## PROJECT 4 WINE ANALYSIS

#### **Problem Statement**

VinoVista, a renowned winery, is committed to producing consistently exceptional wines. However, they have observed variability in the quality of their white wine batches. To address this challenge, VinoVista seeks a predictive model that can accurately assess the quality of wine batches based on their chemical properties before bottling.

#### **Project Goal:**

The primary objective of this project is to develop a robust machine learning model capable of predicting wine quality on a scale of 0-10. This model will utilize a dataset containing various chemical properties of wine, such as acidity, pH, residual sugar, chlorides, free sulfur dioxide, total sulfur dioxide, density, sulfates, and alcohol content.

#### **Resources:**

Wine Quality Dataset from UC Irvine Machine Learning Repository with 4899 samples

### **Predicting Wine Quality**

#### **Collaborators:**

**Chuck Bui** 

**Jack Jeffries** 

**Beau Massie** 

**Christopher Turner** 

# **QUICK TAKE**

- This data set was chosen because it contains information about the chemical propterties of white wines, such as acidity, sugar levels, and alcohol content and the data is modeled after the physicochemical wine tests.
- The number of samples in our dataset after filtering is 3,090. We used Quality as our target and there
  are 11 total features.
- We cleaned our database by checking for null values, dropping any outliers in 'total sulfur dioxide' above 150 as that is likely a data entry error. Our quality values range from 1-10, so we created binary classification to use in our machine learning models. We chose 7 and above as good; 6 and below as not good.
- We checked for multi collinearity among our features for possible features that could be dropped to increase our precision and recall values and filtered our data to only keep wines that pH values between 3 and 4, as this is the nominal level for white wines.

### LOAD THE DATA AND BEGIN CLEANING

df.info()

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	рΗ	sulphates	alcohol	quality
0	7.0	0.27	0.36	20.7	0.045	45.0	170.0	1.0010	3.00	0.45	8.8	6
1	6.3	0.30	0.34	1.6	0.049	14.0	132.0	0.9940	3.30	0.49	9.5	6
2	8.1	0.28	0.40	6.9	0.050	30.0	97.0	0.9951	3.26	0.44	10.1	6
3	7.2	0.23	0.32	8.5	0.058	47.0	186.0	0.9956	3.19	0.40	9.9	6
4	7.2	0.23	0.32	8.5	0.058	47.0	186.0	0.9956	3.19	0.40	9.9	6

```
# check for nulls ( no nulls), and data type
df.info()
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 4898 entries, 0 to 4897
Data columns (total 12 columns):

# Column Non-Null Count Dtype fixed acidity 4898 non-null float64 volatile acidity 4898 non-null citric acid 4898 non-null float64 residual sugar 4898 non-null float64 chlorides 4898 non-null float64 free sulfur dioxide 4898 non-null float64 total sulfur dioxide 4898 non-null float64 density 4898 non-null float64 4898 non-null float64 9 sulphates 4898 non-null float64 4898 non-null float64 10 alcohol 11 quality 4898 non-null int64

dtypes: float64(11), int64(1) memory usage: 459.3 KB

```
# The scale for wine quality is 1-10, so we create a binary classification to use for our ML models
df['quality_label'] = np.where(df['quality'] >= 7, 'good', 'not good')
df.drop('quality', axis=1, inplace=True)
```

```
<class 'pandas.core.frame.DataFrame'>
Index: 3090 entries, 1 to 4897
Data columns (total 12 columns):
    Column
                          Non-Null Count Dtype
    fixed acidity
                          3090 non-null
                                          float64
    volatile acidity
                          3090 non-null
                                          float64
    citric acid
                          3090 non-null
                                          float64
    residual sugar
                          3090 non-null
                                          float64
    chlorides
                                          float64
                          3090 non-null
    free sulfur dioxide
                          3090 non-null
                                          float64
    total sulfur dioxide
                                          float64
                          3090 non-null
                                          float64
    density
                          3090 non-null
    pН
                          3090 non-null
                                          float64
    sulphates
                          3090 non-null
                                          float64
    alcohol
                          3090 non-null
                                          float64
11 quality label
                          3090 non-null
                                          object
dtypes: float64(11), object(1)
memory usage: 313.8+ KB
```

### **CODING PAGE 2**

