

model-summary-and-eda-1

February 6, 2024

LOADING DATA FROM GITHUB INTO SINGLESTORE NOTEBOOK

```
[4272]: import pandas as pd

github_url = 'https://raw.githubusercontent.com/aniruddhachoudhury/
↳Red-Wine-Quality/master/winequality-red.csv'
wine_data = pd.read_csv(github_url)
print('Wine Quality Dataset:')
print(wine_data.head())
```

Wine Quality Dataset:

	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	

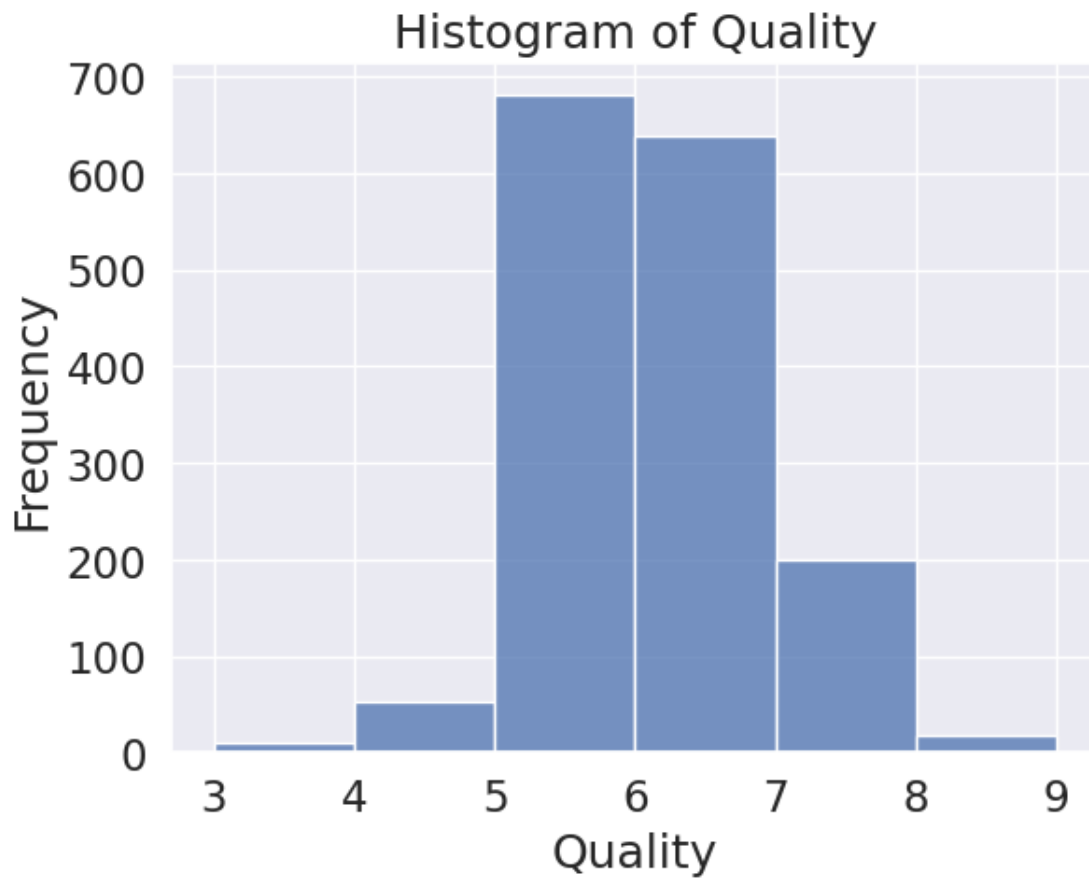
	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
0	11.0	34.0	0.9978	3.51	0.56	
1	25.0	67.0	0.9968	3.20	0.68	
2	15.0	54.0	0.9970	3.26	0.65	
3	17.0	60.0	0.9980	3.16	0.58	
4	11.0	34.0	0.9978	3.51	0.56	

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5

```
[4273]: import seaborn as sns
import matplotlib.pyplot as plt

sns.histplot(wine_data['quality'], bins=range(3, 10), kde=False)
plt.xlabel('Quality')
plt.ylabel('Frequency')
```

