Neural Network EDA, Model and Prediction on Wine Data

April 30, 2023

1 Brief

Welcome to the EDA, Model and Prediction of wine data. One of the ways to determine wine quality is by its physiocochemistry. The purpose of this study was to look at the physicochemical properties of the wine and analyze which model, of the ones we learned in the course, can give us the best fit, prediction and accuracy score of the quality of the wine.

2 Introduction

Ever wondered if a wine is good based on it's phiscochemical measurements? We will look into building a set of models to analyze exactly that!

In this project, the data consisted of 6497 observations across 11 physicochemical properties and the corresponding quality. The datasets were cleaned and explored. Then, we build a deep neural network and predict the quality of wine based on the model built.

After that, the accuracy test showed how well the model did at predicting the wine quality based on the physicochemical properties of the wine.

3 The Data

3.1 Sources

Data Set Source: > "Wine Quality Data Set." UCI Machine Learning Repository, 7 Oct. 2009, https://archive.ics.uci.edu/ml/datasets/Wine+Quality

Data Set Research: > P. Cortez, A. Cerdeira, F. Almeida, T. Matos and J. Reis. Modeling wine preferences by data mining from physicochemical properties. In Decision Support Systems, Elsevier, 47(4):547-553, 2009.

Further Research on Wine properties: > Mor, Nuriel S., et al. "Wine Quality and Type Prediction from Physicochemical Properties Using Neural Networks for Machine Learning: A Free Software for Winemakers and Customers." AgriRxiv, vol. 2022, 30 Jan. 2022, https://doi.org/10.31220/agrirxiv.2022.00125.

Further Research on Winemaking > "Winemaking." Wikipedia: The Free Encyclopedia. Wikimedia Foundation, Inc, 22 July 2004, https://en.wikipedia.org/wiki/Winemaking. Accessed 7 Dec. 2022.

3.2 Description of Data

The data, acquired from UCI, is related with wine physicochemical inputs and sensory outputs for a Portuguese "Vinho Verde" wine.

3.2.1 Physicochemical Properties

In this dataset, the wine quality is determined by 11 physicochemical qualities: fixed acidity, volatile acidity, citric acid, residual sugar, chlorides, free sulfur dioxide, total sulfur dioxide, density, pH, sulphates, and alcohol. I did some outside research to understand what these qualities are and what their values mean.

Fixed Acidity: The fixed acids found in wines are tartaric, malic, citric, and succinic. Values are how much acid is in the wine based on the type of acid.

Volatile acidity: This is a measure of the low molecular weight of fatty acids.

Critic Acid: This is an organic acid, added to give wine its sour taste amongst other things.

Residual Sugar: The sugar that is leftover after the fermentation process.

Chlorides: This measures how much salt is in the wine.

Free Sulfur Dioxide: These are another type of salty additive generally used for oxidation. A fraction of this will react with sugars and fraction will not. This is the fraction that did not react.

Total Sulfur Dioxide: The total of reacted and free sulfur dioxide.

Density: The measure of mass per unit volume.

pH: The scale used to measure acidity and basicity of wine.

Sulfites: The chemical compounds that contain sulfite ions for preservation.

Alcohol: The Alcohol by Volume of wine.

3.2.2 Other Variables

Aside from the physicochemcial properties, there are a couple other variables in this dataset.

Quality: This is a score or grade given to a wine in part based on the above physicochemical properties.

Color: Red wine or White wine.

3.3 Data Import

To start building models, the data must be first imported.

3.3.1 Import Python Libraries

Begin by importing all the libraries that will be used.

```
[26]: import pandas as pd
      import matplotlib.pyplot as plt
      %matplotlib inline
      import numpy as np
      import math
      import seaborn as sns
      sns.set()
      from sklearn.model_selection import train_test_split
      from sklearn.preprocessing import StandardScaler
      from sklearn.metrics import accuracy_score
      from sklearn import preprocessing, model_selection
      import sklearn.metrics as metrics
      from sklearn.metrics import roc_auc_score, roc_curve, confusion_matrix, u
       \hookrightarrow classification_report
      from tensorflow import keras
      from tensorflow.keras import layers, optimizers, regularizers
      from tensorflow.keras.models import Model, Sequential
      from tensorflow.keras.layers import Dense, Dropout, Input, Add,
       →BatchNormalization, \
                                           Softmax, Activation
      from tensorflow.keras.optimizers import Adam
      from keras.utils import plot_model
      #from kt utils import *
      import keras.backend as K
      # Ignore Future Warnings
      import warnings
      warnings.simplefilter(action='ignore', category=FutureWarning)
```

3.3.2 Import Wine Quality Data

Next, There are 2 CSV files to import that identify the quality of wines: one for red wines and one for white wines. These will be concatenated and this forms our complete DataFrame.

Since 'color' is a string label, it can be encoded of using the label as a number so it can be used for analysis.

