

# Wine Quality EDA

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## Getting started

### Importing libraries

```
In [ ]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
plt.style.available
plt.style.use('seaborn-pastel')
import warnings
warnings.filterwarnings("ignore")
```

```
C:\Users\Thanh\AppData\Local\Temp\ipykernel_77288\1651659601.py:6: MatplotlibDeprecationWarning: The seaborn styles shipped by Matplotlib are deprecated since 3.6, as they no longer correspond to the styles shipped by seaborn. However, they will remain available as 'seaborn-v0_8-<style>'. Alternatively, directly use the seaborn API instead.
plt.style.use('seaborn-pastel')
```

## Importing dataset


```
In [ ]: df = pd.read_csv('wineqt.csv', index_col='Id')
```

## Quick overview about data

```
In [ ]: df.head()
```

```
Out[ ]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
<b>Id</b>										
<b>0</b>	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56
<b>1</b>	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68
<b>2</b>	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65
<b>3</b>	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58
<b>4</b>	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56



No categoricial data within the dataset

## Check Null

```
In [ ]: null_counts = df.isnull().sum()

print(null_counts)
```

```
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
```

## Check for missing data and data types

```
In [ ]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Index: 1143 entries, 0 to 1597
Data columns (total 12 columns):
 #   Column                Non-Null Count  Dtype  
---  -
 0   fixed acidity          1143 non-null   float64
 1   volatile acidity       1143 non-null   float64
 2   citric acid            1143 non-null   float64
 3   residual sugar         1143 non-null   float64
 4   chlorides              1143 non-null   float64
 5   free sulfur dioxide    1143 non-null   float64
 6   total sulfur dioxide   1143 non-null   float64
 7   density                1143 non-null   float64
 8   pH                    1143 non-null   float64
 9   sulphates              1143 non-null   float64
10   alcohol                1143 non-null   float64
11   quality                1143 non-null   int64  
dtypes: float64(11), int64(1)
memory usage: 116.1 KB
```

There are no missing values in the dataset. The target column consists of integer values which have an ordinary character. The rest of the values are float. <\html>

## Check Duplicate

```
In [ ]: data = df.drop(['quality'], axis=1)
duplicate_count = data.duplicated().sum()
print("Number of duplicate rows:", duplicate_count)
```

Number of duplicate rows: 125

## Analysis of statistic distribution


Conduct exploratory data analysis (EDA) to gain insights into the dataset

a) Calculate basic statistics for numerical columns:

```
In [ ]: df.describe()
```

Out[ ]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide
count	1143.000000	1143.000000	1143.000000	1143.000000	1143.000000	1143.000000	1143.000000
mean	8.311111	0.531339	0.268364	2.532152	0.086933	15.615486	45.000000
std	1.747595	0.179633	0.196686	1.355917	0.047267	10.250486	32.000000
min	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	6.000000
25%	7.100000	0.392500	0.090000	1.900000	0.070000	7.000000	21.000000
50%	7.900000	0.520000	0.250000	2.200000	0.079000	13.000000	37.000000
75%	9.100000	0.640000	0.420000	2.600000	0.090000	21.000000	61.000000
max	15.900000	1.580000	1.000000	15.500000	0.611000	68.000000	289.000000



b) Generate summary statistics for categorical columns

In [ ]:

```
# Calculate the frequency counts for each quality value
quality_counts = df['quality'].value_counts()

# Calculate the percentage distribution of quality values
quality_percentage = (quality_counts / len(df)) * 100

# Display the results
print("Frequency counts of quality values:")
print(quality_counts)
print("\nPercentage distribution of quality values:")
print(quality_percentage)
```

Frequency counts of quality values:

```
quality
5      483
6      462
7      143
4       33
8       16
3        6
```

Name: count, dtype: int64

Percentage distribution of quality values:

```
quality
5      42.257218
6      40.419948
7      12.510936
4       2.887139
8       1.399825
3       0.524934
```

Name: count, dtype: float64

c) Explore the relationships between variables using appropriate EDA techniques such as correlation analysis or cross-tabulation.

