confusion_matrix(y_test, y_pred)
    df_cm = pd.DataFrame(cm)
    sns.set(font_scale=1.2) # for label size
    sns.heatmap(df_cm, annot=True, annot_kws={"size": 16}) # font size
    plt.show()
    clas = classification_report(y_test, y_pred)
    print(title, "\n", clas)

def report(X_train, y_train, X_test, y_test, model, title):
    model.fit(X_train, y_train)
    y_pred = model.predict(X_test)
    print(title, "\n", model.score(X_test, y_test))
```

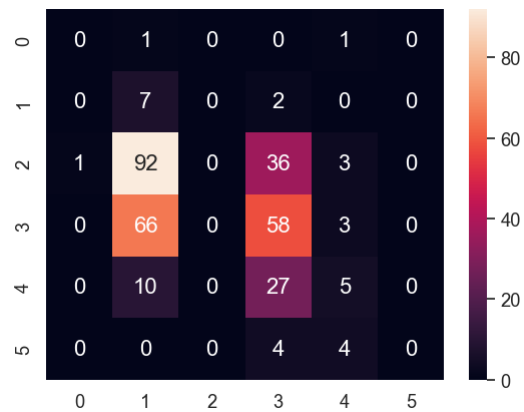
### 1. Random Forest Classifier