Finally, the column names are updated with underscores to enable easier coding.

```
[27]: # Import red wines
      df_red = pd.read_csv('https://archive.ics.uci.edu/ml/machine-learning-databases/
      ⇔wine-quality/winequality-red.csv', sep=';')
      df red['color'] = "red"
      # # Import white wines
      df_white = pd.read_csv('https://archive.ics.uci.edu/ml/
      →machine-learning-databases/wine-quality/winequality-white.csv', sep=';')
      df_white['color'] = "white"
      # # Combine red and white
      df = pd.concat([df_red, df_white], ignore_index=True)
      df.reindex()
      # # Encode color label
      df['color'] = df['color'].astype('category')
      df['color_enc'] = df['color'].cat.codes
      # Use underscores instead of spaces in column names
      df.columns = [c.replace(' ', '_') for c in df.columns]
      df
[27]:
            fixed_acidity volatile_acidity citric_acid residual_sugar chlorides \
      0
                      7.4
                                       0.70
                                                    0.00
                                                                      1.9
                                                                               0.076
                      7.8
                                                    0.00
                                                                      2.6
      1
                                       0.88
                                                                               0.098
      2
                      7.8
                                       0.76
                                                    0.04
                                                                      2.3
                                                                               0.092
      3
                     11.2
                                                                      1.9
                                       0.28
                                                    0.56
                                                                               0.075
      4
                      7.4
                                       0.70
                                                    0.00
                                                                      1.9
                                                                               0.076
      6492
                      6.2
                                       0.21
                                                    0.29
                                                                      1.6
                                                                               0.039
      6493
                      6.6
                                       0.32
                                                    0.36
                                                                     8.0
                                                                               0.047
                      6.5
                                       0.24
                                                    0.19
                                                                      1.2
      6494
                                                                               0.041
      6495
                      5.5
                                       0.29
                                                    0.30
                                                                      1.1
                                                                               0.022
      6496
                      6.0
                                       0.21
                                                    0.38
                                                                      0.8
                                                                               0.020
            free_sulfur_dioxide total_sulfur_dioxide density
                                                                  pH sulphates \
      0
                           11.0
                                                 34.0 0.99780
                                                                3.51
                                                                            0.56
                           25.0
      1
                                                 67.0 0.99680
                                                                3.20
                                                                            0.68
      2
                           15.0
                                                 54.0 0.99700
                                                                3.26
                                                                            0.65
                           17.0
      3
                                                 60.0 0.99800
                                                                3.16
                                                                            0.58
      4
                           11.0
                                                 34.0 0.99780
                                                                3.51
                                                                            0.56
                                                 92.0 0.99114
      6492
                           24.0
                                                                3.27
                                                                            0.50
                           57.0
                                                                            0.46
      6493
                                                168.0 0.99490
                                                                3.15
                                                                2.99
      6494
                           30.0
                                                111.0 0.99254
                                                                            0.46
      6495
                           20.0
                                                110.0 0.98869 3.34
                                                                            0.38
```

6496	22.0	98.0	0.98941	3.26	0.32
0-100	22.0	50.0	0.00041	0.20	0.02

	alcohol	quality	color	color_enc
0	9.4	5	red	0
1	9.8	5	red	0
2	9.8	5	red	0
3	9.8	6	red	0
4	9.4	5	red	0
•••	•••		•••	
6492	11.2	6	white	1
6493	9.6	5	white	1
6494	9.4	6	white	1
6495	12.8	7	white	1
6496	11.8	6	white	1

[6497 rows x 14 columns]

The data has been imported into a DataFrame and is now ready for cleaning and analysis.

4 Cleaning the Data and Exploratory Data Analysis

Conduct some exploratory data analysis to understand the data a little better.

First, the shape of the dataset.

```
[28]: print(f"Shape of Dataset: {df.shape}")
```

Shape of Dataset: (6497, 14)

To understand the spread of the data in the columns, look at the column ranges (min and max).

```
[29]: # Print min and max for each column
for c in df.columns:
    # color is a Categorical column and doesn't have a min and max
    if c == "color":
        continue

print(f"{c:<22}: (min, max) = ({df[c].min():.2f}, {df[c].max():.2f})")</pre>
```

```
fixed acidity
                      : (min, max) = (3.80, 15.90)
volatile_acidity
                      : (min, max) = (0.08, 1.58)
citric_acid
                      : (min, max) = (0.00, 1.66)
residual_sugar
                      : (min, max) = (0.60, 65.80)
chlorides
                      : (min, max) = (0.01, 0.61)
free_sulfur_dioxide
                      : (min, max) = (1.00, 289.00)
total_sulfur_dioxide
                     : (min, max) = (6.00, 440.00)
density
                      : (min, max) = (0.99, 1.04)
                      : (min, max) = (2.72, 4.01)
Нq
```

```
      sulphates
      : (min, max) = (0.22, 2.00)

      alcohol
      : (min, max) = (8.00, 14.90)

      quality
      : (min, max) = (3.00, 9.00)

      color_enc
      : (min, max) = (0.00, 1.00)
```

And then lets take a look at how many unique values are in each column.

```
[30]: df.nunique()
```

```
[30]: fixed_acidity
                                106
      volatile_acidity
                                187
      citric_acid
                                 89
      residual_sugar
                                316
      chlorides
                                214
      free_sulfur_dioxide
                                135
      total_sulfur_dioxide
                                276
      density
                                998
                                108
      рΗ
      sulphates
                                111
      alcohol
                                111
      quality
                                  7
      color
                                  2
                                  2
      color_enc
      dtype: int64
```

The data is then visually observed to check for any missing data or any values that should not be there. Any unknowns are converted to NA values and depending on how many NAs there are, those cells or features are dropped or imputed. Here is a tally of how many null values are in the data.

Column Name	Real Nulls	Null-like
fixed_acidity	0	0
volatile_acidity	0	0
citric_acid	0	0
residual_sugar	0	0
chlorides	0	0
<pre>free_sulfur_dioxide</pre>	0	0
total_sulfur_dioxide	0	0
density	0	0

рН	0	0
sulphates	0	0
alcohol	0	0
quality	0	0
color	0	0
color enc	0	0

There are certain columns that do not affect the quality of wine. For example "color". Whether a wine is white or red by itself has no bearing on its quality.

As a note sometimes the color of the grape does have a different average for a particular physicochemical property than the other.

Therefore, before we start analyzing the data, the color column. the encoded color column can be used instead, allowing for some forms of analysis.

```
[32]: df = df.drop('color', axis=1)
df = df.drop('color_enc', axis=1)
df
```