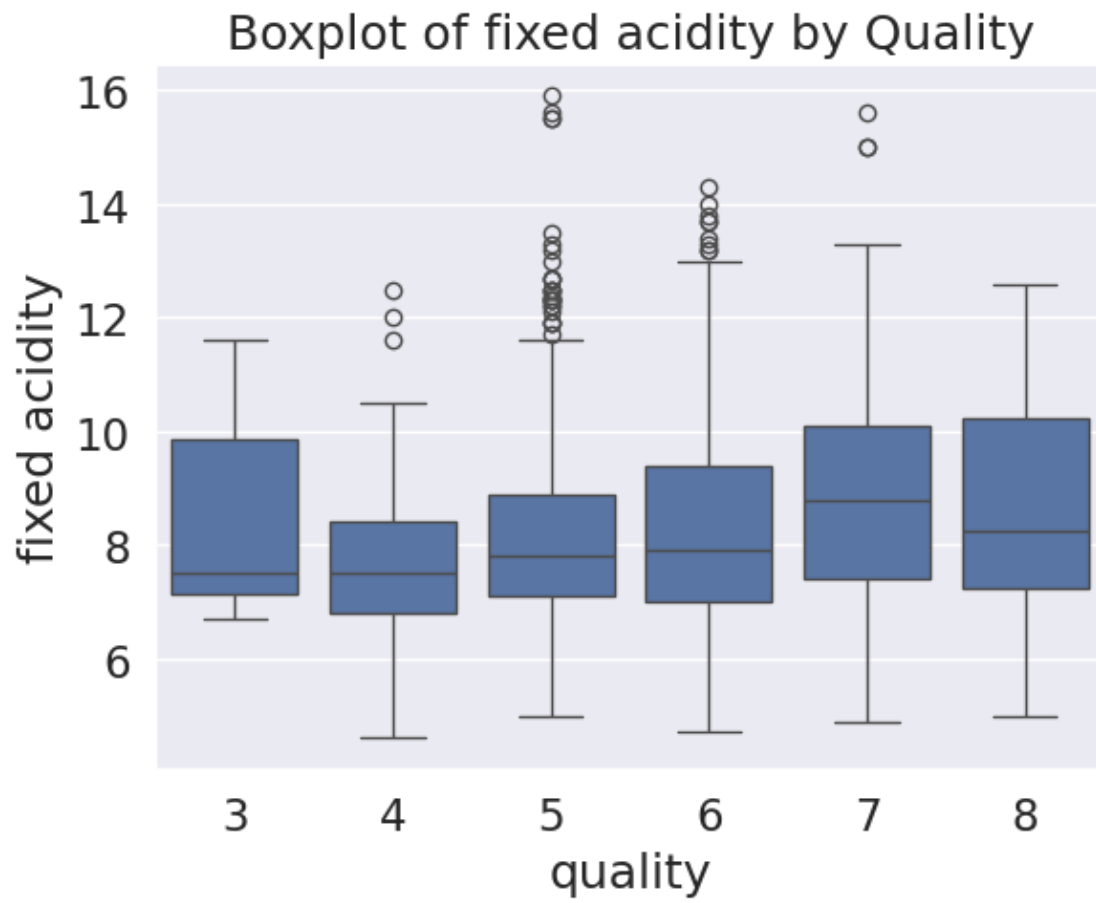
```
plt.title('Histogram of Quality')
plt.show()
```