```
In [ ]: wineRF = RandomForestClassifier()
clas_report(X_over, y_over, X_test, y_test, wineRF, 'Oversampled Data')
clas_report(X_under, y_under, X_test, y_test, wineRF, 'Undersampled Data')
clas_report(X_smote, y_smote, X_test, y_test, wineRF, 'SMOTE Data')
clas_report(X_impute, y_impute, X_test, y_test, wineRF, 'Imputed Data')
```



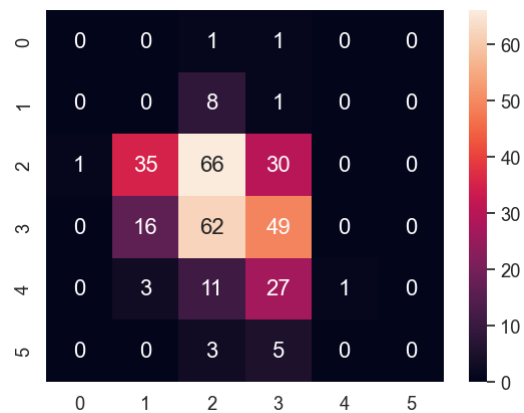
Oversampled Data

	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.49	0.70	0.57	132
6	0.42	0.39	0.40	127
7	0.00	0.00	0.00	42
8	0.00	0.00	0.00	8
accuracy			0.44	320
macro avg	0.15	0.18	0.16	320
weighted avg	0.37	0.44	0.40	320



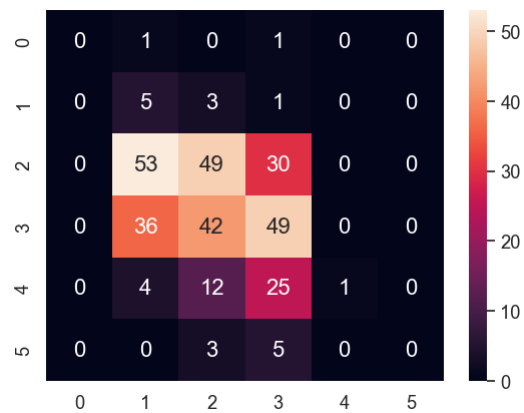
Undersampled Data

	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.04	0.78	0.08	9
5	0.00	0.00	0.00	132
6	0.46	0.46	0.46	127
7	0.31	0.12	0.17	42
8	0.00	0.00	0.00	8
accuracy			0.22	320
macro avg	0.13	0.23	0.12	320
weighted avg	0.22	0.22	0.21	320



SMOTE Data

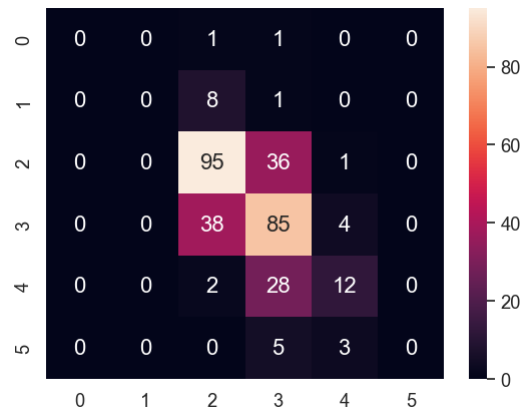
	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.44	0.50	0.47	132
6	0.43	0.39	0.41	127
7	1.00	0.02	0.05	42
8	0.00	0.00	0.00	8
accuracy			0.36	320
macro avg	0.31	0.15	0.15	320
weighted avg	0.48	0.36	0.36	320



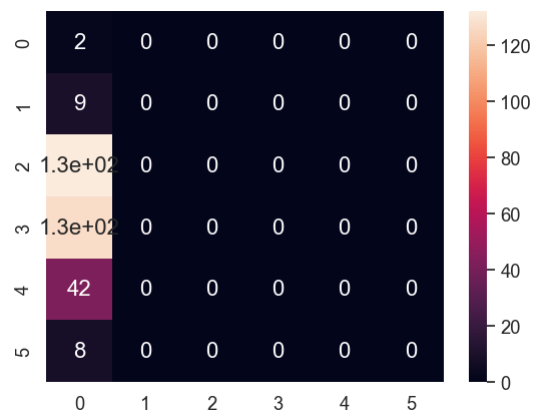
Imputed Data	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.05	0.56	0.09	9
5	0.45	0.37	0.41	132
6	0.44	0.39	0.41	127
7	1.00	0.02	0.05	42
8	0.00	0.00	0.00	8
accuracy			0.33	320
macro avg	0.32	0.22	0.16	320
weighted avg	0.49	0.33	0.34	320

## 2. Support Vector Machine

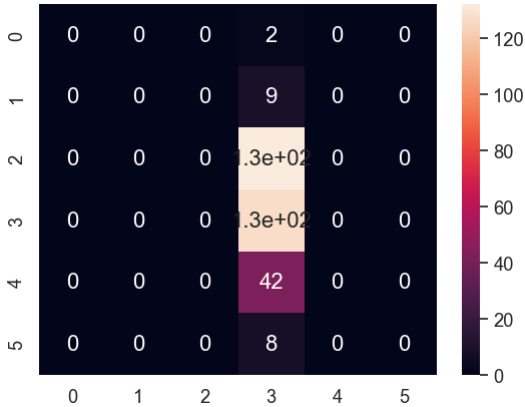
```
In [ ]: wineSVM = SVC()
clas_report(X_train, y_train, X_test, y_test, wineSVM, 'Regular Data')
clas_report(X_over, y_over, X_test, y_test, wineSVM, 'Oversampled Data')
clas_report(X_under, y_under, X_test, y_test, wineSVM, 'Undersampled Data')
clas_report(X_smote, y_smote, X_test, y_test, wineSVM, 'SMOTE Data')
clas_report(X_impute, y_impute, X_test, y_test, wineSVM, 'Imputed Data')
```



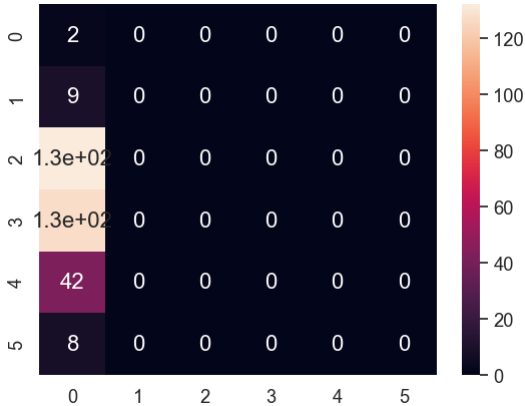
Regular Data	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.66	0.72	0.69	132
6	0.54	0.67	0.60	127
7	0.60	0.29	0.39	42
8	0.00	0.00	0.00	8
accuracy			0.60	320
macro avg	0.30	0.28	0.28	320
weighted avg	0.57	0.60	0.57	320



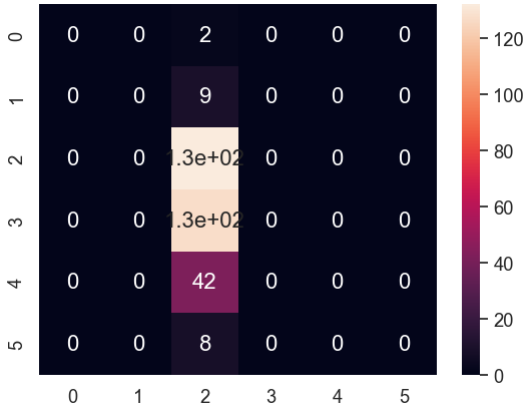
Oversampled Data				
	precision	recall	f1-score	support
3	0.01	1.00	0.01	2
4	0.00	0.00	0.00	9
5	0.00	0.00	0.00	132
6	0.00	0.00	0.00	127
7	0.00	0.00	0.00	42
8	0.00	0.00	0.00	8
accuracy			0.01	320
macro avg	0.00	0.17	0.00	320
weighted avg	0.00	0.01	0.00	320



Undersampled Data				
	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.00	0.00	0.00	132
6	0.40	1.00	0.57	127
7	0.00	0.00	0.00	42
8	0.00	0.00	0.00	8
accuracy			0.40	320
macro avg	0.07	0.17	0.09	320
weighted avg	0.16	0.40	0.23	320



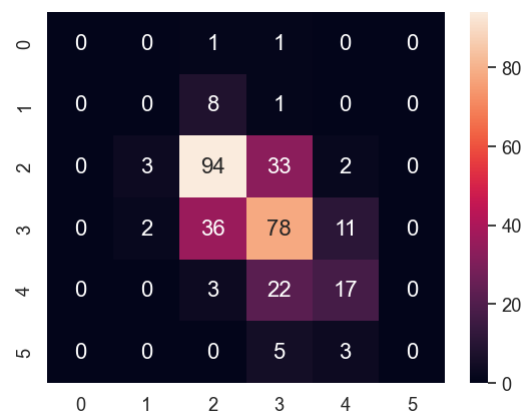
SMOTE Data				
	precision	recall	f1-score	support
3	0.01	1.00	0.01	2
4	0.00	0.00	0.00	9
5	0.00	0.00	0.00	132
6	0.00	0.00	0.00	127
7	0.00	0.00	0.00	42
8	0.00	0.00	0.00	8
accuracy			0.01	320
macro avg	0.00	0.17	0.00	320
weighted avg	0.00	0.01	0.00	320



Imputed Data		precision	recall	f1-score	support
3		0.00	0.00	0.00	2
4		0.00	0.00	0.00	9
5		0.41	1.00	0.58	132
6		0.00	0.00	0.00	127
7		0.00	0.00	0.00	42
8		0.00	0.00	0.00	8
accuracy				0.41	320
macro avg		0.07	0.17	0.10	320
weighted avg		0.17	0.41	0.24	320

### 3. Artificial Neural Network

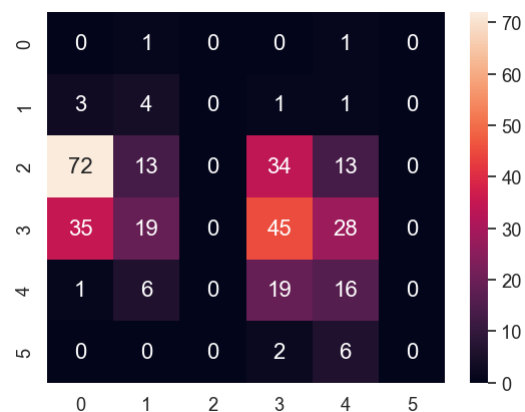
```
In [ ]: wineMLP = MLPClassifier()
clas_report(X_train, y_train, X_test, y_test, wineMLP, 'Regular Data')
clas_report(X_over, y_over, X_test, y_test, wineMLP, 'Oversampled Data')
clas_report(X_under, y_under, X_test, y_test, wineMLP, 'Undersampled Data')