```
In [ ]: # Correlation matrix
correlation_matrix = df.corr()
print(correlation_matrix)
```

	fixed acidity	volatile acidity	citric acid \	
fixed acidity	1.000000	-0.250728	0.673157	
volatile acidity	-0.250728	1.000000	-0.544187	
citric acid	0.673157	-0.544187	1.000000	
residual sugar	0.171831	-0.005751	0.175815	
chlorides	0.107889	0.056336	0.245312	
free sulfur dioxide	-0.164831	-0.001962	-0.057589	
total sulfur dioxide	-0.110628	0.077748	0.036871	
density	0.681501	0.016512	0.375243	
pH	-0.685163	0.221492	-0.546339	
sulphates	0.174592	-0.276079	0.331232	
alcohol	-0.075055	-0.203909	0.106250	
quality	0.121970	-0.407394	0.240821	

	residual sugar	chlorides	free sulfur dioxide \	
fixed acidity	0.171831	0.107889	-0.164831	
volatile acidity	-0.005751	0.056336	-0.001962	
citric acid	0.175815	0.245312	-0.057589	
residual sugar	1.000000	0.070863	0.165339	
chlorides	0.070863	1.000000	0.015280	
free sulfur dioxide	0.165339	0.015280	1.000000	
total sulfur dioxide	0.190790	0.048163	0.661093	
density	0.380147	0.208901	-0.054150	
pH	-0.116959	-0.277759	0.072804	
sulphates	0.017475	0.374784	0.034445	
alcohol	0.058421	-0.229917	-0.047095	
quality	0.022002	-0.124085	-0.063260	

	total sulfur dioxide	density	pH	sulphates	\
fixed acidity	-0.110628	0.681501	-0.685163	0.174592	
volatile acidity	0.077748	0.016512	0.221492	-0.276079	
citric acid	0.036871	0.375243	-0.546339	0.331232	
residual sugar	0.190790	0.380147	-0.116959	0.017475	
chlorides	0.048163	0.208901	-0.277759	0.374784	
free sulfur dioxide	0.661093	-0.054150	0.072804	0.034445	
total sulfur dioxide	1.000000	0.050175	-0.059126	0.026894	
density	0.050175	1.000000	-0.352775	0.143139	
pH	-0.059126	-0.352775	1.000000	-0.185499	
sulphates	0.026894	0.143139	-0.185499	1.000000	
alcohol	-0.188165	-0.494727	0.225322	0.094421	
quality	-0.183339	-0.175208	-0.052453	0.257710	

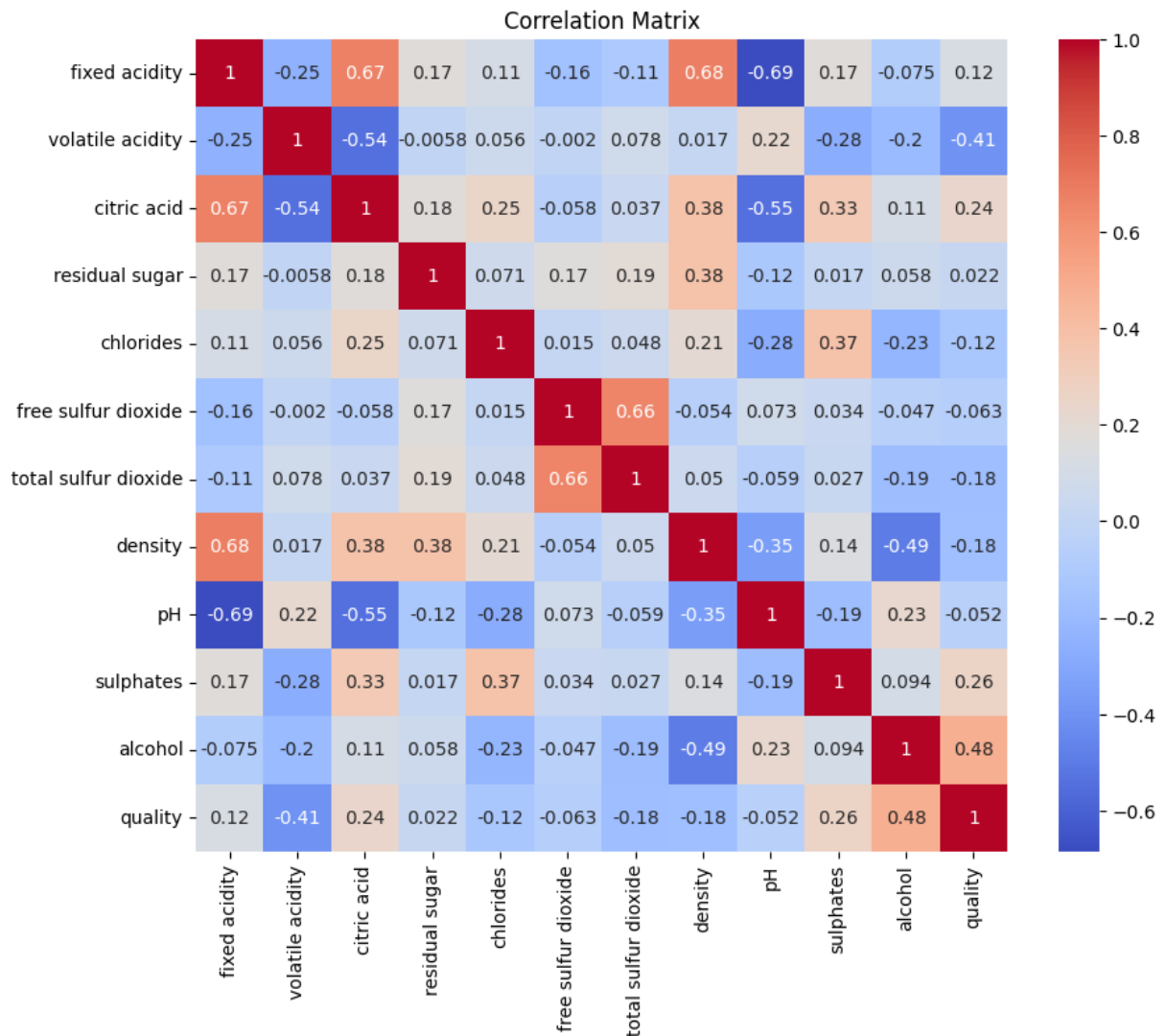
  

	alcohol	quality
fixed acidity	-0.075055	0.121970
volatile acidity	-0.203909	-0.407394
citric acid	0.106250	0.240821
residual sugar	0.058421	0.022002
chlorides	-0.229917	-0.124085
free sulfur dioxide	-0.047095	-0.063260
total sulfur dioxide	-0.188165	-0.183339
density	-0.494727	-0.175208
pH	0.225322	-0.052453
sulphates	0.094421	0.257710
alcohol	1.000000	0.484866
quality	0.484866	1.000000

## Heatmap of correlation matrix