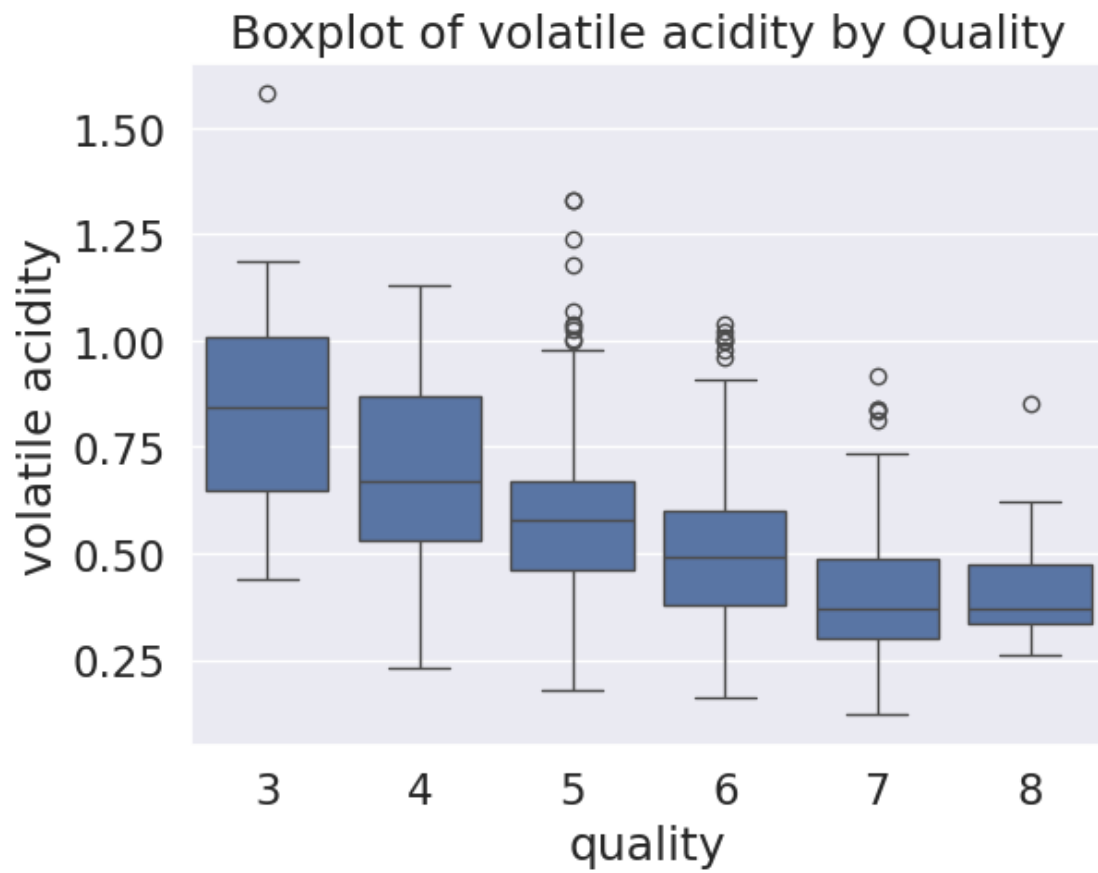


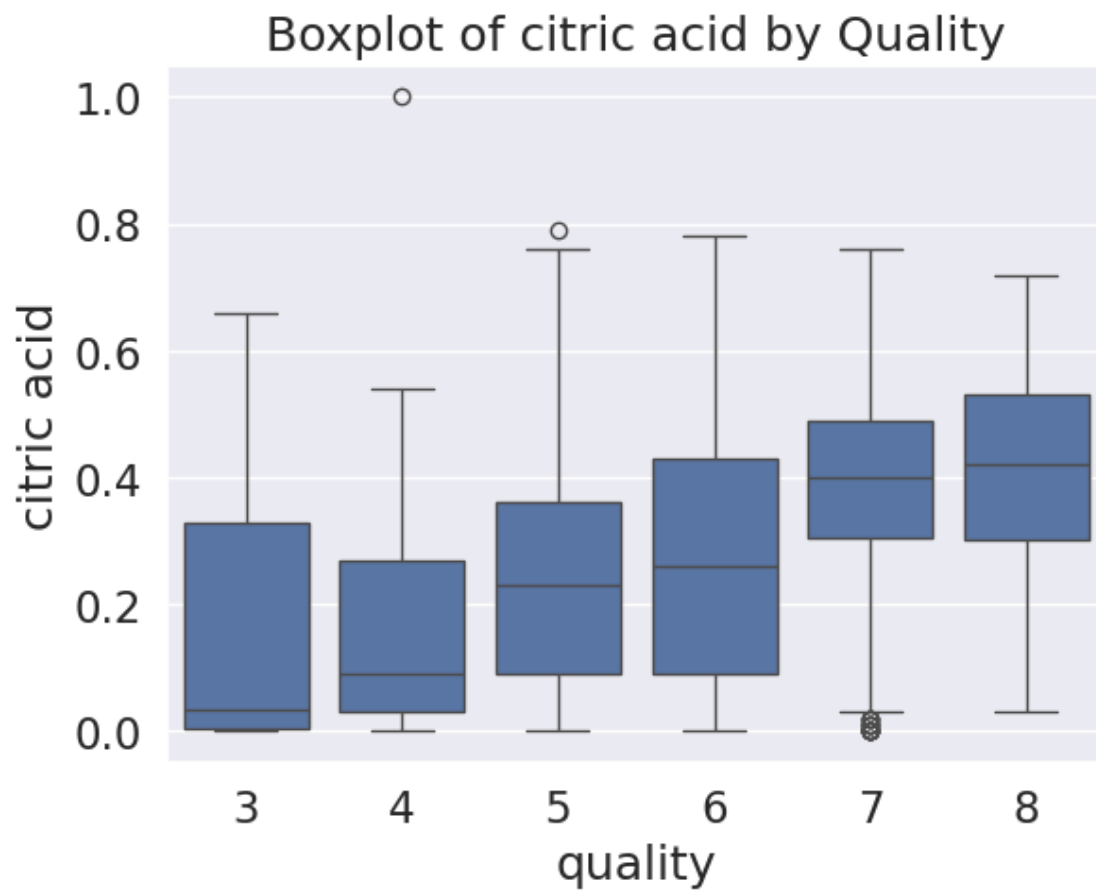
```
[4274]: import seaborn as sns
import matplotlib.pyplot as plt

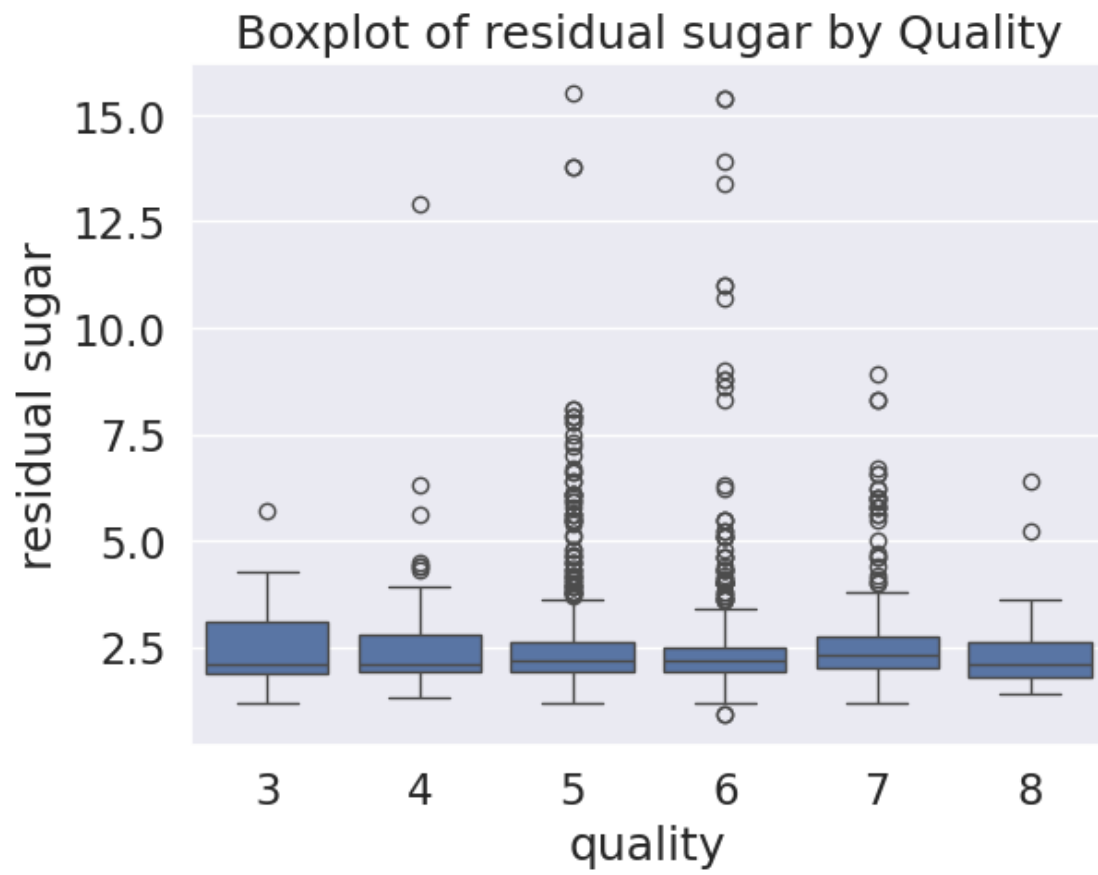
predictors = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual_