clas_report(X_smote, y_smote, X_test, y_test, wineMLP, 'SMOTE Data')
clas_report(X_impute, y_impute, X_test, y_test, wineMLP, 'Imputed Data')
```



Regular Data		precision	recall	f1-score	support
3		0.00	0.00	0.00	2
4		0.00	0.00	0.00	9
5		0.66	0.71	0.69	132
6		0.56	0.61	0.58	127
7		0.52	0.40	0.45	42
8		0.00	0.00	0.00	8
accuracy				0.59	320
macro avg		0.29	0.29	0.29	320
weighted avg		0.56	0.59	0.57	320



Oversampled Data		precision	recall	f1-score	support
3		0.01	0.50	0.01	2
4		0.00	0.00	0.00	9
5		0.51	0.21	0.30	132
6		0.46	0.09	0.16	127
7		0.27	0.69	0.39	42
8		0.00	0.00	0.00	8
accuracy				0.22	320
macro avg		0.21	0.25	0.14	320
weighted avg		0.43	0.22	0.24	320



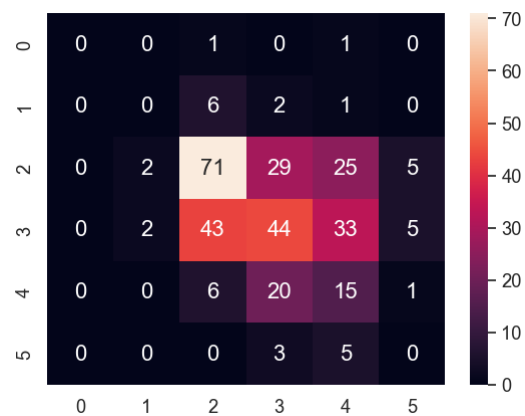
Undersampled Data

	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.09	0.44	0.15	9
5	0.00	0.00	0.00	132
6	0.45	0.35	0.39	127
7	0.25	0.38	0.30	42
8	0.00	0.00	0.00	8
accuracy			0.20	320
macro avg	0.13	0.20	0.14	320
weighted avg	0.21	0.20	0.20	320



SMOTE Data

	precision	recall	f1-score	support
3	0.02	1.00	0.03	2
4	0.03	0.22	0.05	9
5	0.45	0.15	0.23	132
6	0.56	0.17	0.27	127
7	0.32	0.29	0.30	42
8	0.05	0.12	0.07	8
accuracy			0.18	320
macro avg	0.24	0.33	0.16	320
weighted avg	0.46	0.18	0.24	320



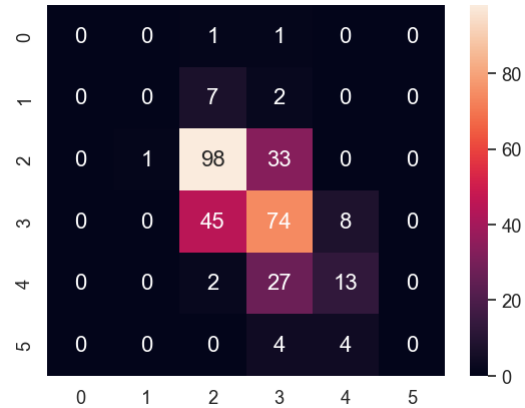
Imputed Data

	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.56	0.54	0.55	132
6	0.45	0.35	0.39	127
7	0.19	0.36	0.25	42
8	0.00	0.00	0.00	8
accuracy			0.41	320
macro avg	0.20	0.21	0.20	320
weighted avg	0.43	0.41	0.41	320

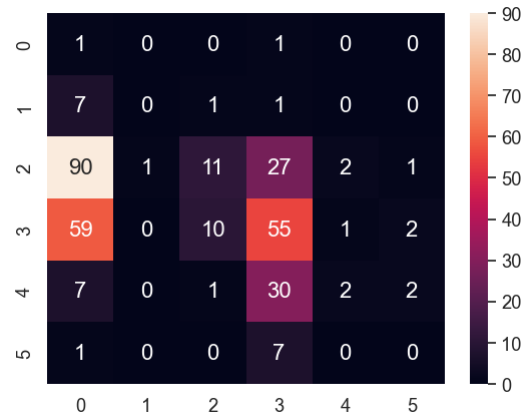


## 4. Logistic Regression

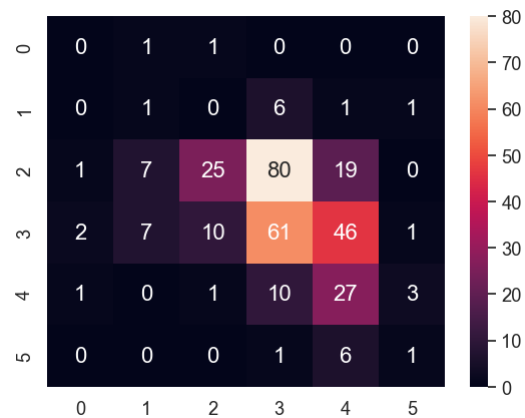
```
In [ ]: wineLR = LogisticRegression()
clas_report(X_train, y_train, X_test, y_test, wineLR, 'Regular Data')
clas_report(X_over, y_over, X_test, y_test, wineLR, 'Oversampled Data')
clas_report(X_under, y_under, X_test, y_test, wineLR, 'Undersampled Data')
clas_report(X_smote, y_smote, X_test, y_test, wineLR, 'SMOTE Data')
clas_report(X_impute, y_impute, X_test, y_test, wineLR, 'Imputed Data')
```



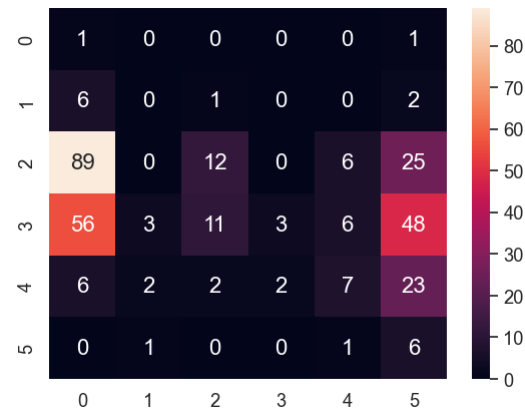
Regular Data					
	precision	recall	f1-score	support	
3	0.00	0.00	0.00	2	
4	0.00	0.00	0.00	9	
5	0.64	0.74	0.69	132	
6	0.52	0.58	0.55	127	
7	0.52	0.31	0.39	42	
8	0.00	0.00	0.00	8	
accuracy			0.58	320	
macro avg	0.28	0.27	0.27	320	
weighted avg	0.54	0.58	0.55	320	



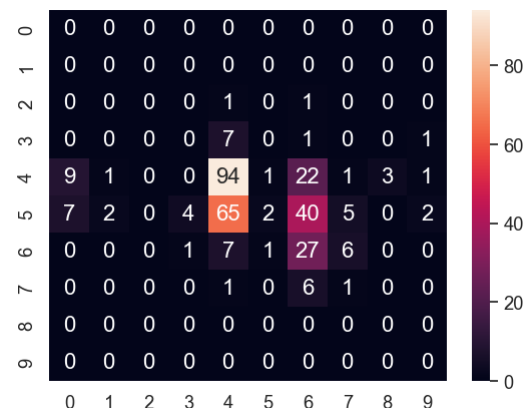
Oversampled Data					
	precision	recall	f1-score	support	
3	0.01	0.50	0.01	2	
4	0.00	0.00	0.00	9	
5	0.48	0.08	0.14	132	
6	0.45	0.43	0.44	127	
7	0.40	0.05	0.09	42	
8	0.00	0.00	0.00	8	
accuracy			0.22	320	
macro avg	0.22	0.18	0.11	320	
weighted avg	0.43	0.22	0.25	320	



Undersampled Data					
	precision	recall	f1-score	support	
3	0.00	0.00	0.00	2	
4	0.06	0.11	0.08	9	
5	0.68	0.19	0.30	132	
6	0.39	0.48	0.43	127	
7	0.27	0.64	0.38	42	
8	0.17	0.12	0.14	8	
accuracy			0.36	320	
macro avg	0.26	0.26	0.22	320	
weighted avg	0.47	0.36	0.35	320	



SMOTE Data					
	precision	recall	f1-score	support	
3	0.01	0.50	0.01	2	
4	0.00	0.00	0.00	9	
5	0.46	0.09	0.15	132	
6	0.60	0.02	0.05	127	
7	0.35	0.17	0.23	42	
8	0.06	0.75	0.11	8	
accuracy			0.09	320	
macro avg	0.25	0.26	0.09	320	
weighted avg	0.48	0.09	0.11	320	



Imputed Data					
	precision	recall	f1-score	support	
1	0.00	0.00	0.00	0	
2	0.00	0.00	0.00	0	
3	0.00	0.00	0.00	2	
4	0.00	0.00	0.00	9	
5	0.54	0.71	0.61	132	
6	0.50	0.02	0.03	127	
7	0.28	0.64	0.39	42	
8	0.08	0.12	0.10	8	
9	0.00	0.00	0.00	0	
10	0.00	0.00	0.00	0	
accuracy			0.39	320	
macro avg	0.14	0.15	0.11	320	
weighted avg	0.46	0.39	0.32	320	

## Baseline Results for comparison after hyperparameter tuning