```
In [ ]: correlation = df.corr()

# Generate a heatmap using seaborn
plt.figure(figsize=(10, 8))
sns.heatmap(correlation, annot=True, cmap='coolwarm', square=True)
plt.title('Correlation Matrix')
plt.show()
```



Some colinearities can be found. PCA could be usefull when creating a machine learning model.

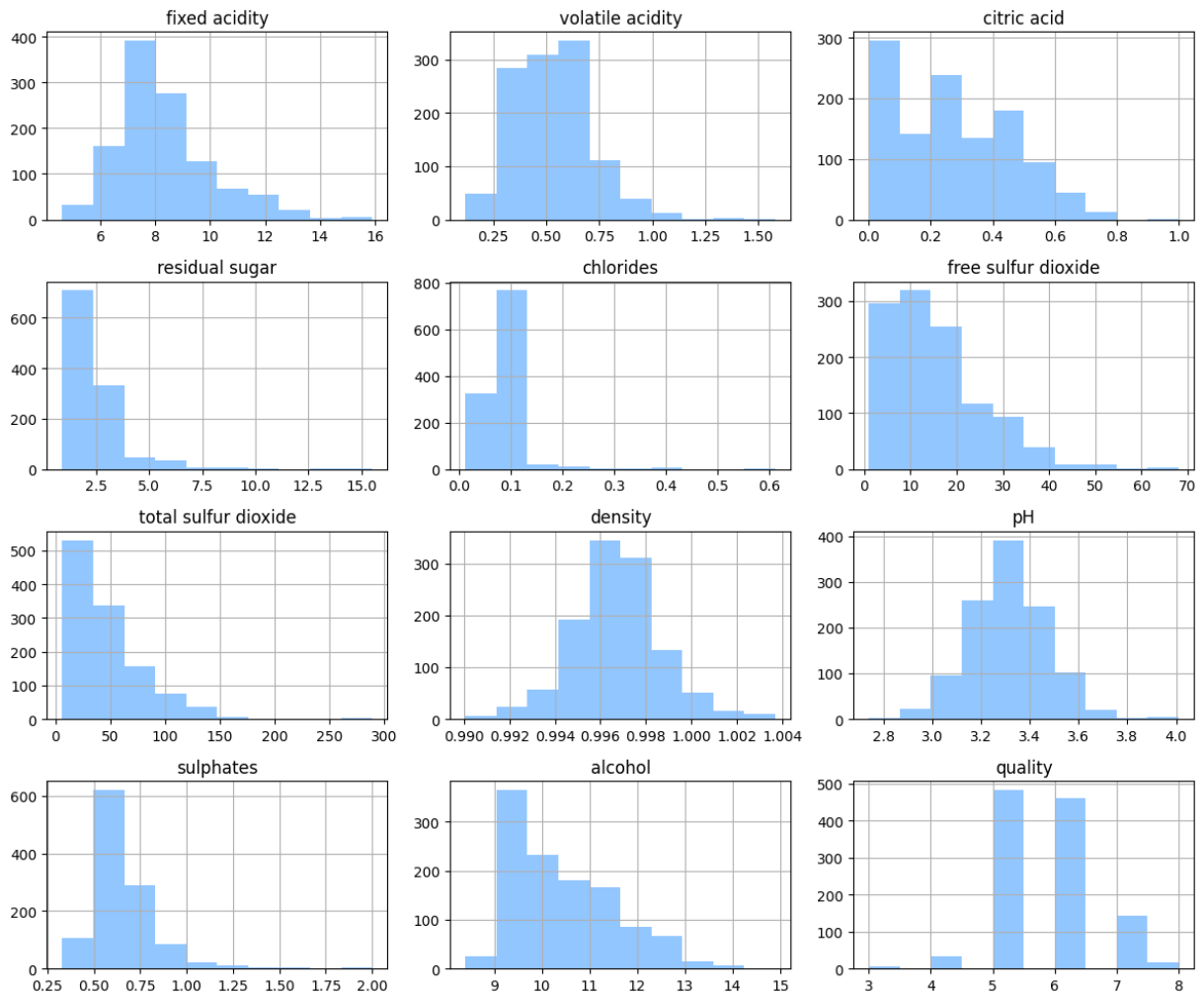
The quality range ranges between 3 and 8. No unrealistic values detectable. <\html>

## Visualisation

# Create meaningful visualizations to understand the dataset better. Include at least three different types of visualization

a) A histogram to visualize the distribution of a numerical variable.

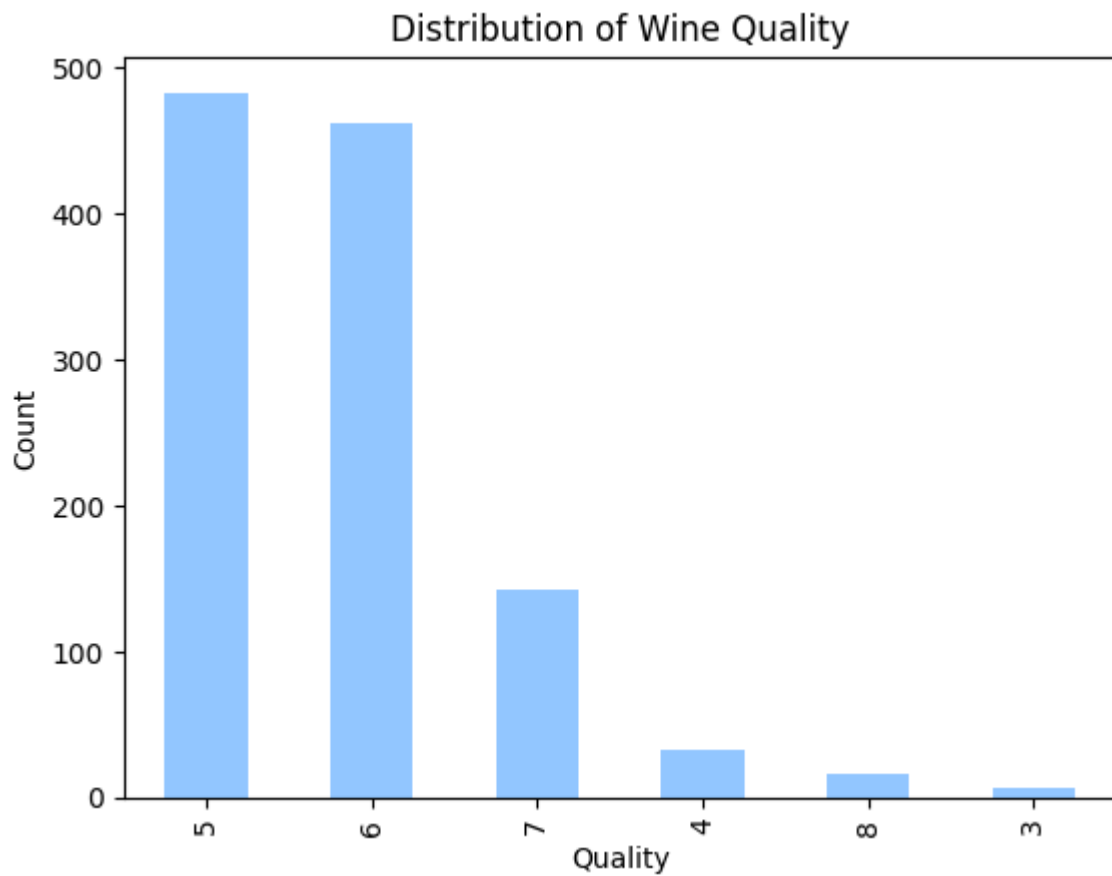
```
In [ ]: # Histograms
df.hist(figsize=(12, 10))
plt.tight_layout()
plt.show()
```



Bar plot for 'quality' column