↪sugar', 'chlorides', 'free sulfur dioxide', 'total sulfur dioxide',
↪'density', 'pH', 'sulphates', 'alcohol']

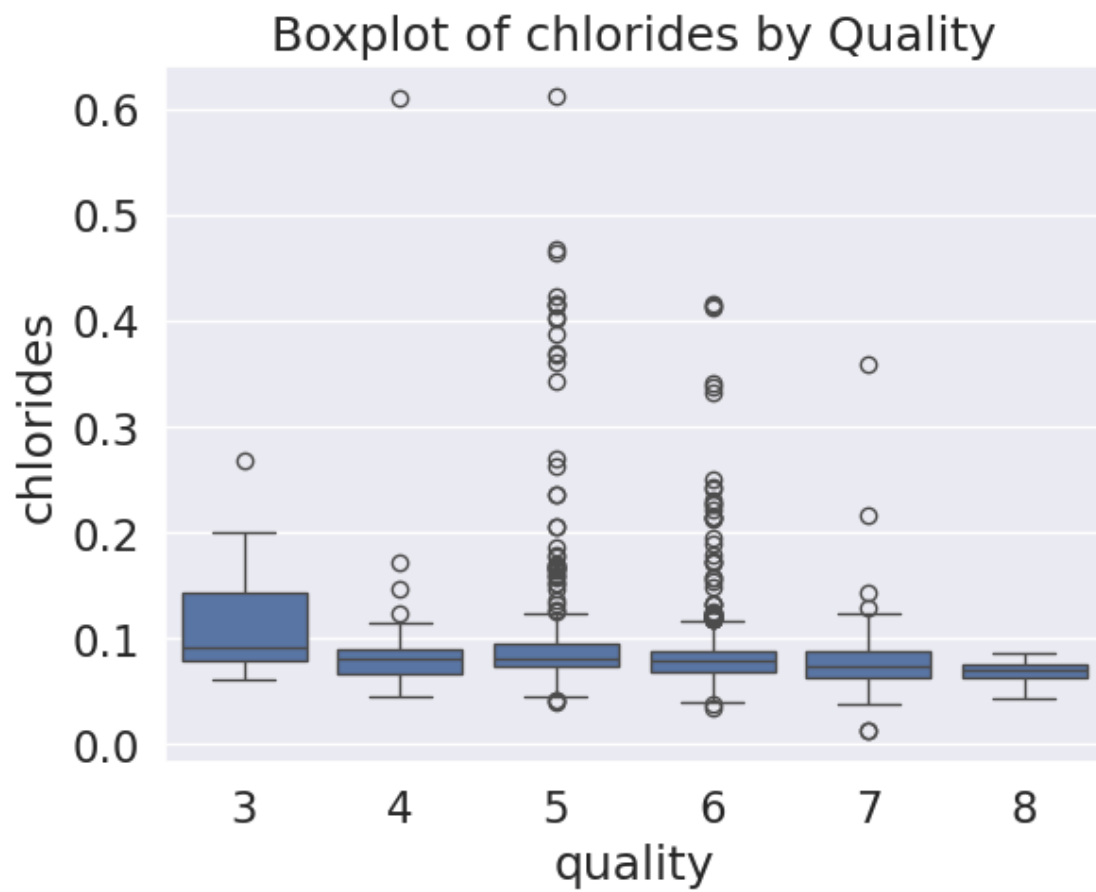
for predictor in predictors:
    sns.boxplot(x='quality', y=predictor, data=wine_data)
    plt.title(f'Boxplot of {predictor} by Quality')
    plt.show()
```