```
In [ ]: report(X_train, y_train, X_test, y_test, wineRF, 'Regular Data')
report(X_over, y_over, X_test, y_test, wineRF, 'Oversampled Data')
report(X_under, y_under, X_test, y_test, wineRF, 'Undersampled Data')
report(X_smote, y_smote, X_test, y_test, wineRF, 'SMOTE Data')
report(X_impute, y_impute, X_test, y_test, wineRF, 'Imputed Data')

report(X_train, y_train, X_test, y_test, wineSVM, 'Regular Data')
report(X_over, y_over, X_test, y_test, wineSVM, 'Oversampled Data')
report(X_under, y_under, X_test, y_test, wineSVM, 'Undersampled Data')
report(X_smote, y_smote, X_test, y_test, wineSVM, 'SMOTE Data')
report(X_impute, y_impute, X_test, y_test, wineSVM, 'Imputed Data')

report(X_train, y_train, X_test, y_test, wineMLP, 'Regular Data')
report(X_over, y_over, X_test, y_test, wineMLP, 'Oversampled Data')
report(X_under, y_under, X_test, y_test, wineMLP, 'Undersampled Data')
report(X_smote, y_smote, X_test, y_test, wineMLP, 'SMOTE Data')
report(X_impute, y_impute, X_test, y_test, wineMLP, 'Imputed Data')
```

```
report(X_train, y_train, X_test, y_test, wineLR, 'Regular Data')
report(X_over, y_over, X_test, y_test, wineLR, 'Oversampled Data')
report(X_under, y_under, X_test, y_test, wineLR, 'Undersampled Data')
report(X_smote, y_smote, X_test, y_test, wineLR, 'SMOTE Data')
report(X_impute, y_impute, X_test, y_test, wineLR, 'Imputed Data')
```

```
Regular Data : 0.653125
Oversampled Data : 0.375
Undersampled Data : 0.284375
SMOTE Data : 0.359375
Imputed Data : 0.29375
Regular Data : 0.6
Oversampled Data : 0.00625
Undersampled Data : 0.396875
SMOTE Data : 0.00625
Imputed Data : 0.4125
Regular Data : 0.5875
Oversampled Data : 0.225
Undersampled Data : 0.2125
SMOTE Data : 0.253125
Imputed Data : 0.165625
Regular Data : 0.578125
Oversampled Data : 0.215625
Undersampled Data : 0.359375
SMOTE Data : 0.090625
Imputed Data : 0.3875
```

## 4. Hyperparameter Tuning

### 1. Random Forest Classifier

#### Manual GridSearch

##### Regular Data GridSearch

Note: These will likely take a very long time to run if you don't remove some of the hyper-parameters we tested here

##### Original Data Gridsearch