	fixed_acidity	volat	ile_acidity	citric_ac	id res	idual_su	ıgar	chlori	des	\
)	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	
1	7.4		0.70	0.	00		1.9	0.	076	
	•••		•••			•••				
3492	6.2		0.21	0.	29		1.6	0.	039	
3493	6.6		0.32	0.	36		8.0	0.	047	
3494	6.5		0.24	0.	19		1.2	0.	041	
3495	5.5		0.29	0.	30		1.1	0.	022	
3496	6.0		0.21	0.	38		0.8	0.	020	
	free_sulfur_di	oxide	total_sulfu	r_dioxide	densit	у рН	sulj	phates	\	
)		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	0 3.26		0.65		
3		17.0		60.0	0.9980	0 3.16		0.58		
1		11.0		34.0	0.9978	3.51		0.56		
••										
3492		24.0		92.0	0.9911	4 3.27		0.50		
3493		57.0		168.0	0.9949	0 3.15		0.46		
3494		30.0		111.0	0.9925	4 2.99		0.46		
3495		20.0		110.0	0.9886	3.34		0.38		
3496		22.0		98.0	0.9894	1 3.26		0.32		
	6492 6493 6494 6495 6496 6492 6493 6494 6495	7.4 7.8 7.8 7.8 7.8 11.2 7.4 	7.4 7.8 7.8 7.8 7.8 7.8 7.4 7.4 7.4 7.4 7.4 7.4 7.4 7.4 7.4 7.4	7.4 0.70 7.8 0.88 7.8 0.76 8 11.2 0.28 7.4 0.70 8 7.4 0	7.4 0.70 0.6 7.8 0.88 0.88 7.8 0.76 0.8 7.4 0.70 0.70 7.4 0.76 0.8 7.4 0.70 0.70 7.4 0.70 0.70 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.4 0.70 0.8 7.5 0.29 0.8 7.6 0.21 0.8 7.6 0.21 0.8 7.7 0.8 7.8 0.8 0.8 7.0 0.8 0.8 7.	7.4 0.70 0.00 7.8 0.88 0.00 7.8 0.76 0.04 8 11.2 0.28 0.56 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 7.4 0.70 0.00 8 11.0 0.29 0.30 8 11.0 0.24 0.19 8 11.0 0.29 0.30 8 11.0 0.21 0.38 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9978 8 11.0 0.9988	7.4 0.70 0.00 7.8 0.88 0.00 7.8 0.76 0.04 7.8 0.70 0.00 7.8 0.76 0.04 7.8 0.70 0.00 7.8 0.76 0.04 7.4 0.70 0.00 7.4 0.70 0.00 7.4 0.70 0.00 7.4 0.70 0.00 7.4 0.70 0.00 7.4 0.70 0.00 7.5 0.28 0.56 7.4 0.70 0.00 7.6 0.00 7.7 0.00 7.8 0.29 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.30 7.8 0.20 0.3	7.4 0.70 0.00 1.9 7.8 0.88 0.00 2.6 7.8 0.78 0.76 0.04 2.3 8 11.2 0.28 0.56 1.9 7.4 0.70 0.00 1.9 7.4 0.70 0.00 1.9 7.4 0.70 0.00 1.9 7.4 0.70 0.00 1.9 7.5 0.21 0.29 1.6 7.6 0.24 0.19 1.2 7.6 0.29 0.30 1.1 7.6 0.0 0.21 0.38 0.8 7 11.0 34.0 0.99780 3.51 7 25.0 67.0 0.99680 3.20 7 15.0 54.0 0.99780 3.51 7 11.0 34.0 0.99780 3.51 7 11.0 34.0 0.99780 3.51 7 11.0 34.0 0.99780 3.51 7 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51 8 11.0 34.0 0.99780 3.51	7.4 0.70 0.00 1.9 0. 7.8 0.88 0.00 2.6 0. 7.8 0.78 0.76 0.04 2.3 0. 7.8 0.70 0.00 1.9 0. 7.8 0.78 0.76 0.04 2.3 0. 7.8 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.4 0.70 0.00 1.9 0. 7.5 0.29 0.36 8.0 0. 7.6 0.24 0.19 1.2 0. 7.6 0.29 0.30 1.1 0.20 1.1 0. 7.6 0.29 0.30 1.1 0. 7.7 0.29 0.30 1.1 0. 7.8 0.29 0.30 0. 7.8 0.29 0.30 0. 7.8 0.29 0.30 0. 7.8 0.29 0.30 0. 7.9 0.29 0.30 0. 7.9 0.20 0.99780 3.51 0.56 0.65 0.65 0.65 0.65 0.65 0.65 0.65	7.4 0.70 0.00 1.9 0.076 7.8 0.88 0.00 2.6 0.098 7.8 0.76 0.04 2.3 0.092 7.4 0.70 0.00 1.9 0.075 7.4 0.70 0.00 1.9 0.075 7.4 0.70 0.00 1.9 0.075 7.4 0.70 0.00 1.9 0.076 7.4 0.70 0.00 1.9 0.076 7.4 0.70 0.00 1.9 0.076 7.4 0.70 0.00 1.9 0.076 7.4 0.70 0.00 1.9 0.076 7.4 0.70 0.00 1.9 0.00 7.4 0.039 7.4 0.70 0.00 1.9 0.00 7.4 0.039 7.4 0.19 1.2 0.041 7.4 0.70 0.29 0.30 1.1 0.022 7.4 0.19 1.2 0.041 7.4 0.7 0.22 0.36 8.0 0.022 7.5 0.29 0.30 1.1 0.022 7.6 0.0 0.21 0.38 0.8 0.020 7.8 0.00 0.21 0.38 0.8 0.020 7.8 0.00 0.21 0.38 0.8 0.020 7.8 0.00 0.99780 3.51 0.56 7.0 0.99680 3.20 0.68 7.0 0.99780 3.16 0.58 7.0 0.99780 3.16 0.58 7.0 0.99780 3.15 0.56 7.0 0.99780 3.15 0.56 7.0 0.99780 3.15 0.56 7.0 0.99780 3.15 0.56 7.0 0.99780 3.15 0.56 7.0 0.99780 3.15 0.46 7.0 0.99780 3.15 0.46 7.0 0.99780 3.15 0.46 7.0 0.99254 2.99 0.46 7.0 0.99554 2.99 0.46