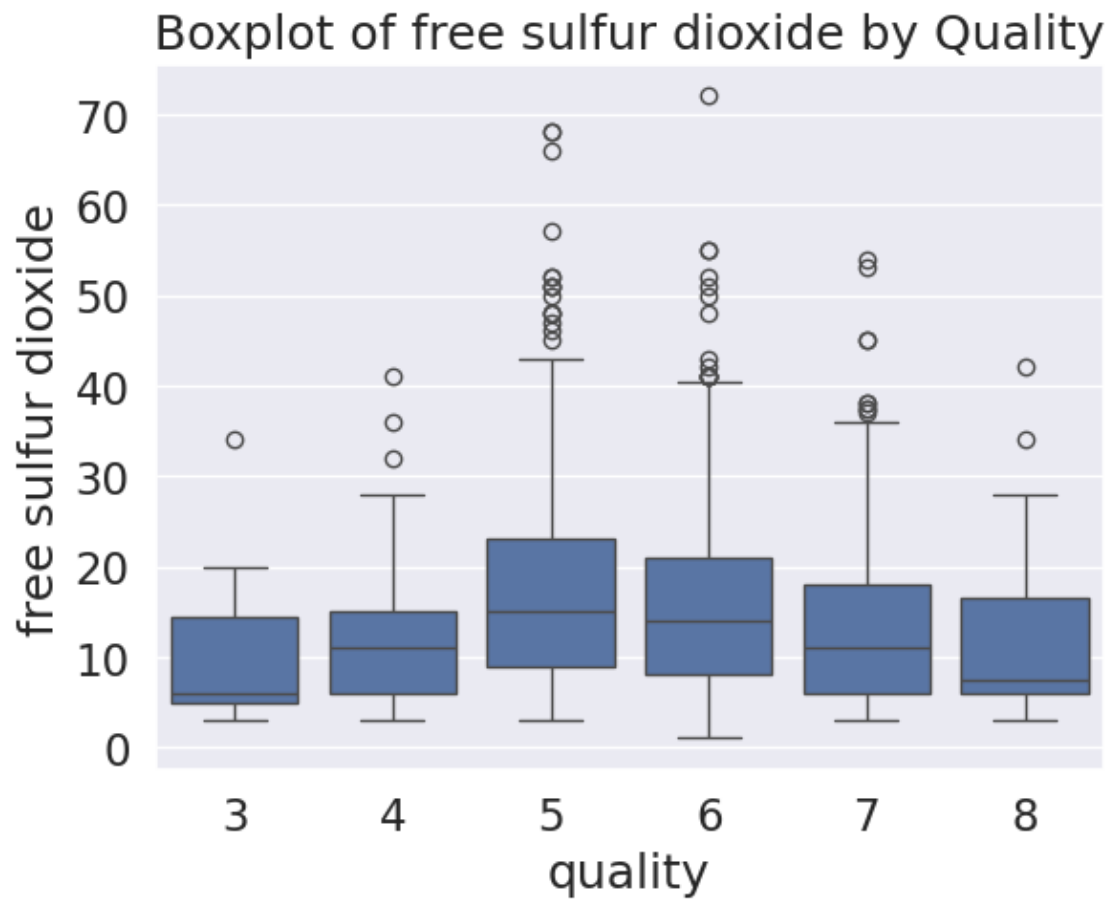


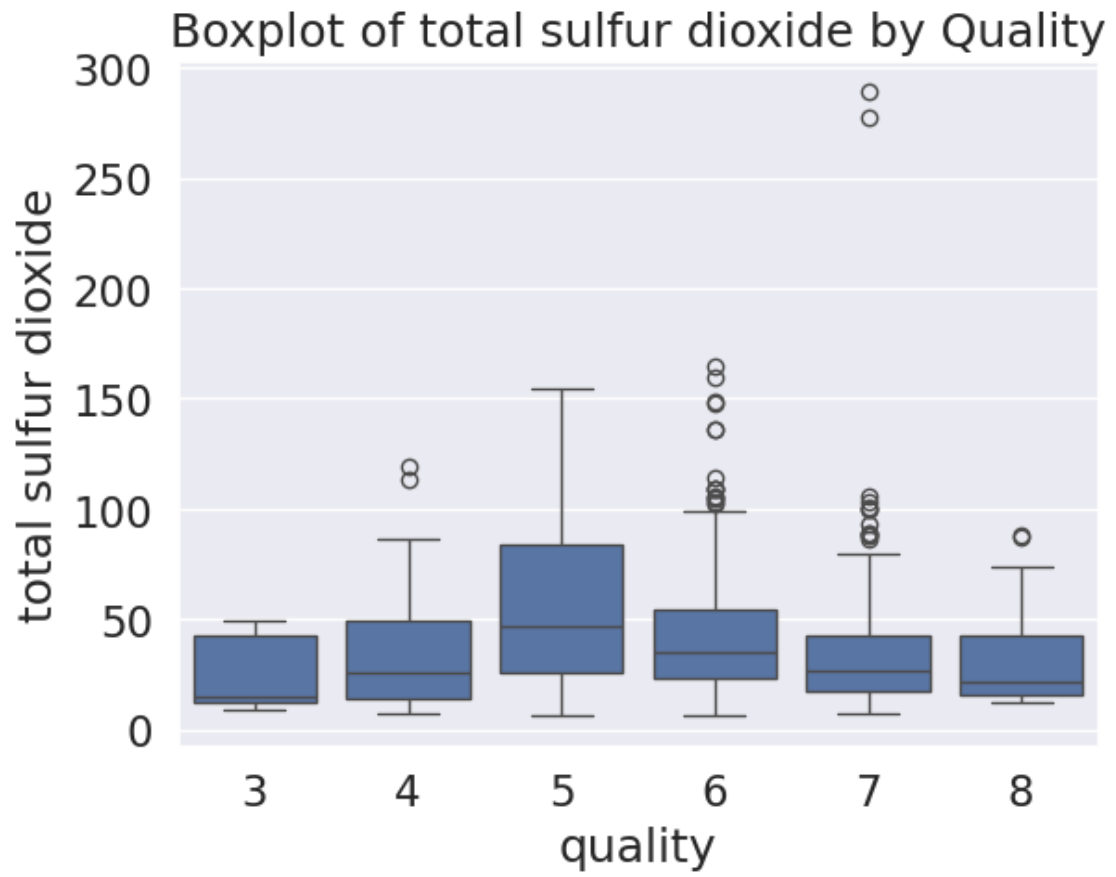


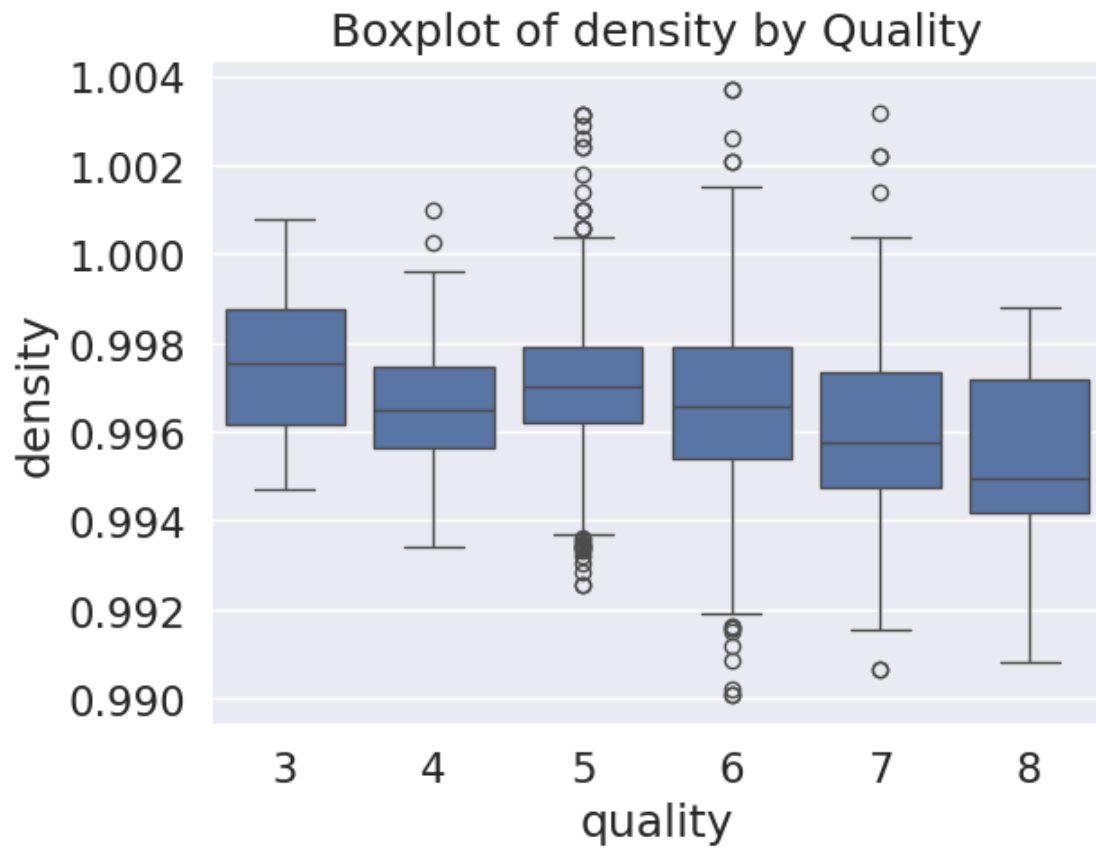


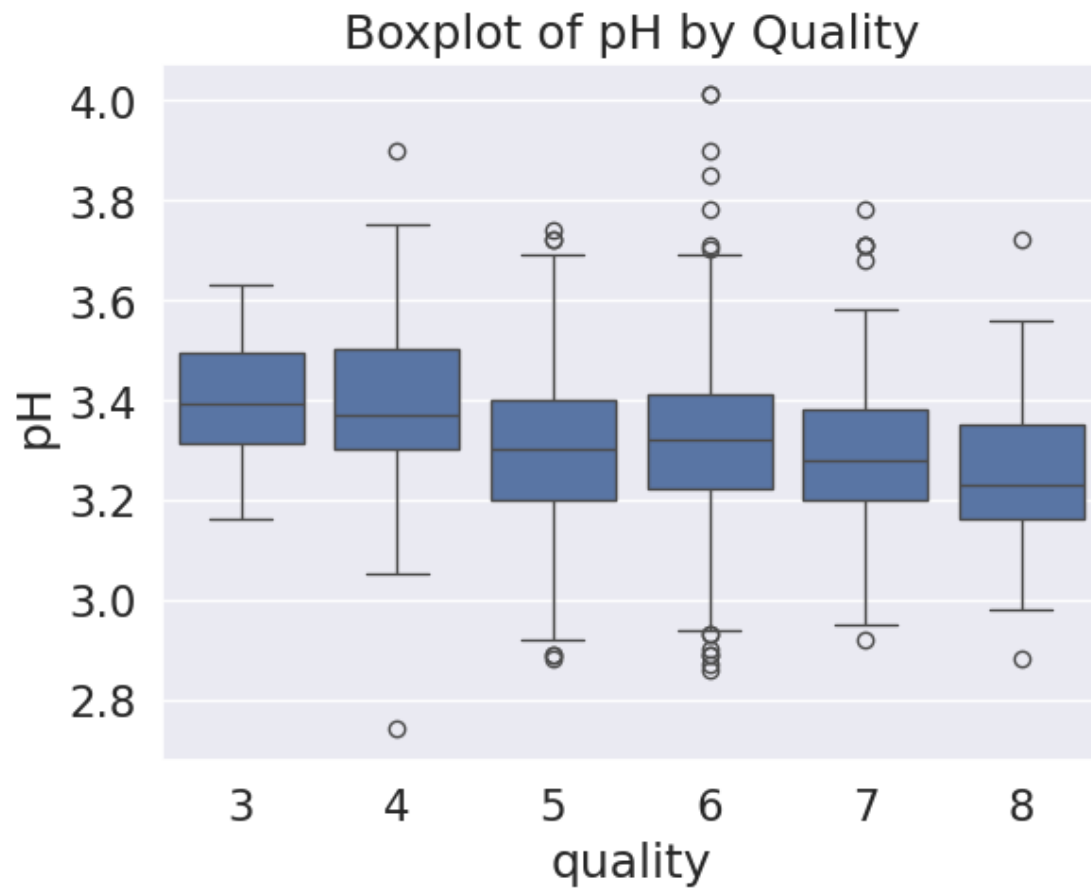


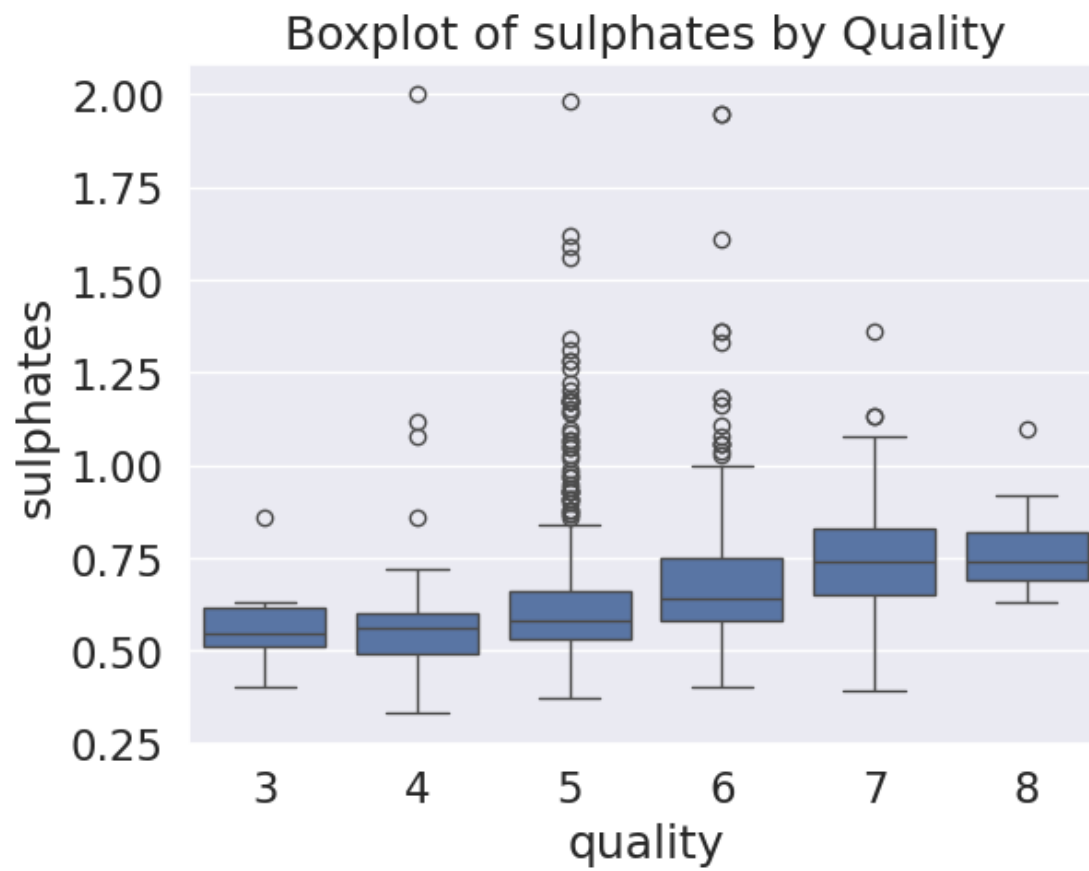