```
In [ ]: import numpy as np
#manual Grid Search (With Cut Features)
m_n_estimators = [20, 50, 75, 100, 200, 450, 700]
m_max_depth = [5,6,7,8, 15]
m_max_features = [.05, .1, .15, .2, .5, .8, 1]
m_max_samples = [.1, .2, .4, .6, .75, 1]
number_of_tests = len(m_n_estimators)*len(m_max_depth)*len(m_max_features)*len(m_max_samples)
hyper_parameter_matrix = np.zeros((number_of_tests, 5))
counter = 0

for f in m_max_features:
    for d in m_max_depth:
        for e in m_n_estimators:
            for s in m_max_samples:
                if f == 15:
                    wineRFC_HP = RandomForestClassifier(max_features=None, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                else:
                    wineRFC_HP = RandomForestClassifier(max_features=f, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                wineRFC_HP.fit(X_train_HP, y_train_HP)
                score = wineRFC_HP.score(X_test_HP, y_test_HP)
                score = np.round(score, 4)
                hyper_parameter_matrix[counter] = [f, d, e, s, score]
                counter += 1
```

```
In [ ]: #manual Grid Search (Original Data)
m_n_estimators = [20, 50, 75, 100, 200, 450, 700]
m_max_depth = [4,6,8,15]
m_max_features = [.05, .1, .15, .2, .5, .8, 1]
m_max_samples = [.1, .2, .4, .6, .75, 1]
number_of_tests = len(m_n_estimators)*len(m_max_depth)*len(m_max_features)*len(m_max_samples)
hyper_parameter_matrix_Original = np.zeros((number_of_tests, 5))
X_HP_train_Original, X_HP_test_Original, y_HP_train_Original, y_HP_test_Original = train_test_split(X_train_Original, y_train_Original, test_size = .2, random_state=10) #split
counter = 0

for f in m_max_features:
    for d in m_max_depth:
        for e in m_n_estimators:
            for s in m_max_samples:
                if f == 15:
                    wineRFC_HP = RandomForestClassifier(max_features=None, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                else:
                    wineRFC_HP = RandomForestClassifier(max_features=f, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                wineRFC_HP.fit(X_HP_train_Original, y_HP_train_Original)
                score = wineRFC_HP.score(X_HP_test_Original, y_HP_test_Original)
                score = np.round(score, 4)
                hyper_parameter_matrix_Original[counter] = [f, d, e, s, score]
                counter += 1
```

##### Smote GridSearch

```
In [ ]: #manual Grid Search (Smote Data)
m_n_estimators = [20, 50, 75, 100, 200, 450, 700]
m_max_depth = [5,6,7,8, 15]
m_max_features = [.05, .1, .15, .2, .5, .8, 1]
m_max_samples = [.1, .2, .4, .6, .75, 1]
number_of_tests = len(m_n_estimators)*len(m_max_depth)*len(m_max_features)*len(m_max_samples)
hyper_parameter_matrix_Smote = np.zeros((number_of_tests, 5))
counter = 0

for f in m_max_features:
    for d in m_max_depth:
        for e in m_n_estimators:
            for s in m_max_samples:
                if f == 15:
                    wineRFC_HP = RandomForestClassifier(max_features=None, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                else:
                    wineRFC_HP = RandomForestClassifier(max_features=f, max_depth=d, n_estimators=e, max_samples=s, oob_score=True, random_state=417)
                wineRFC_HP.fit(X_smote, y_smote)
                score = wineRFC_HP.score(X_test_HP, y_test_HP)
                score = np.round(score, 4)
                hyper_parameter_matrix_Smote[counter] = [f, d, e, s, score]
                counter += 1
```

Find best Hyper Parameters for each model

## Regular Data

```
In [ ]: np.set_printoptions(suppress=True)
np.set_printoptions(threshold=10000)
best_HP = hyper_parameter_matrix[0]

for i in range(len(hyper_parameter_matrix)):
    if hyper_parameter_matrix[i][4] > best_HP[4]:
        # print(best_HP)
        best_HP = hyper_parameter_matrix[i]

print(best_HP)

[ 0.05  15.   200.    0.75   0.6562]
```

## Original Data

```
In [ ]: # print(hyper_parameter_matrix_Original)
best_HP_Original = hyper_parameter_matrix_Original[0]

for i in range(len(hyper_parameter_matrix_Original)):
    if hyper_parameter_matrix_Original[i][4] > best_HP_Original[4]:
        # print(best_HP_Original)
        best_HP_Original = hyper_parameter_matrix_Original[i]

print(best_HP_Original)

[ 0.05  15.    75.    0.75   0.7266]
```

## Smote Data

```
In [ ]: best_HP_Smote = hyper_parameter_matrix_Smote[0]

for i in range(len(hyper_parameter_matrix_Smote)):
    if hyper_parameter_matrix_Smote[i][4] > best_HP_Smote[4]:
        # print(best_HP_Smote)
        best_HP_Smote = hyper_parameter_matrix_Smote[i]

print(best_HP_Smote)

[ 0.5    8.    50.    0.75   0.4922]
```

## Visualize hyper-parameter around local maxima for best performer (Original)

```
In [ ]: #creating hyper-parameter-matrices (Just for Original)
n_estimators_oop_matrix_2 = np.zeros((1,2))
max_depth_oop_matrix_2 = np.zeros((1,2))
max_features_oop_matrix_2 = np.zeros((1,2))
max_samples_oop_matrix_2 = np.zeros((1,2))

for i in hyper_parameter_matrix_Original:
    for j in range(0,4):
        HP_to_check = {0, 1, 2, 3}
        HP_to_check.remove(j)
        all_good=True
        for l in HP_to_check:
            if i[l] != best_HP_Original[l]:
                all_good=False
        if all_good:
            if j==0:
                max_features_oop_matrix_2 = np.append(max_features_oop_matrix_2,[[i[0], i[4]]], axis=0)
            elif j==1:
                max_depth_oop_matrix_2 = np.append(max_depth_oop_matrix_2 ,[[i[1], i[4]]], axis=0)
            elif j==2:
                n_estimators_oop_matrix_2 = np.append(n_estimators_oop_matrix_2 ,[[i[2], i[4]]], axis=0)
            elif j==3:
                max_samples_oop_matrix_2 = np.append(max_samples_oop_matrix_2 ,[[i[3], i[4]]], axis=0)