```
In [ ]: df['quality'].value_counts().plot(kind='bar')
plt.xlabel('Quality')
plt.ylabel('Count')
plt.title('Distribution of Wine Quality')
plt.show()
```



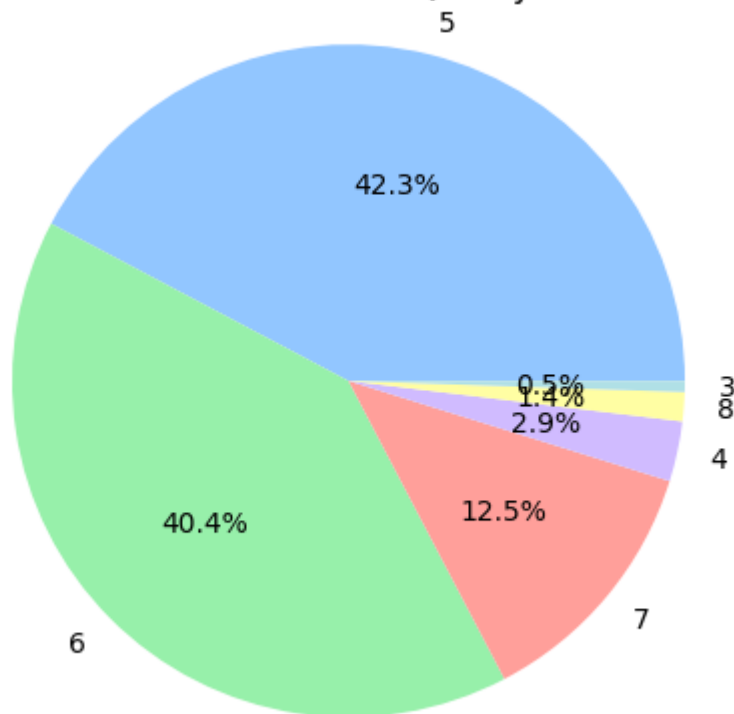


Distribution of Quality