alcohol quality 0 9.4 5

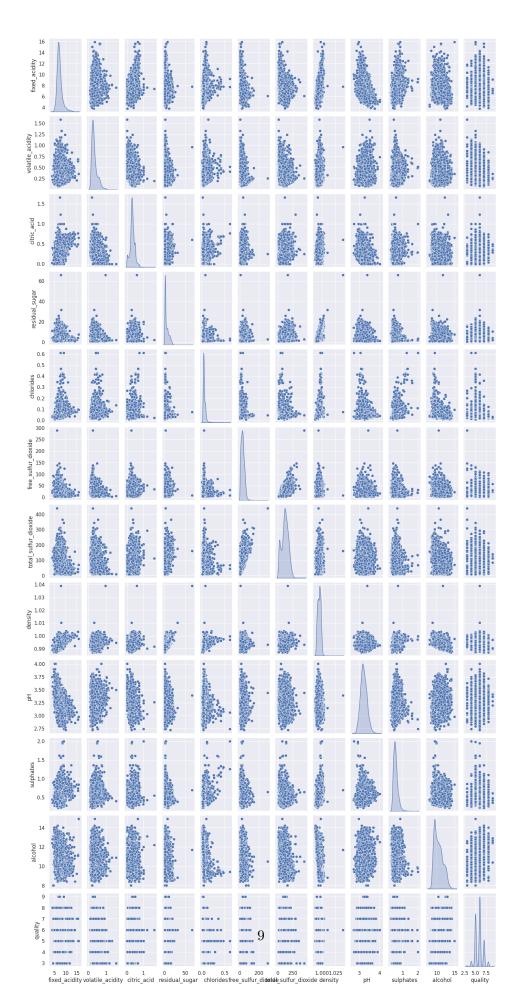
```
9.8
                       5
1
2
           9.8
                       5
3
           9.8
                       6
                       5
4
           9.4
6492
          11.2
                       6
           9.6
                       5
6493
6494
           9.4
                       6
                       7
6495
          12.8
6496
          11.8
                       6
```

[6497 rows x 12 columns]

Inspecting the relationships between the features or components using a pair plot and a heatmap

```
[33]: # pair plot
sns.pairplot(df, aspect=0.5 , diag_kind='kde')
```

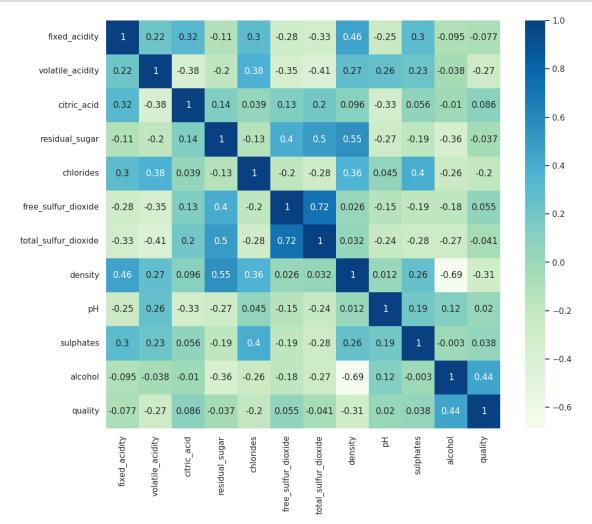
[33]: <seaborn.axisgrid.PairGrid at 0x7f57c79521a0>



This is a histogram pairplot of quality vs each individual property. The chemical property is on the x-axis, while the quality is on the y-axis. This histogram pairplot shows the count of a particular rank at a particular value of the property. For example for fixed acidity, there are a of quality=6 wines in the range of 5 to 10 for fixed acidity vs, the range of 10 to 15. As the wine quality increases, more the lower range of the fixed acidity has more points than the upper range.

Next, plot a heatmap to visualize features are correlated with each other.

```
[34]: # Heatmap
plt.figure(figsize=(12,10))
cor = df.corr()
sns.heatmap(cor, annot=True, cmap=plt.cm.GnBu)
plt.show()
```

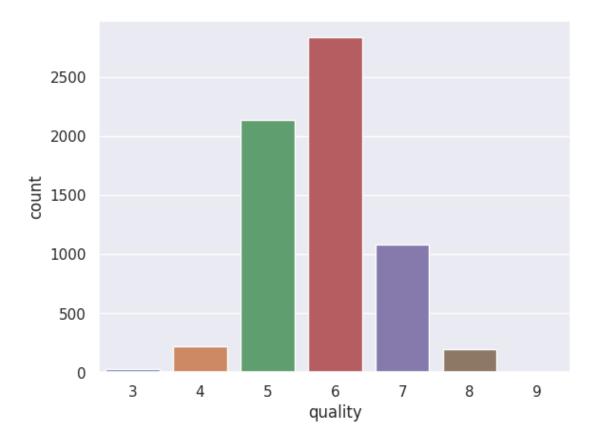


The Heat map shows the correlation between each of the properties. The highest correlation seems

to be between total sulfur dioxide and the color. This could mean that a particular color of wine has more total sulfur dioxide than the other color. For determining, which factor affects the quality the most, the highest correlation seems to be between quality and alcohol content.

```
[35]: sns.countplot(x=df["quality"])
```

[35]: <Axes: xlabel='quality', ylabel='count'>



Here is is distribution of the wines by quality. They are centered around 5, 6 and 7. This could be a downside for our model building.

```
[36]: df["quality"] =df["quality"].astype(int)
df = pd.get_dummies(df, columns=["quality"])

df.head(5)
```

```
[36]:
         fixed_acidity
                         volatile_acidity
                                             citric_acid residual_sugar
                                                                            chlorides
      0
                    7.4
                                       0.70
                                                     0.00
                                                                       1.9
                                                                                 0.076
                    7.8
                                       0.88
                                                     0.00
                                                                       2.6
                                                                                 0.098
      1
                                                     0.04
      2
                    7.8
                                      0.76
                                                                       2.3
                                                                                 0.092
      3
                   11.2
                                      0.28
                                                     0.56
                                                                       1.9
                                                                                 0.075
      4
                    7.4
                                       0.70
                                                     0.00
                                                                                 0.076
                                                                       1.9
```

```
free_sulfur_dioxide
                         total_sulfur_dioxide
                                                  density
                                                                  sulphates
                                                              рΗ
0
                   11.0
                                           34.0
                                                   0.9978
                                                            3.51
                                                                        0.56
                   25.0
                                           67.0
                                                                        0.68
1
                                                   0.9968
                                                            3.20
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
   alcohol
            quality_3
                         quality_4 quality_5 quality_6
                                                            quality_7
                                                                         quality 8
0
       9.4
       9.8
                     0
                                  0
                                              1
                                                          0
                                                                      0
                                                                                  0
1
2
       9.8
                     0
                                  0
                                              1
                                                          0
                                                                      0
                                                                                  0
3
       9.8
                      0
                                  0
                                              0
                                                          1
                                                                      0
                                                                                  0
       9.4
4
                      0
                                  0
                                              1
                                                          0
                                                                      0
                                                                                  0
   quality_9
0
            0
            0
1
2
            0
           0
3
            0
```