```
# convert categorical variables into numerical ones using one-hot encoding to use in our model
df = pd.get_dummies(df, columns=['quality_label'], drop_first=True)
```

df.head()

:	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	рН	sulphates	alcohol	quality_label_not good
1	6.3	0.30	0.34	1.6	0.049	14.0	132.0	0.9940	3.30	0.49	9.5	True
2	8.1	0.28	0.40	6.9	0.050	30.0	97.0	0.9951	3.26	0.44	10.1	True
5	8.1	0.28	0.40	6.9	0.050	30.0	97.0	0.9951	3.26	0.44	10.1	True
6	6.2	0.32	0.16	7.0	0.045	30.0	136.0	0.9949	3.18	0.47	9.6	True
8	6.3	0.30	0.34	1.6	0.049	14.0	132.0	0.9940	3.30	0.49	9.5	True

# check that shape
df.shape

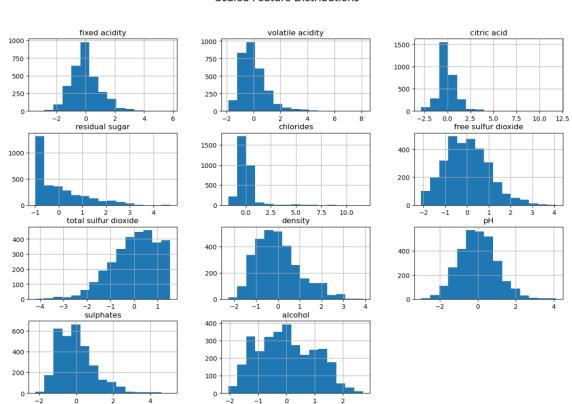
(3090, 12)

```
# separate features from target
y=df['quality_label_not good']
X=df.drop(columns='quality_label_not good')
```

```
# clean it up, scale it etc
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
X_scaled_df = pd.DataFrame(X_scaled, columns=X.columns)
```

```
# bin our data to view distributions of our scaled data
X_scaled_df.hist(bins=15, figsize=(15, 10))
plt.suptitle('Scaled Feature Distributions', fontsize=16)
plt.show()
```

#### Scaled Feature Distributions



### **CODING PAGE 3**

(2317, 11)

```
LogisticRegression

LogisticRegression(max_iter=200, random_state=1776)
```

```
# predictions and confusion matrix
testing_predictions = classifier.predict(X_test)
test_matrix = confusion_matrix(y_test, testing_predictions)
print(test_matrix)
```