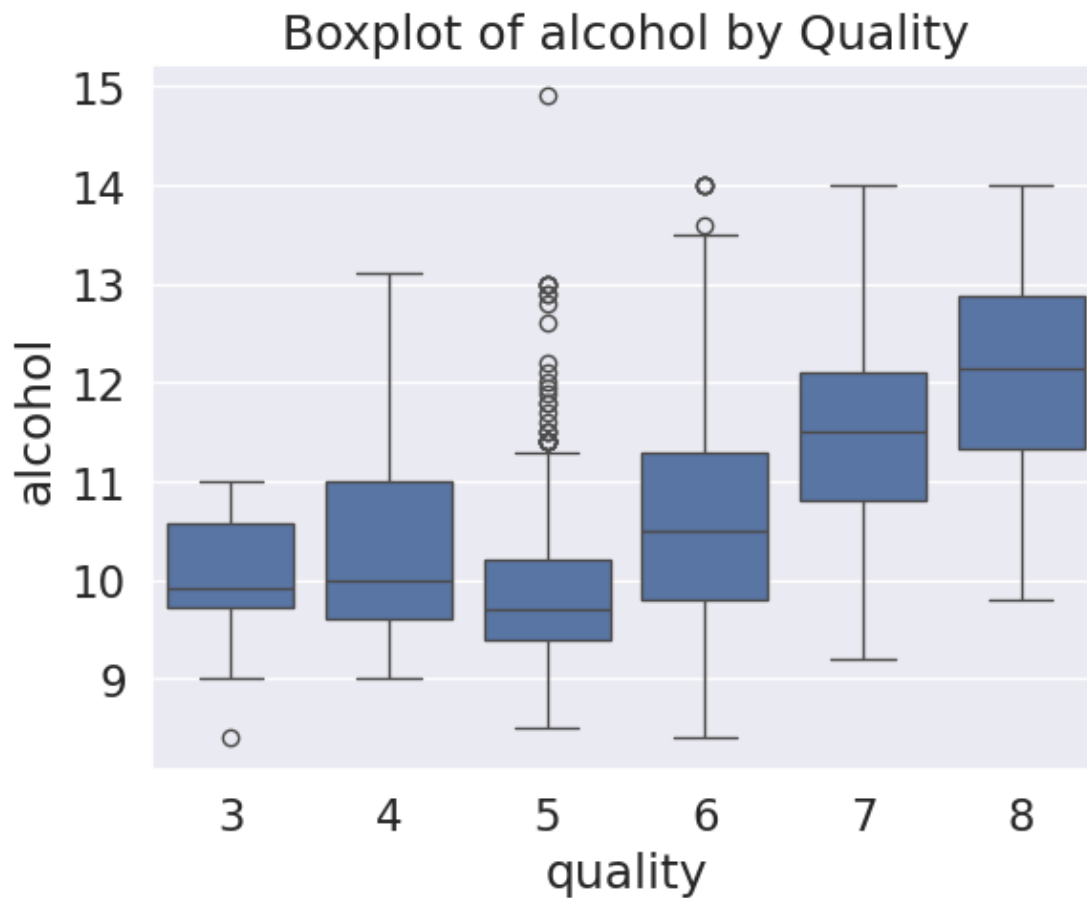








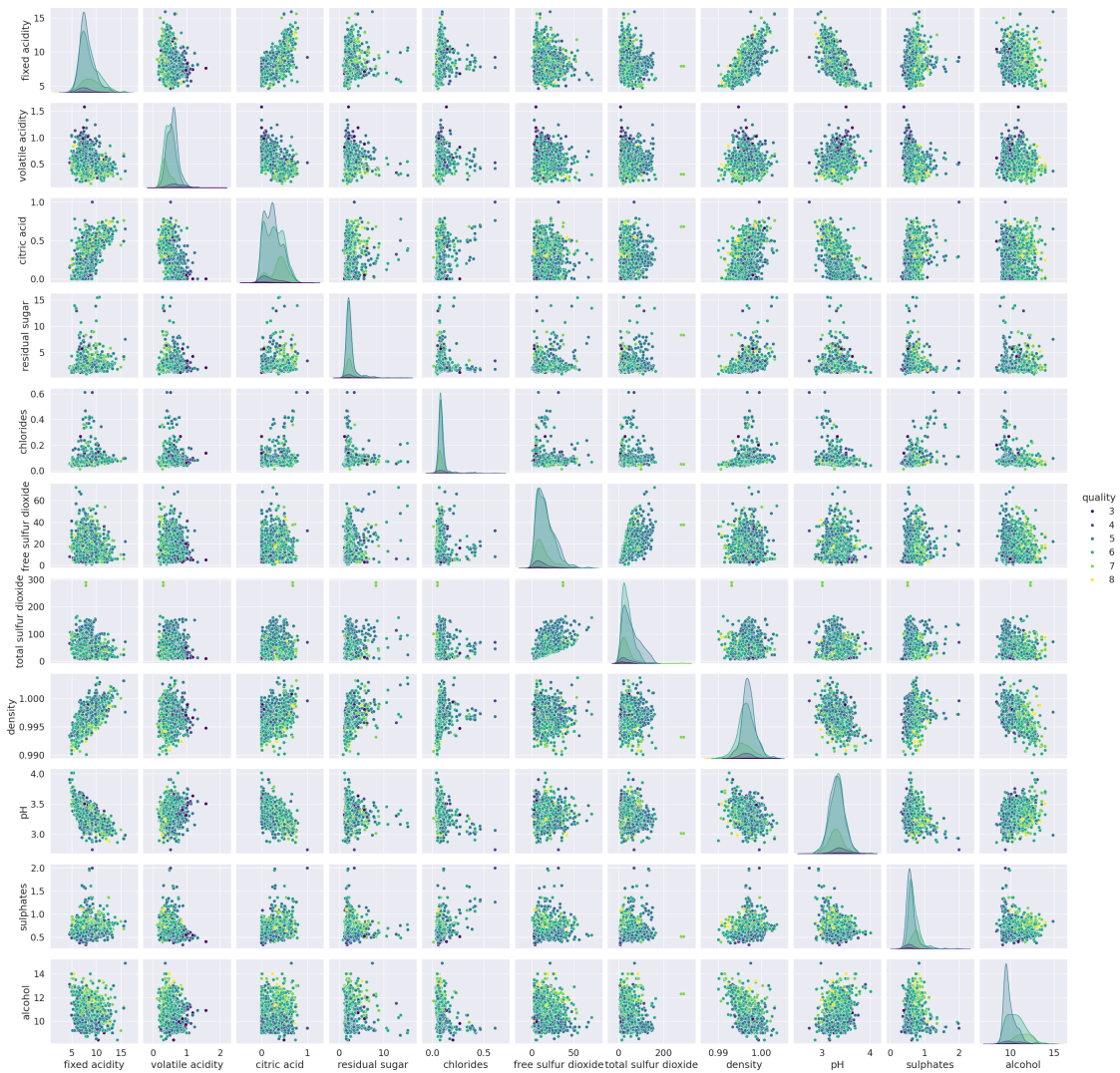




```
[4275]: import seaborn as sns

selected_predictors = ['fixed acidity', 'volatile acidity', 'citric acid', '
    ↪ 'residual sugar', 'chlorides', 'free sulfur dioxide', 'total sulfur
    ↪ dioxide', 'density', 'pH', 'sulphates', 'alcohol', 'quality']

sns.pairplot(wine_data[selected_predictors], hue='quality', palette='viridis')
plt.show()
```



```
[4276]: import plotly.express as px
import pickle
```

```
[4277]: plt.figure(figsize=(18, 8))
sns.heatmap(wine_data.corr(), vmin=-1, vmax=1, annot=True, cmap='BrBG')
plt.title('Correlation Map Of Red Wine Quality', fontdict={'fontsize':12},
          pad=12);
```