5 Modeling the Data

5.1 Preparing the Data

First we prepare the data so that we do not have bias later. We will also split the data into training and test set. The training set will be 75% and the test set will be 25%.

```
(4872, 11) (4872, 6) (1625, 11) (1625, 6)
```

5.2 Method

1. Evaluate Models Build a model

- 2. Make Predictions or Plot the Accuracy Make predictions and find accuracy or metrics of the model.
- 3. Analyze Analyze and make conclusion about the fit and accuracy of the model.

5.2.1 Model Architecture

With a stronger sense of the training data and the goal, the next discussion is which model architecture to use and train for best results. Selecting the right model architecture is important because it will use the right amount of resources for the task, will not overfit the data, and will give accurate classification results.

5.2.2 Options for Architecture

The loss function must be selected, with possible choices including mean absolute error (mae), mean squared error (mse), and Huber loss for regression loss calculations. This problem is a categorical classification problem so there are the options of using Categorical Cross Entropy.

Additionally the activation function for the linear units must be selected. Choices here include Sigmoid, TanH, and ReLU. Since ReLU has shown great promise with training Neural Network models and since it also can increase the speed of the model learning, that will be one of the activation function selected here.

Additionally, it is possible to select strategies to optimize the hyperparameters of an Neural Network model and one of the most popular and effective strategies is Stochastic Gradient Descent. This will be the selected strategy for optimizing the Neural Network to reduce loss.

```
fig, axes = plt.subplots(1, 2, figsize = (8, 3))

axes[0].plot(classifier.epoch, classifier.history[acc], label = 'acc')
axes[0].set_title('Accuracy vs Epochs', fontsize = 12)
axes[0].set_xlabel('Epochs', fontsize = 10)
axes[0].set_ylabel('Accuracy', fontsize = 10)
axes[0].legend()

axes[1].plot(classifier.epoch, classifier.history['loss'], label = 'loss')
axes[1].set_title("Loss Curve",fontsize=12)
axes[1].set_xlabel("Epochs",fontsize=10)
axes[1].set_ylabel("Loss",fontsize=10)
axes[1].legend()

plt.show()
```

5.3 Basic Stochastic Gradient Descent

Stochastic gradient descent (SGD) is a method for finding a minimum of a function by iteratively taking steps in the direction of the negative gradient of the function at the current point. The steps are chosen randomly, hence the name "stochastic". SGD is a popular algorithm for training neural networks.

SGD works by repeatedly taking steps in the direction of the negative gradient of the loss function. The loss function is a measure of how well the model fits the training data. The gradient of the loss function is a vector that points in the direction of greatest increase in the loss function. By taking steps in the opposite direction of the gradient, SGD can find a minimum of the loss function.

```
[39]: mod1 = keras.Sequential([
    layers.Dense(512, activation='relu', input_shape=[11]),
    layers.Dense(512, activation='relu'),
    layers.Dense(512, activation='relu'),
    layers.Dense(1),
])
```

[40]: mod1.summary()

Model: "sequential_2"

Layer (type)	Output Shape	Param #
dense_4 (Dense)	(None, 512)	6144
dense_5 (Dense)	(None, 512)	262656
dense_6 (Dense)	(None, 512)	262656
dense_7 (Dense)	(None, 1)	513

Total params: 531,969 Trainable params: 531,969 Non-trainable params: 0

```
batch_size=256,
  epochs=100,
)
```

```
Epoch 1/100
0.8340 - val_loss: 0.1664 - val_accuracy: 0.8344
Epoch 2/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 3/100
0.8340 - val_loss: 0.1656 - val_accuracy: 0.8344
Epoch 4/100
0.8340 - val_loss: 0.1664 - val_accuracy: 0.8344
Epoch 5/100
0.8340 - val_loss: 0.1662 - val_accuracy: 0.8344
Epoch 6/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 7/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 8/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 9/100
20/20 [============= ] - 1s 29ms/step - loss: 0.1663 - accuracy:
0.8340 - val_loss: 0.1660 - val_accuracy: 0.8344
Epoch 10/100
0.8340 - val_loss: 0.1660 - val_accuracy: 0.8344
Epoch 11/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 12/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 13/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 14/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 15/100
```

```
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 16/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 17/100
20/20 [============= ] - 1s 27ms/step - loss: 0.1663 - accuracy:
0.8340 - val_loss: 0.1661 - val_accuracy: 0.8344
Epoch 18/100
0.8340 - val_loss: 0.1662 - val_accuracy: 0.8344
Epoch 19/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 20/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 21/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 22/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 23/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 24/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 25/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 26/100
0.8340 - val_loss: 0.1661 - val_accuracy: 0.8344
Epoch 27/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 28/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 29/100
20/20 [============ ] - 1s 28ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 30/100
0.8340 - val_loss: 0.1660 - val_accuracy: 0.8344
Epoch 31/100
```

```
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 32/100
0.8340 - val_loss: 0.1660 - val_accuracy: 0.8344
Epoch 33/100
20/20 [============= ] - 1s 27ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 34/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 35/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 36/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 37/100
0.8340 - val_loss: 0.1656 - val_accuracy: 0.8344
Epoch 38/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 39/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 40/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 41/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 42/100
20/20 [============ ] - 1s 28ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 43/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 44/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 45/100
20/20 [============ ] - 1s 28ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 46/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 47/100
```