max_features_oop_matrix_Original = np.delete(max_features_oop_matrix_2, obj=0, axis=0)
max_depth_oop_matrix_Original = np.delete(max_depth_oop_matrix_2, obj=0, axis=0)
n_estimators_oop_matrix_Original = np.delete(n_estimators_oop_matrix_2, obj=0, axis=0)
max_samples_oop_matrix_Original = np.delete(max_samples_oop_matrix_2, obj=0, axis=0)
```

## Making the plots

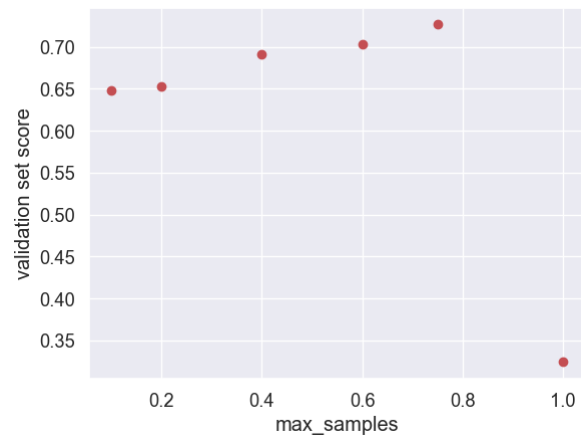
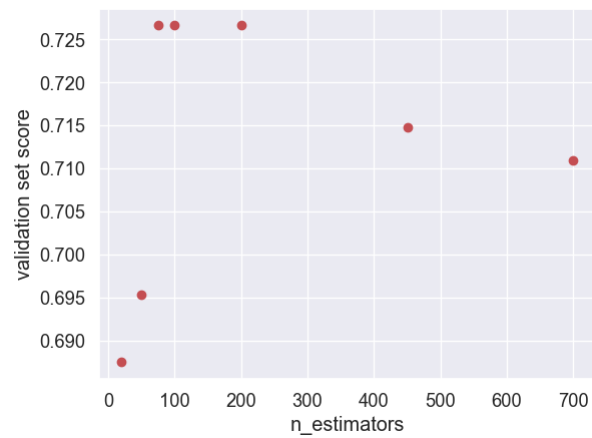
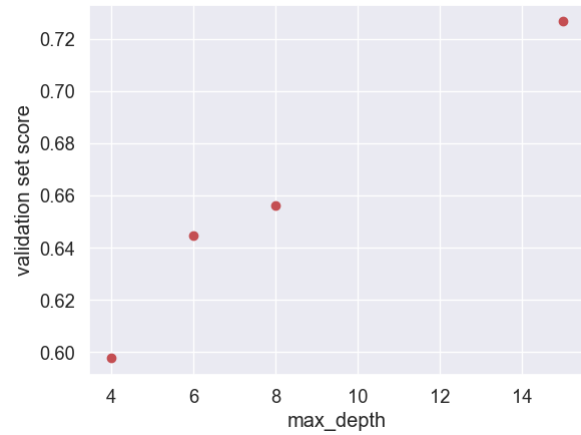
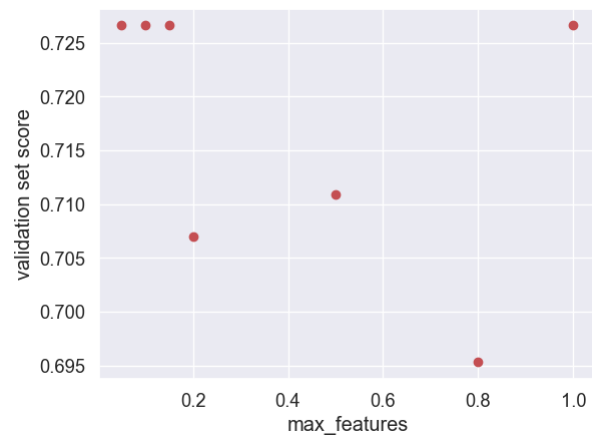
```
In [ ]: plt.plot(max_features_oop_matrix_Original[:, 0],max_features_oop_matrix_Original[:, 1], 'ro')
plt.xlabel('max_features')
plt.ylabel('validation set score')
plt.show()

plt.plot(max_depth_oop_matrix_Original[:, 0], max_depth_oop_matrix_Original[:, 1], 'ro')
plt.xlabel('max_depth')
plt.ylabel('validation set score')
plt.show()

plt.plot(n_estimators_oop_matrix_Original[:, 0], n_estimators_oop_matrix_Original[:, 1], 'ro')
plt.xlabel('n_estimators')
plt.ylabel('validation set score')
plt.show()

plt.plot(max_samples_oop_matrix_Original[:, 0], max_samples_oop_matrix_Original[:, 1], 'ro')
plt.xlabel('max_samples')
plt.ylabel('validation set score')
plt.show()

print(best_HP_Original)
```



[ 0.05 15. 75. 0.75 0.7266]

```
In [ ]: def grid_search(X_train, y_train, model, param_grid, title):
        grid = GridSearchCV(estimator=model, param_grid=param_grid, cv=5)
        grid.fit(X_train, y_train)
        print(title, ":", grid.best_params_)
        print("best score:", grid.best_score_)
```

```
In [ ]: # wineRFC = RandomForestClassifier(random_state=417)
#
# param_gridRFC = {
#     'n_estimators': [100, 200, 500, 1000],
#     'max_depth': [5,6,7,8],
# }
#
# grid_search(X_train, y_train, wineRFC, param_gridRFC, 'Regular Data')
# grid_search(X_over, y_over, wineRFC, param_gridRFC, 'Oversampled Data')
# grid_search(X_under, y_under, wineRFC, param_gridRFC, 'Undersampled Data')
# grid_search(X_smote, y_smote, wineRFC, param_gridRFC, 'SMOTE Data')
# grid_search(X_impute, y_impute, wineRFC, param_gridRFC, 'Imputed Data')
```

## 2. Support Vector Machine