```
In [ ]: value_counts = df['quality'].value_counts()

# Plot the distribution of the "quality" column as a pie chart
plt.pie(value_counts, labels=value_counts.index, autopct='%1.1f%%')
plt.title('Distribution of Quality')
plt.axis('equal') # Ensure pie is drawn as a circle
plt.show()
```

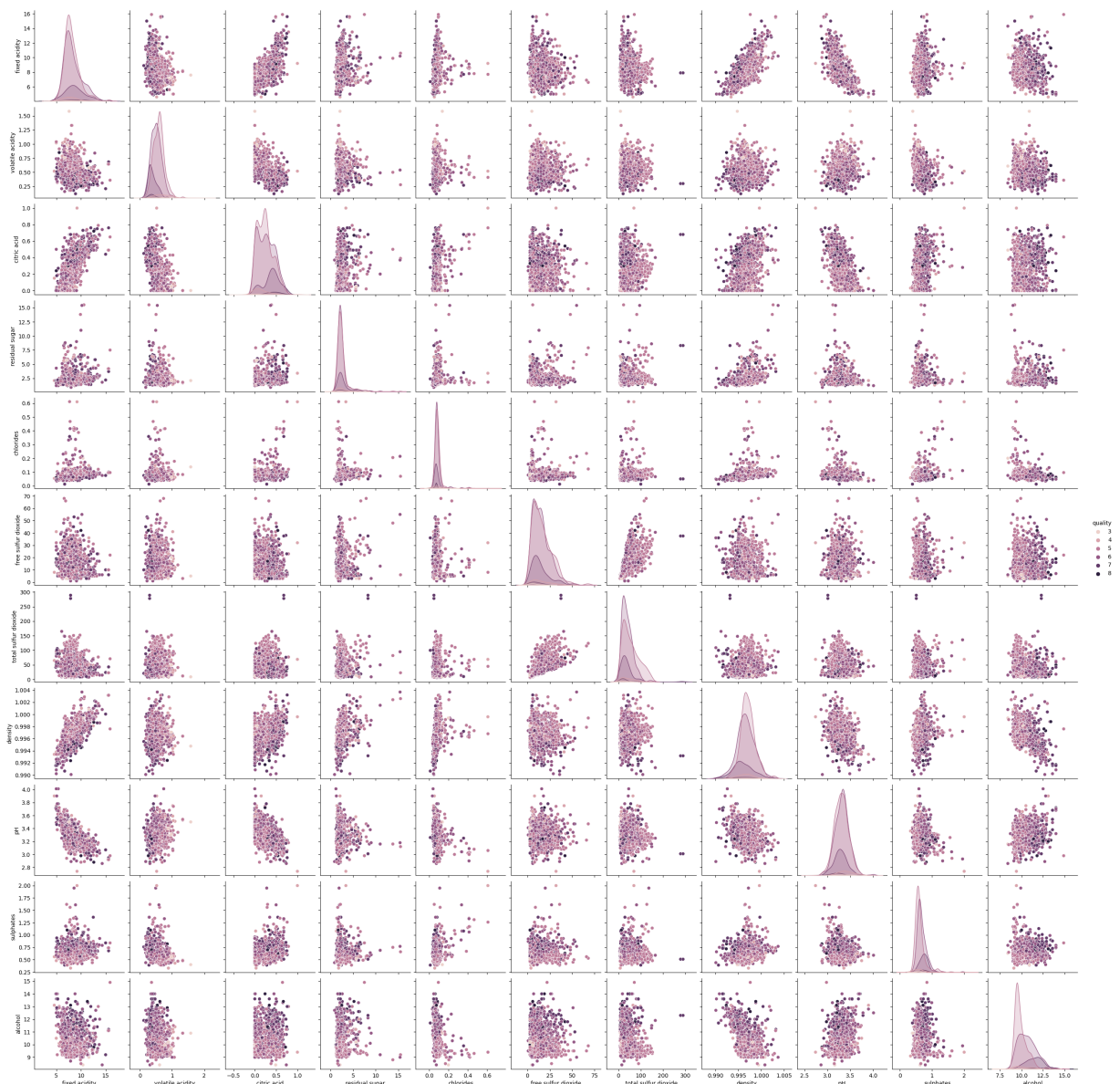
Distribution of Quality



Pairplot