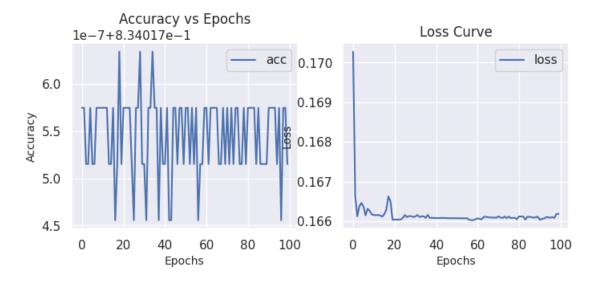
```
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 48/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 49/100
20/20 [============= ] - 1s 48ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 50/100
20/20 [============= ] - 1s 30ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 51/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 52/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 53/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 54/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 55/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 56/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 57/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 58/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 59/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 60/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 61/100
20/20 [============ ] - 1s 27ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 62/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 63/100
```

```
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 64/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 65/100
20/20 [============ ] - 1s 26ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 66/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 67/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 68/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 69/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 70/100
20/20 [============= ] - 1s 48ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 71/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 72/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 73/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 74/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 75/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 76/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 77/100
20/20 [============ ] - 1s 27ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 78/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 79/100
```

```
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 80/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 81/100
20/20 [============ ] - 1s 28ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 82/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 83/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 84/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 85/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 86/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 87/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 88/100
0.8340 - val_loss: 0.1656 - val_accuracy: 0.8344
Epoch 89/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 90/100
20/20 [============ ] - 1s 39ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1656 - val_accuracy: 0.8344
Epoch 91/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 92/100
0.8340 - val_loss: 0.1659 - val_accuracy: 0.8344
Epoch 93/100
20/20 [============ ] - 1s 39ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 94/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 95/100
```

```
0.8340 - val_loss: 0.1656 - val_accuracy: 0.8344
Epoch 96/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 97/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 98/100
20/20 [=====
           =========] - Os 25ms/step - loss: 0.1661 - accuracy:
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
Epoch 99/100
0.8340 - val_loss: 0.1658 - val_accuracy: 0.8344
Epoch 100/100
0.8340 - val_loss: 0.1657 - val_accuracy: 0.8344
```

[43]: model_performance_graphs(history1, "accuracy")



```
[44]: train1_acc = mod1.evaluate(x = X_train, y = Y_train)
    print()
    print ("Train Accuracy = " + str(train1_acc[1]))
    print()
    test1_acc = mod1.evaluate(x = X_test, y = Y_test)
    print()
    print ("Loss = " + str(test1_acc[0]))
    print ("Test Accuracy = " + str(test1_acc[1]))
```

The accuracy of our model is pretty high. We can attempt to improve the accuracy by utilizing other techniques

5.4 Dropout and Batch Normalization

Dropout and batch normalization are two techniques used to improve the performance of neural networks. Dropout works by randomly dropping out (setting to zero) a certain percentage of neurons during each training epoch. This forces the network to learn more robust features that are not dependent on any single neuron. Batch normalization works by normalizing the inputs to each layer of the network, which helps to prevent the values from becoming too large or too small.

```
[45]: mod2 = Sequential()
      # layer 1
      mod2.add(Dense(30, input_dim=11, activation='relu',__
       →name='fc0',kernel_regularizer=regularizers.12(0.01)))
      mod2.add(BatchNormalization(momentum=0.99, epsilon=0.001))
      #layer 2
      mod2.add(Dense(50, name='fc1',bias_initializer='zeros'))
      mod2.add(Activation('tanh'))
      mod2.add(BatchNormalization(momentum=0.99, epsilon=0.001))
      mod2.add(Dropout(0.5))
      #layer 3
      mod2.add(Dense(100, name='fc2',bias_initializer='zeros'))
      mod2.add(Activation('relu'))
      mod2.add(BatchNormalization(momentum=0.99, epsilon=0.001))
      mod2.add(Dropout(0.5))
      #layer 4
      mod2.add(Dense(6, name='fc3',bias_initializer='zeros'))
      mod2.add(Activation('softmax'))
      mod2.add(BatchNormalization(momentum=0.99, epsilon=0.001))
```

```
[46]: mod2.summary()
```

```
Model: "sequential_3"
```