```
[[ 78 133]
[ 56 506]]
```

# classification report for our standard data
test\_report = classification\_report(y\_test, testing\_predictions)
print(test\_report)

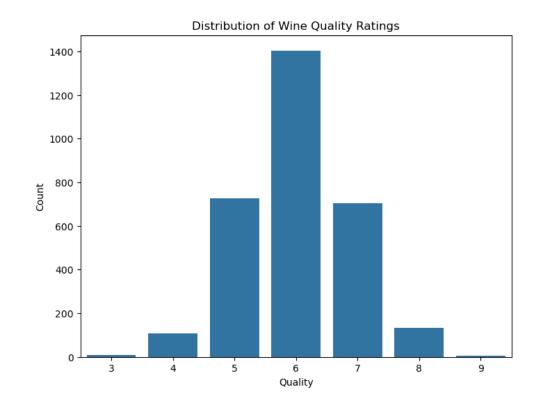
precision	recall	†1-score	support
0.58	0.37	0.45	211
0.79	0.90	0.84	562
		0.76	773
0.69	0.64	0.65	773
0.73	0.76	0.74	773
	0.79	0.58 0.37 0.79 0.90 0.69 0.64	0.58 0.37 0.45 0.79 0.90 0.84 0.76 0.69 0.64 0.65

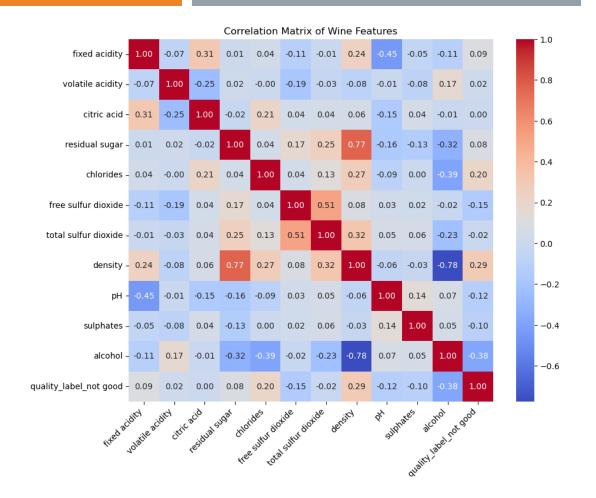
### CODING PAGE 4

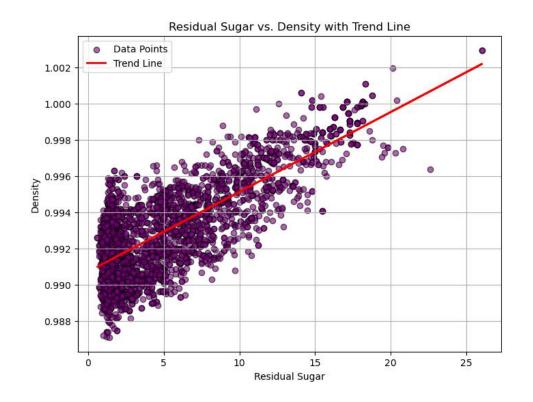
```
# xgboost model - our best performing model
from xgboost import XGBClassifier
|
xgb_clf = XGBClassifier(random_state=1776, scale_pos_weight=len(y_train[y_train == 0]) / len(y_train[y_train == 1]))
xgb_clf.fit(X_train, y_train)
predictions = xgb_clf.predict(X_test)
print(classification_report(y_test, predictions))
```

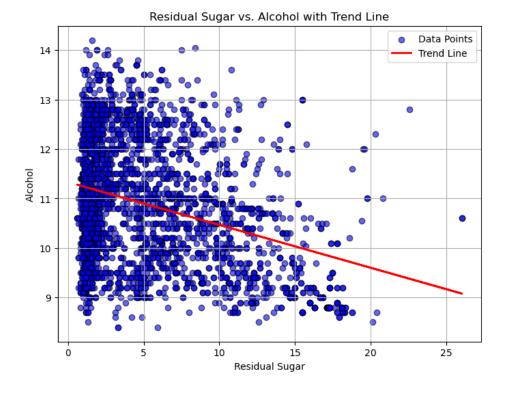
	precision	recall	f1-score	support
False	0.71	0.68	0.69	211
True	0.88	0.90	0.89	562
accuracy			0.84	773
macro avg	0.80	0.79	0.79	773
weighted avg	0.83	0.84	0.84	773

### VIZ WALKTHROUGH PAGE I

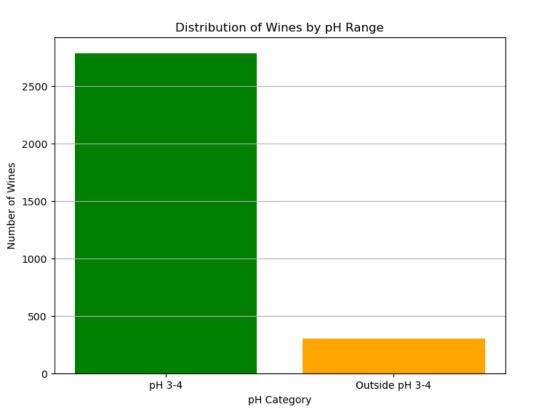


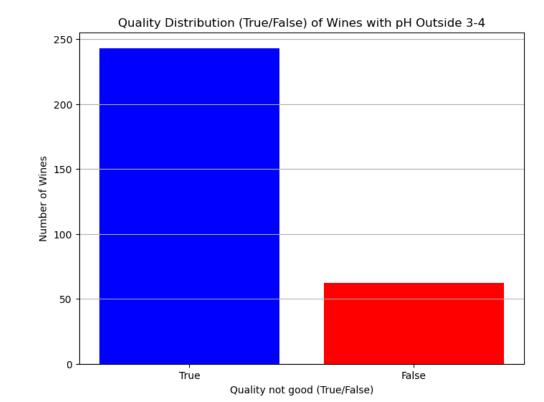






## VIZ WALKTHROUGH PAGE 3





## **CONCLUSION**

# **Data Insights**

- Residual sugar and density are critical features for predicting wine quality.
- Chemicals like pH, alcohol content, and chlorides also play a significant role

# **Future Steps**

- Incorporate external features like region or grape variety
- Chemicals like pH, alcohol content, and chlorides also play a significant role

## WRAP AND SUGGESTIONS TO COMPANY

# suggestions

- Suggestions for packaging wine or marketing directly to the consumer
- Teach the consumer with features on the labels and allow them to be part of the process, thus making them feel more loyal to the brand