```
In [ ]: sns.pairplot(df, hue='quality')
```

```
Out[ ]: <seaborn.axisgrid.PairGrid at 0x2d5cf7f7c90>
```

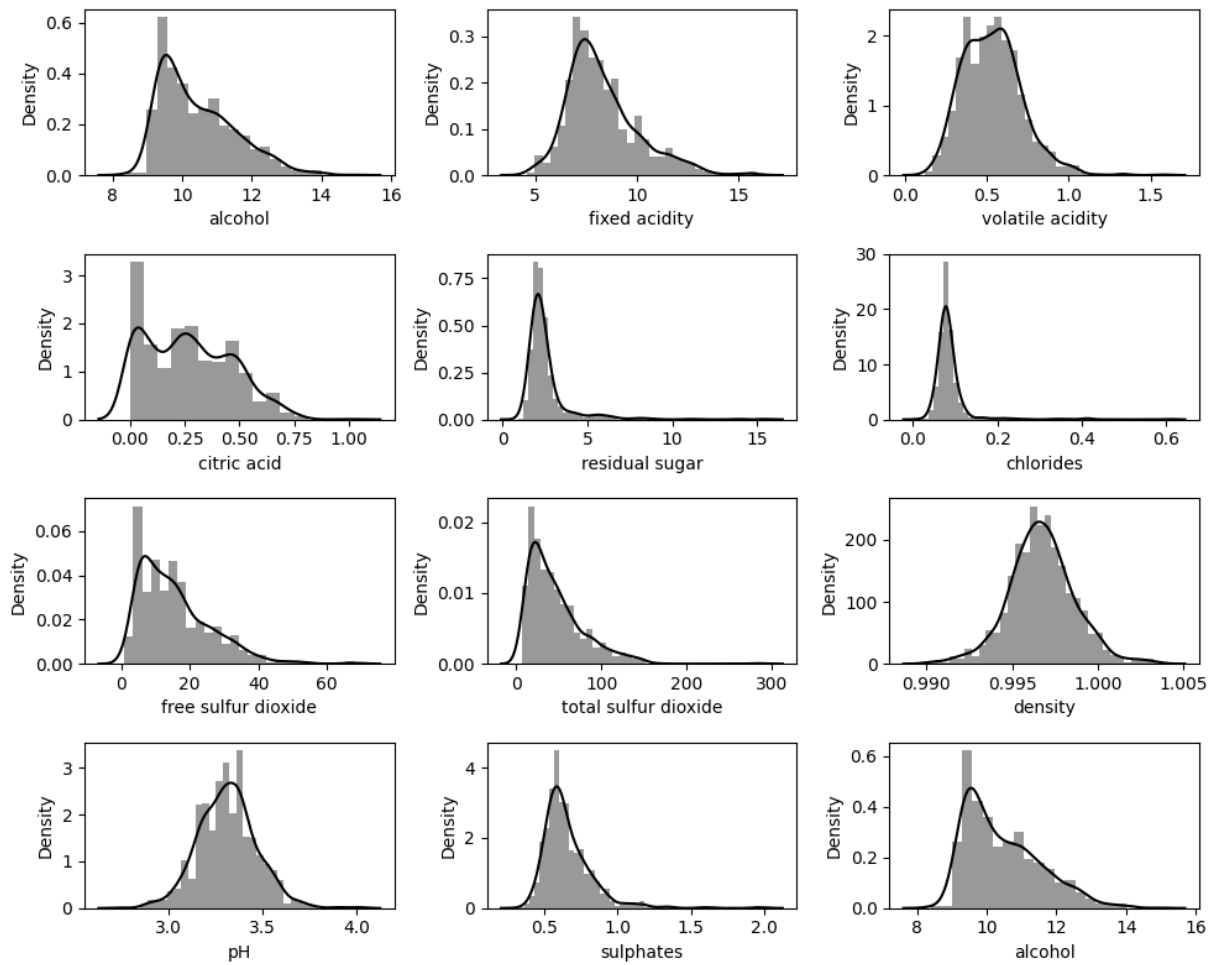


Some correlations can be observed, which can be looked at in more detail in a heatmap. For quality column (target), no strong linear trend can be observed for single attributes.

```
In [ ]: fig, axes = plt.subplots(4,3,figsize=(10,8))