Layer (type)	Output Shape	Param #
fc0 (Dense)	(None, 30)	360

```
batch_normalization_4 (Batc (None, 30)
                                               120
    hNormalization)
     fc1 (Dense)
                           (None, 50)
                                               1550
     activation 3 (Activation)
                           (None, 50)
                                               0
     batch_normalization_5 (Batc (None, 50)
                                               200
    hNormalization)
     dropout_2 (Dropout)
                           (None, 50)
                                               0
    fc2 (Dense)
                           (None, 100)
                                               5100
     activation_4 (Activation)
                           (None, 100)
                                               400
     batch_normalization_6 (Batc (None, 100)
    hNormalization)
                           (None, 100)
     dropout_3 (Dropout)
                                               0
     fc3 (Dense)
                           (None, 6)
                                               606
    activation_5 (Activation)
                           (None, 6)
    batch_normalization_7 (Batc (None, 6)
                                               24
    hNormalization)
    ______
    Total params: 8,360
    Trainable params: 7,988
    Non-trainable params: 372
    ______
[47]: Adam = optimizers.Adam(learning_rate=0.001, beta_1=0.9, beta_2=0.999,__
     ⇒epsilon=1e-08)
    mod2.compile(optimizer = Adam, loss = "mae", metrics = ["accuracy"])
[48]: history2 = mod2.fit(x = X_train, y = Y_train, epochs = 100, verbose=1,__
     →batch_size = 64,validation_data=(X_test, Y_test))
    Epoch 1/100
    0.1882 - val_loss: 0.3982 - val_accuracy: 0.1631
    Epoch 2/100
    0.2377 - val_loss: 0.3616 - val_accuracy: 0.1729
```

```
Epoch 3/100
0.2917 - val_loss: 0.3279 - val_accuracy: 0.4215
Epoch 4/100
0.3502 - val_loss: 0.3253 - val_accuracy: 0.4215
Epoch 5/100
0.3890 - val_loss: 0.2613 - val_accuracy: 0.4215
Epoch 6/100
0.4438 - val_loss: 0.2515 - val_accuracy: 0.4289
Epoch 7/100
0.4516 - val_loss: 0.2518 - val_accuracy: 0.4843
Epoch 8/100
0.4645 - val_loss: 0.2365 - val_accuracy: 0.4720
Epoch 9/100
0.4750 - val_loss: 0.2075 - val_accuracy: 0.5058
Epoch 10/100
0.4873 - val_loss: 0.2112 - val_accuracy: 0.4751
Epoch 11/100
0.4971 - val_loss: 0.2494 - val_accuracy: 0.4185
Epoch 12/100
0.4984 - val_loss: 0.2180 - val_accuracy: 0.4412
Epoch 13/100
0.5082 - val_loss: 0.1904 - val_accuracy: 0.4929
Epoch 14/100
0.5064 - val_loss: 0.1846 - val_accuracy: 0.4862
Epoch 15/100
0.5101 - val_loss: 0.1884 - val_accuracy: 0.4609
Epoch 16/100
0.5080 - val_loss: 0.1873 - val_accuracy: 0.4695
Epoch 17/100
0.5113 - val_loss: 0.1799 - val_accuracy: 0.4837
Epoch 18/100
0.5226 - val_loss: 0.1684 - val_accuracy: 0.5182
```

```
Epoch 19/100
0.5205 - val_loss: 0.1763 - val_accuracy: 0.4917
Epoch 20/100
0.5261 - val_loss: 0.1622 - val_accuracy: 0.5329
Epoch 21/100
0.5168 - val_loss: 0.1657 - val_accuracy: 0.5255
Epoch 22/100
0.5287 - val_loss: 0.1687 - val_accuracy: 0.5108
Epoch 23/100
0.5220 - val_loss: 0.1686 - val_accuracy: 0.5175
Epoch 24/100
0.5174 - val_loss: 0.1839 - val_accuracy: 0.4597
Epoch 25/100
0.5255 - val_loss: 0.1599 - val_accuracy: 0.5274
Epoch 26/100
0.5277 - val_loss: 0.1902 - val_accuracy: 0.2849
Epoch 27/100
0.5244 - val_loss: 0.1879 - val_accuracy: 0.2154
Epoch 28/100
0.5271 - val_loss: 0.1669 - val_accuracy: 0.5108
Epoch 29/100
0.5189 - val_loss: 0.1599 - val_accuracy: 0.5286
Epoch 30/100
0.5308 - val_loss: 0.1855 - val_accuracy: 0.4505
Epoch 31/100
0.5086 - val_loss: 0.1647 - val_accuracy: 0.5040
Epoch 32/100
0.5216 - val_loss: 0.1697 - val_accuracy: 0.4228
0.5183 - val_loss: 0.1700 - val_accuracy: 0.4234
Epoch 34/100
0.5193 - val_loss: 0.1698 - val_accuracy: 0.4209
```