```
In [ ]: wineSVM = SVC()
param_grid_svm = {
    'kernel': ['rbf', 'sigmoid', 'poly'],
    'C': [1, 10, 100, 1000]
}

grid_search(X_train, y_train, wineSVM, param_grid_svm, 'Regular Data')
grid_search(X_over, y_over, wineSVM, param_grid_svm, 'Oversampled Data')
grid_search(X_under, y_under, wineSVM, param_grid_svm, 'Undersampled Data')
grid_search(X_smote, y_smote, wineSVM, param_grid_svm, 'SMOTE Data')
grid_search(X_impute, y_impute, wineSVM, param_grid_svm, 'Imputed Data')

Regular Data : {'C': 1, 'kernel': 'rbf'}
best score: 0.6067493872549019
Oversampled Data : {'C': 1000, 'kernel': 'rbf'}
best score: 0.6841409691629956
Undersampled Data : {'C': 1, 'kernel': 'rbf'}
best score: 0.6875094449287997
SMOTE Data : {'C': 1000, 'kernel': 'rbf'}
best score: 0.6771942792672817
Imputed Data : {'C': 1000, 'kernel': 'rbf'}
best score: 0.5710942367601245
```

## 3. Artificial Neural Network

```
In [ ]: wineANN = MLPClassifier(activation='logistic')
param_grid_ann = {
    'learning_rate_init': [0.001, 0.01, 0.1],
    'max_iter': [100, 300, 500],
    'hidden_layer_sizes': [(10,), (100,), (300,)]
}

grid_search(X_train, y_train, wineANN, param_grid_ann, 'Regular Data')
grid_search(X_over, y_over, wineANN, param_grid_ann, 'Oversampled Data')
grid_search(X_under, y_under, wineANN, param_grid_ann, 'Undersampled Data')
grid_search(X_smote, y_smote, wineANN, param_grid_ann, 'SMOTE Data')
grid_search(X_impute, y_impute, wineANN, param_grid_ann, 'Imputed Data')

Regular Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.001, 'max_iter': 100}
best score: 0.6083180147058823
Oversampled Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.001, 'max_iter': 500}
best score: 0.6823788546255507
Undersampled Data : {'hidden_layer_sizes': (100,), 'learning_rate_init': 0.001, 'max_iter': 300}
best score: 0.7058238884045337
SMOTE Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.01, 'max_iter': 500}
best score: 0.715860399278175
Imputed Data : {'hidden_layer_sizes': (300,), 'learning_rate_init': 0.001, 'max_iter': 500}
best score: 0.5685669781931464
```

## 4. Logistic Regression

```
In [ ]: wineLR = LogisticRegression()
param_grid_lr = {
    'penalty': ['l1', 'l2'],
    'solver': ['lbfgs', 'liblinear', 'saga'],
    'max_iter': [100, 300, 500]
}

grid_search(X_train, y_train, wineLR, param_grid_lr, 'Regular Data')
grid_search(X_over, y_over, wineLR, param_grid_lr, 'Oversampled Data')
grid_search(X_under, y_under, wineLR, param_grid_lr, 'Undersampled Data')
grid_search(X_smote, y_smote, wineLR, param_grid_lr, 'SMOTE Data')
grid_search(X_impute, y_impute, wineLR, param_grid_lr, 'Imputed Data')

Regular Data : {'max_iter': 100, 'penalty': 'l2', 'solver': 'lbfgs'}
best score: 0.6004993872549019
Oversampled Data : {'max_iter': 300, 'penalty': 'l1', 'solver': 'liblinear'}
best score: 0.6533039647577092
Undersampled Data : {'max_iter': 500, 'penalty': 'l2', 'solver': 'lbfgs'}
best score: 0.7112060447544318
SMOTE Data : {'max_iter': 100, 'penalty': 'l1', 'solver': 'liblinear'}
best score: 0.5051326787429711
Imputed Data : {'max_iter': 500, 'penalty': 'l2', 'solver': 'lbfgs'}
best score: 0.5748033489096572
```

## 5. Testing Final Models

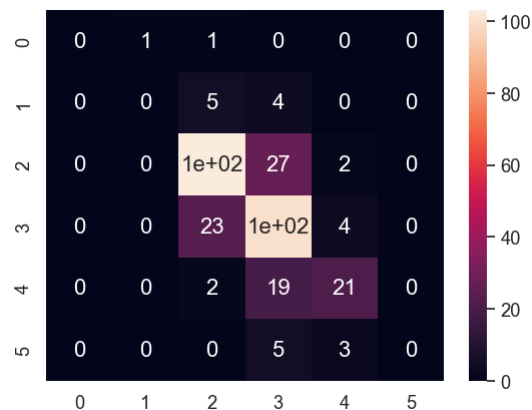
```
In [ ]: def final_report(X_train, y_train, X_test, y_test, model, title):
    model.fit(X_train, y_train)
    y_pred = model.predict(X_test)
    print(title, "\n", model.score(X_test, y_test))
    # clas = classification_report(y_pred, y_test)
    # print(clas)
```

### 1. Random Forest Classifier

```
In [ ]: #True Final RFC model (the rest are for data analysis)
finalRFC = RandomForestClassifier(max_features=best_HP_Original[0], max_depth=int(best_HP_Original[1]), n_estimators=int(best_HP_Original[2]), max_samples=best_HP_Original[3])
finalRFC.fit(X_train_Original, y_train_Original)
print(finalRFC.score(X_test_Original, y_test_Original))
clas_report(X_train_Original, y_train_Original, X_test_Original, y_test_Original, finalRFC, "Final RFC W/ Original Data")

0.7
```

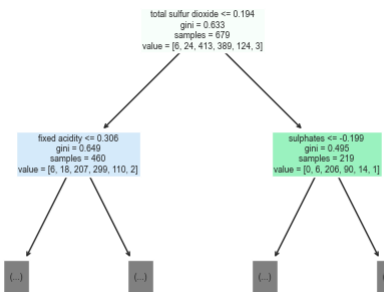




Final RFC W/	Original Data			
	precision	recall	f1-score	support
3	0.00	0.00	0.00	2
4	0.00	0.00	0.00	9
5	0.77	0.78	0.77	132
6	0.65	0.79	0.71	127
7	0.70	0.50	0.58	42
8	0.00	0.00	0.00	8
accuracy			0.70	320
macro avg	0.35	0.34	0.34	320
weighted avg	0.66	0.70	0.68	320