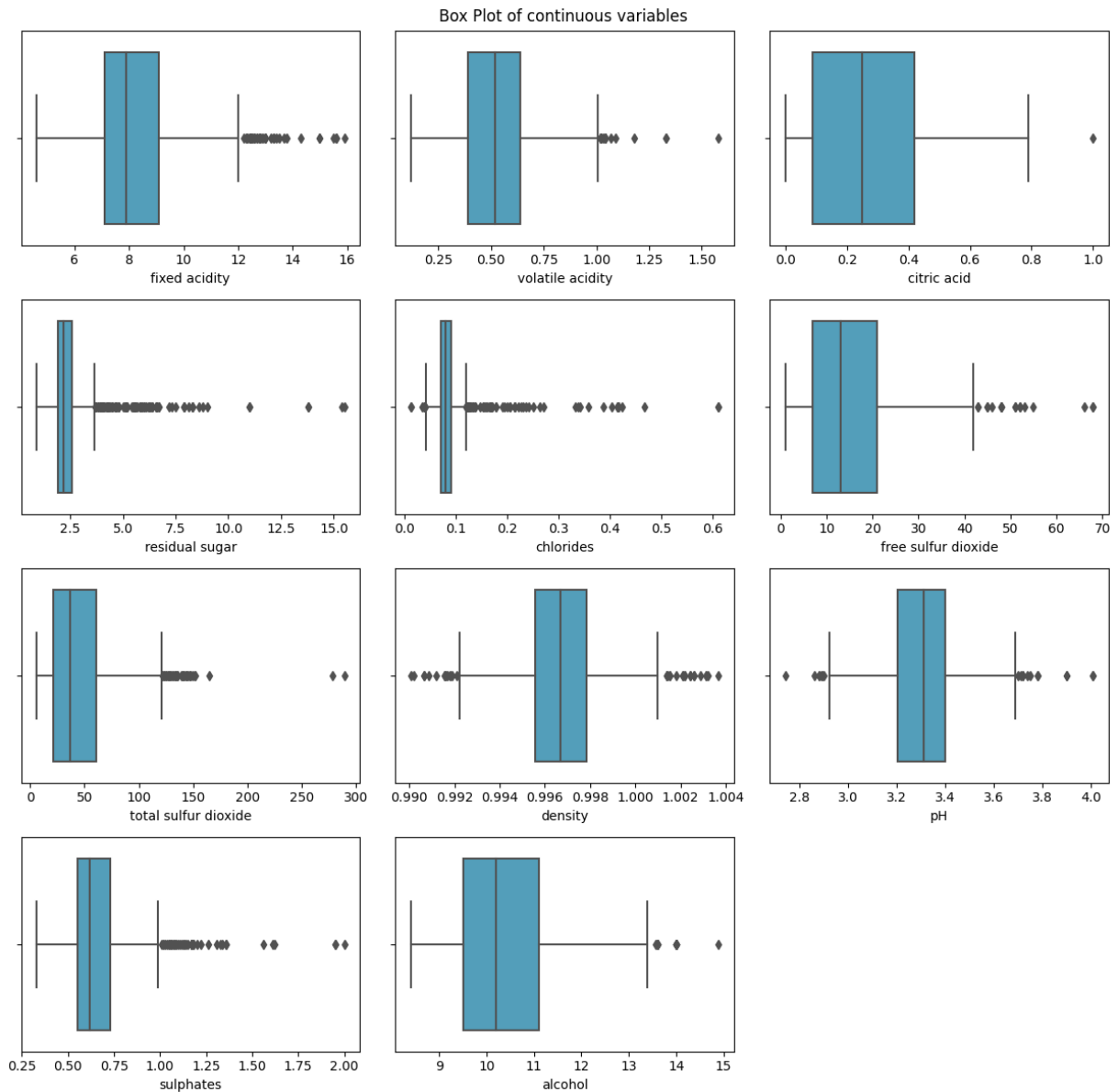
i=-1
for row in range(4):
    for col in range(3):
        if i<11:
            sns.distplot(df[df.columns[:-1][i]], ax=axes[row][col], color='black')
            i+=1

fig.tight_layout()
```



Box plot for each continuous variable

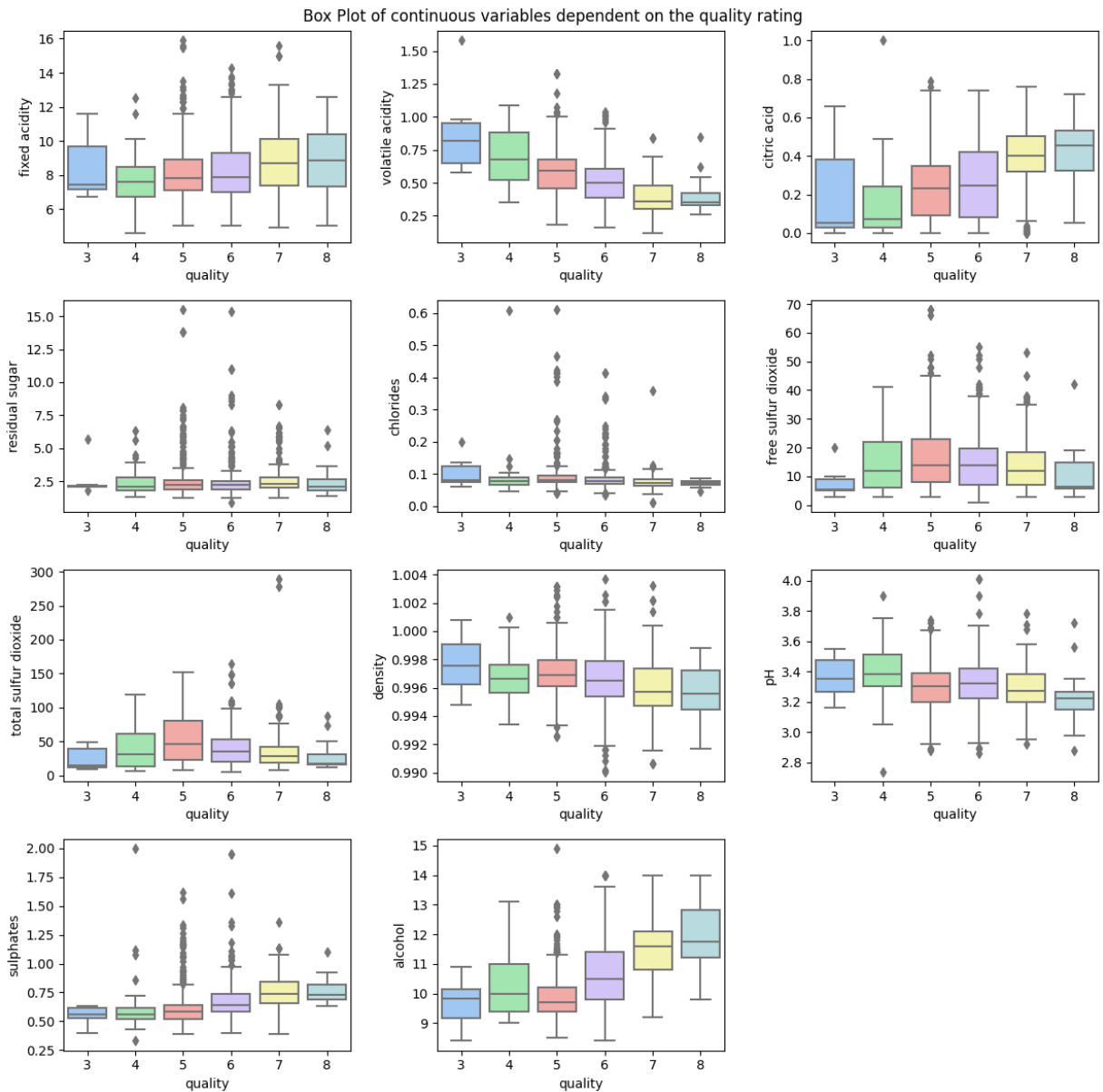
```
In [ ]: plt.figure(figsize=(12,12))
for i, col in enumerate(df.select_dtypes(include=['float64']).columns):
    ax = plt.subplot(4,3, i+1)
    sns.boxplot(data=df, x=col, ax=ax,palette="GnBu_d")
plt.suptitle('Box Plot of continuous variables')
plt.tight_layout()
```



Several outliers can be detected in the dataset using boxplots. However, values don't seem to be unusual. For machine learning models, I would keep them in the first place.

Box plot for each continuous variable in dependence of target column