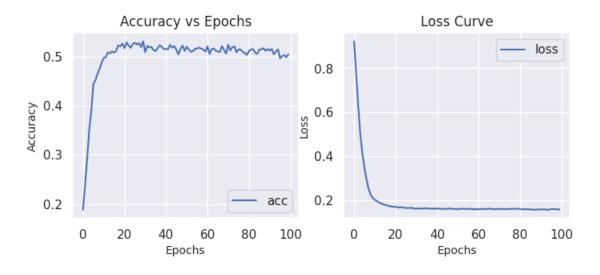
```
Epoch 35/100
0.5129 - val_loss: 0.1691 - val_accuracy: 0.4332
Epoch 36/100
0.5109 - val_loss: 0.1556 - val_accuracy: 0.5440
Epoch 37/100
0.5170 - val_loss: 0.1706 - val_accuracy: 0.4295
Epoch 38/100
0.5226 - val_loss: 0.1758 - val_accuracy: 0.4215
Epoch 39/100
0.5195 - val_loss: 0.1692 - val_accuracy: 0.3520
Epoch 40/100
0.5148 - val_loss: 0.1746 - val_accuracy: 0.4178
Epoch 41/100
0.5150 - val_loss: 0.1659 - val_accuracy: 0.4732
Epoch 42/100
0.5148 - val_loss: 0.1563 - val_accuracy: 0.5372
Epoch 43/100
0.5236 - val_loss: 0.1684 - val_accuracy: 0.5028
Epoch 44/100
0.5181 - val_loss: 0.1556 - val_accuracy: 0.5311
Epoch 45/100
0.5209 - val_loss: 0.1695 - val_accuracy: 0.4283
Epoch 46/100
0.5135 - val_loss: 0.1703 - val_accuracy: 0.4166
Epoch 47/100
0.5041 - val_loss: 0.1690 - val_accuracy: 0.4222
Epoch 48/100
0.5156 - val_loss: 0.1686 - val_accuracy: 0.4437
Epoch 49/100
0.5222 - val_loss: 0.1654 - val_accuracy: 0.4615
Epoch 50/100
0.5113 - val_loss: 0.1614 - val_accuracy: 0.5200
```

```
Epoch 51/100
0.5195 - val_loss: 0.1802 - val_accuracy: 0.4597
Epoch 52/100
0.5142 - val_loss: 0.1984 - val_accuracy: 0.4037
Epoch 53/100
0.5096 - val_loss: 0.1634 - val_accuracy: 0.5083
Epoch 54/100
0.5105 - val_loss: 0.1673 - val_accuracy: 0.4997
Epoch 55/100
0.5154 - val_loss: 0.1553 - val_accuracy: 0.5397
Epoch 56/100
0.5160 - val_loss: 0.1672 - val_accuracy: 0.5034
Epoch 57/100
0.5181 - val_loss: 0.1557 - val_accuracy: 0.5323
Epoch 58/100
0.5156 - val_loss: 0.1686 - val_accuracy: 0.4295
Epoch 59/100
0.5142 - val_loss: 0.1605 - val_accuracy: 0.5225
Epoch 60/100
0.5099 - val_loss: 0.1583 - val_accuracy: 0.5182
Epoch 61/100
0.5203 - val_loss: 0.1555 - val_accuracy: 0.5262
Epoch 62/100
0.5049 - val_loss: 0.1863 - val_accuracy: 0.4418
Epoch 63/100
0.5150 - val_loss: 0.1596 - val_accuracy: 0.5286
Epoch 64/100
0.5160 - val_loss: 0.1528 - val_accuracy: 0.5391
0.5117 - val_loss: 0.1652 - val_accuracy: 0.4812
Epoch 66/100
0.5099 - val_loss: 0.1609 - val_accuracy: 0.4991
```

```
Epoch 67/100
0.5092 - val_loss: 0.1690 - val_accuracy: 0.3458
Epoch 68/100
0.5209 - val_loss: 0.1569 - val_accuracy: 0.5385
Epoch 69/100
0.5131 - val_loss: 0.1559 - val_accuracy: 0.5372
Epoch 70/100
0.5055 - val_loss: 0.1888 - val_accuracy: 0.4369
Epoch 71/100
0.5238 - val_loss: 0.1523 - val_accuracy: 0.5397
Epoch 72/100
0.5123 - val_loss: 0.1531 - val_accuracy: 0.5446
Epoch 73/100
0.5189 - val_loss: 0.1554 - val_accuracy: 0.5194
Epoch 74/100
0.5203 - val_loss: 0.1684 - val_accuracy: 0.4985
Epoch 75/100
0.5084 - val_loss: 0.1600 - val_accuracy: 0.5132
Epoch 76/100
0.5144 - val_loss: 0.1699 - val_accuracy: 0.4794
Epoch 77/100
0.5140 - val_loss: 0.1576 - val_accuracy: 0.5372
Epoch 78/100
0.5092 - val_loss: 0.1674 - val_accuracy: 0.4978
Epoch 79/100
0.5072 - val_loss: 0.1623 - val_accuracy: 0.5243
Epoch 80/100
0.5031 - val_loss: 0.1565 - val_accuracy: 0.5323
0.5107 - val_loss: 0.1567 - val_accuracy: 0.5298
Epoch 82/100
0.5142 - val_loss: 0.1645 - val_accuracy: 0.4862
```

```
Epoch 83/100
0.5148 - val_loss: 0.1613 - val_accuracy: 0.4991
Epoch 84/100
0.5076 - val_loss: 0.1617 - val_accuracy: 0.4898
Epoch 85/100
0.5049 - val_loss: 0.1563 - val_accuracy: 0.5120
Epoch 86/100
0.5142 - val_loss: 0.1548 - val_accuracy: 0.5274
Epoch 87/100
0.5142 - val_loss: 0.1551 - val_accuracy: 0.5169
Epoch 88/100
0.5168 - val_loss: 0.1541 - val_accuracy: 0.5225
Epoch 89/100
0.5111 - val_loss: 0.1596 - val_accuracy: 0.5095
Epoch 90/100
0.5152 - val_loss: 0.1545 - val_accuracy: 0.5243
Epoch 91/100
0.5121 - val_loss: 0.1540 - val_accuracy: 0.5231
Epoch 92/100
0.5152 - val_loss: 0.1529 - val_accuracy: 0.5243
Epoch 93/100
0.5043 - val_loss: 0.1610 - val_accuracy: 0.4985
Epoch 94/100
0.5101 - val_loss: 0.1516 - val_accuracy: 0.5323
Epoch 95/100
0.5144 - val_loss: 0.1631 - val_accuracy: 0.4948
Epoch 96/100
0.4963 - val_loss: 0.1639 - val_accuracy: 0.4874
Epoch 97/100
0.5006 - val_loss: 0.1587 - val_accuracy: 0.5286
Epoch 98/100
0.5027 - val_loss: 0.1567 - val_accuracy: 0.5120
```

[49]: model_performance_graphs(history2, "accuracy")



```
[50]: train2_acc = mod2.evaluate(x = X_train, y = Y_train)
     print()
     print ("Train Accuracy = " + str(train2_acc[1]))
     print()
     test2_acc = mod2.evaluate(x = X_test, y = Y_test)
     print()
     print ("Loss = " + str(test2_acc[0]))
     print ("Test Accuracy = " + str(test2_acc[1]))
     153/153 [============ ] - Os 2ms/step - loss: 0.1566 -
     accuracy: 0.5230
     Train Accuracy = 0.522988498210907
     51/51 [=====
                               =======] - Os 3ms/step - loss: 0.1520 - accuracy:
     0.5311
     Loss = 0.15203529596328735
     Test Accuracy = 0.5310769081115723
```

Unfortunately, the accuracy of our model reduced when utilizing dropout and batch normalization. When the network is small relative to the dataset, regularization is usually unnecessary. If the

model capacity is already low, lowering it further by adding regularization will hurt performance. Additionally, there are some cases where Batch Normalization can actually hurt the performance of a neural network. For example, if the batch size is too small, Batch Normalization can introduce noise into the training process. Additionally, if the network is very deep, Batch Normalization can make it difficult for the network to learn long-range dependencies.

6 Conclusion

In conclusion, I first used a basic stochastic gradient descent algorithm with our wine data. We got an accuracy of nearly 85%. When I introduced dropout and batch normalization, I saw a decrease in our accuracy.

This is the best I could obtain without overfitting. The algorithm might perform better with a better distribution among the different scores. An other explanation might be that wine testing is strongly subjective.

Neural network models are a powerful tool for machine learning. They can be used to solve a wide variety of problems, including classification. However, neural network models can be difficult to train and can be prone to overfitting.

Overfitting is a phenomenon in machine learning where a model performs well on the training data but poorly on the test data. This happens when the model learns the noise in the training data instead of the underlying patterns. While there was a disparity in the accuracies of the two different models, there was little to no difference between the training and testing accuracies of each model.

Overall, neural network models are a powerful tool for machine learning. However, it is important to be aware of their limitations and to use them carefully.

6.0.1 References

- https://www.kaggle.com/code/ryanholbrook/a-single-neuron
- https://www.kaggle.com/code/davidzarebski/a-neural-net-for-red-wine-quality-estimation -https://www.kaggle.com/code/rprkh15/red-wine-quality-xgboost-optuna-neural-networks#Prediction-Using-Neural-Networks