```
In [ ]: from sklearn import tree
original_tree_features = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density', 'pH', '']
estimator = final RFC.estimators_[2]
tree.plot_tree(final RFC.estimators_[2], max_depth=1, feature_names=original_tree_features, filled = True)

Out [ ]: [Text(0.5, 0.8333333333333334, 'total sulfur dioxide <= 0.194\nngini = 0.633\nnsamples = 679\nvalue = [6, 24, 413, 389, 124, 3]'),
Text(0.25, 0.5, 'fixed acidity <= 0.306\nngini = 0.649\nnsamples = 460\nvalue = [6, 18, 207, 299, 110, 2]'),
Text(0.125, 0.16666666666666666, '\n (...) \n'),
Text(0.375, 0.16666666666666666, '\n (...) \n'),
Text(0.75, 0.5, 'sulphates <= -0.199\nngini = 0.495\nnsamples = 219\nvalue = [0, 6, 206, 90, 14, 1]'),
Text(0.625, 0.16666666666666666, '\n (...) \n'),
Text(0.875, 0.16666666666666666, '\n (...) \n')]
```



```
In [ ]: # wineRF_regular_final = RandomForestClassifier(n_estimators=100, max_depth=8, random_state=417)
# wineRF_over_final = RandomForestClassifier(n_estimators=200, max_depth=8, random_state=417)
# wineRF_under_final = RandomForestClassifier(n_estimators=1000, max_depth=7, random_state=417)
# wineRF_smote_final = RandomForestClassifier(n_estimators=200, max_depth=8, random_state=417)
# wineRF_impute_final = RandomForestClassifier(n_estimators=100, max_depth=8, random_state=417)
#
# final_report(X_train, y_train, X_test, y_test, wineRF_regular_final, 'Regular Data')
# final_report(X_over, y_over, X_test, y_test, wineRF_over_final, 'Oversampled Data')
# final_report(X_under, y_under, X_test, y_test, wineRF_under_final, 'Undersampled Data')
# final_report(X_smote, y_smote, X_test, y_test, wineRF_smote_final, 'SMOTE Data')
# final_report(X_impute, y_impute, X_test, y_test, wineRF_impute_final, 'Imputed Data')
```

## 2. Support Vector Machine

```
In [ ]: wineSVM_regular_final = SVC(kernel = 'rbf', C = 1)
wineSVM_over_final = SVC(kernel = 'rbf', C = 10)
wineSVM_under_final = SVC(kernel = 'rbf', C = 1)
wineSVM_smote_final = SVC(kernel = 'rbf', C = 1000)
wineSVM_impute_final = SVC(kernel = 'rbf', C = 1)

final_report(X_train, y_train, X_test, y_test, wineSVM_regular_final, 'Regular Data')
final_report(X_over, y_over, X_test, y_test, wineSVM_over_final, 'Oversampled Data')
final_report(X_under, y_under, X_test, y_test, wineSVM_under_final, 'Undersampled Data')
final_report(X_smote, y_smote, X_test, y_test, wineSVM_smote_final, 'SMOTE Data')
final_report(X_impute, y_impute, X_test, y_test, wineSVM_impute_final, 'Imputed Data')
```

Regular Data : 0.6  
Oversampled Data : 0.00625  
Undersampled Data : 0.396875  
SMOTE Data : 0.00625  
Imputed Data : 0.4125

## 3. Artificial Neural Network

```
In [ ]: wineANN_regular_final = MLPClassifier(learning_rate_init=0.01, max_iter=100, hidden_layer_sizes=(100,))
wineANN_over_final = MLPClassifier(learning_rate_init=0.01, max_iter=500, hidden_layer_sizes=(300,))
wineANN_under_final = MLPClassifier(learning_rate_init=0.1, max_iter=100, hidden_layer_sizes=(100,))
wineANN_smote_final = MLPClassifier(learning_rate_init=0.01, max_iter=500, hidden_layer_sizes=(300,))
wineANN_impute_final = MLPClassifier(learning_rate_init=0.1, max_iter=300, hidden_layer_sizes=(300,))
```

```

final_report(X_train, y_train, X_test, y_test, wineANN_regular_final, 'Regular Data')
final_report(X_over, y_over, X_test, y_test, wineANN_over_final, 'Oversampled Data')
final_report(X_under, y_under, X_test, y_test, wineANN_under_final, 'Undersampled Data')
final_report(X_smote, y_smote, X_test, y_test, wineANN_smote_final, 'SMOTE Data')
final_report(X_impute, y_impute, X_test, y_test, wineANN_impute_final, 'Imputed Data')

```

```

Regular Data : 0.578125
Oversampled Data : 0.065625
Undersampled Data : 0.221875
SMOTE Data : 0.359375
Imputed Data : 0.184375

```

## 4. Logistic Regression

```

In [ ]: wineLR_regular_final = LogisticRegression(penalty='l2', solver='lbfgs', max_iter=100)
wineLR_over_final = LogisticRegression(penalty='l1', solver='saga', max_iter=100)
wineLR_under_final = LogisticRegression(penalty='l2', solver='lbfgs', max_iter=100)
wineLR_smote_final = LogisticRegression(penalty='l2', solver='lbfgs', max_iter=100)
wineLR_impute_final = LogisticRegression(penalty='l2', solver='lbfgs', max_iter=100)

```

```

final_report(X_train, y_train, X_test, y_test, wineLR_regular_final, 'Regular Data')
final_report(X_over, y_over, X_test, y_test, wineLR_over_final, 'Oversampled Data')
final_report(X_under, y_under, X_test, y_test, wineLR_under_final, 'Undersampled Data')
final_report(X_smote, y_smote, X_test, y_test, wineLR_smote_final, 'SMOTE Data')
final_report(X_impute, y_impute, X_test, y_test, wineLR_impute_final, 'Imputed Data')

```

```

Regular Data : 0.578125
Oversampled Data : 0.2
Undersampled Data : 0.359375
SMOTE Data : 0.090625
Imputed Data : 0.3875

```