```
In [ ]: plt.figure(figsize=(12,12))
for i, col in enumerate(df.select_dtypes(include=['float64']).columns):
    ax = plt.subplot(4,3, i+1)
    sns.boxplot(data=df, x='quality', y=col, ax=ax)
plt.suptitle('Box Plot of continuous variables dependent on the quality rating')
plt.tight_layout()
```



We get several insights of this figure:

1. Median of fixed acidity increases with increasing quality rating, whereas volatile acidity decreases.
2. median of the amount of citric acid increases with increasing quality rating.
3. residual sugar, chlorides and density seem to have little effect on quality rating.
4. low and high rated wines seem to be low in free sulfur dioxide and total sulfur dioxide.
5. better rated wines seem to have a lower pH.
6. wines with higher ratings seem to be higher in the amount of sulphates and alcohol.

## Conclusion

### Report - Wine Data

**Introduction:** This report focuses on the analysis of a dataset related to wine, including attributes such as fixed acidity, volatile acidity, citric acid, residual sugar, chloride, free sulfur

dioxide, density, pH, sulphates, alcohol, quality, and several other attributes. Below is a summary of the findings after data wrangling, exploratory data analysis (EDA), and visualizations.

### **1. Data Inspection and Preprocessing:**

- The raw data appeared to be clean, with no significant missing values or obvious errors in the given columns.
- During preprocessing, there was little need for data cleaning.

### **2. Data Distribution:**

- The dataset contains both continuous and categorical attributes, with 'quality' as the target variable. 'Quality' is a categorical variable representing the quality of wine on a scale from 3 to 9.
- The distribution of wine quality shows signs of skewness, with a high concentration of wines having quality around 5 and 6.

### **3. Data Visualization:**

- Visualizations such as frequency plots, box plots, and scatter plots were created to explore the distribution and relationships between attributes and wine quality.
- Notable findings include a potential relationship between alcohol content and wine quality, as well as the influence of certain chemical attributes (e.g., volatile acidity, citric acid, and sulphates) on wine quality.

### **4. Feature Engineering Techniques:**

- Additional feature engineering may be required to better understand the relationships between attributes and wine quality. For example, creating new features or normalizing data for modeling purposes.

### **5. Modeling:**

- After completing data preprocessing and exploratory data analysis, the next steps typically involve building predictive models to understand the key factors influencing wine quality.

### **6. Conclusion:**

This project has provided a foundation for in-depth analysis and modeling, allowing you to identify the key factors that influence wine quality and make predictions based on this information. Subsequent steps may include building predictive models to provide quality predictions based on chemical and mechanical attributes.

This dataset shows the rating of >1000 different wines and their chemical parameters. The dataset is unbalanced regarding the different quality ratings. >80 % of the wines get a rating of "5" or "6" which can be translated as average rated wines.

There are no missing values within the dataset and no categorical columns.

Colinearity between different features can be observed. For machine learning models, a PCA could be helpful.

There are some outliers within the values. However, the statistical analysis does not show unrealistic values, so for machine learning, I would prefer to not delete or manipulate them in the first place.

When we look at the different features and their impact on the wine rating, different trends, as well as apparently little effect can be observed(see chapter before). When building a machine learning model, alcohol and volatile acidity will probably have the highest feature importance as the trend can be clearly seen.