

# Overview

## Input Variables:

**fixed acidity:** most acids involved with wine or fixed or nonvolatile

**volatile acidity:** the amount of acetic acid in wine

**citric acid:** found in small quantities, citric acid can add 'freshness' and flavor to wines

**residual sugar:** the amount of sugar remaining after fermentation stops

**chlorides:** the amount of salt in the wine

**free sulfur dioxide:** the free form of SO<sub>2</sub> exists in equilibrium between molecular SO<sub>2</sub> (as a dissolved gas) and bisulfite ion

**total sulfur dioxide:** amount of free and bound forms of SO<sub>2</sub>

**density:** the density of water is close to that of water depending on the percent alcohol and sugar content

**pH:** describes how acidic or basic a wine is on a scale from 0 (very acidic) to 14 (very basic)

**sulphates:** a wine additive which can contribute to sulfur dioxide gas (SO<sub>2</sub>) levels

**alcohol:** the percent alcohol content of the wine

## Output Variable:

**quality:** output variable (based on sensory data, score between 0 and 10)

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.metrics import accuracy_score
from sklearn.metrics import mean_squared_error
from sklearn.model_selection import cross_val_score
from sklearn.metrics import confusion_matrix
from collections import Counter
from IPython.core.display import display, HTML
sns.set_style('darkgrid')
import warnings
warnings.filterwarnings('ignore')
```

```
In [2]: wine = pd.read_csv('Red_wine.csv')
```

```
In [3]: wine.head()
```

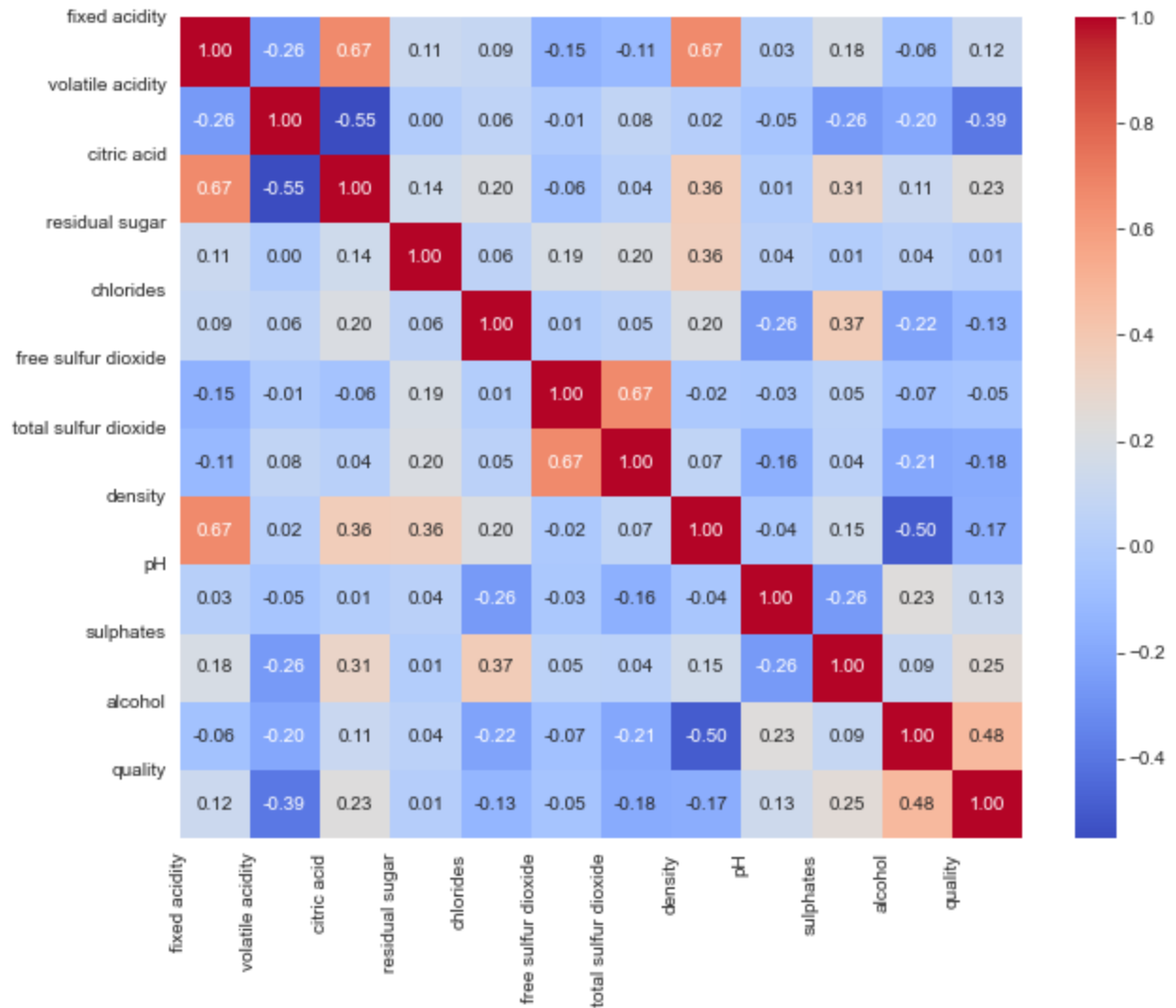
```
Out[3]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4	5.0

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	9.8	5.0
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	9.8	5.0
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	9.8	6.0
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4	5.0

In [4]:

```
corr = wine.corr()
#Plot figsize
fig, ax = plt.subplots(figsize=(10, 8))
#Generate Heat Map, allow annotations and place floats in map
sns.heatmap(corr, cmap='coolwarm', annot=True, fmt=".2f")
#Apply xticks
plt.xticks(range(len(corr.columns)), corr.columns);
#Apply yticks
plt.yticks(range(len(corr.columns)), corr.columns)
#show plot
plt.show()
```



In [5]:

```
pred_test = wine.iloc[3]
```

In [6]:

```
pred_test['type'] = 1
```

```
pred_test.drop(['quality','total sulfur dioxide'],inplace=True)
#pred_test.drop('total_sulfur_dioxide',inplace=True)
pred_test
```

```
Out[6]: fixed acidity      11.200
volatile acidity    0.280
citric acid         0.560
residual sugar      1.900
chlorides           0.075
free sulfur dioxide 17.000
density             0.998
pH                  3.160
sulphates           0.580
alcohol             9.800
type                1.000
Name: 3, dtype: float64
```

```
In [7]: wine.shape
```

```
Out[7]: (1599, 12)
```

```
In [8]: wine.isnull().sum()
```

```
Out[8]: fixed acidity      0
volatile acidity          0
citric acid               0
residual sugar            0
chlorides                 0
free sulfur dioxide        0
total sulfur dioxide       1
density                   0
pH                        1
sulphates                 0
alcohol                   0
quality                   1
dtype: int64
```

```
In [9]: wine2 = wine.fillna(value=wine['total sulfur dioxide'].mean())
wine2 = wine.fillna(value=wine['pH'].mean())
wine2 = wine.fillna(value=wine['quality'].mean())
```

```
In [10]: wine2.isnull().sum()
```

```
Out[10]: 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
```

```
In [11]: wine2.describe()
```

```
Out[11]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	
count	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	159
mean	8.319637	0.527821	0.270976	2.538806	0.087467	15.874922	46.407527	0.996747	
std	1.741096	0.179060	0.194801	1.409928	0.047065	10.460157	32.881792	0.001887	
min	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	5.636421	0.990070	
25%	7.100000	0.390000	0.090000	1.900000	0.070000	7.000000	22.000000	0.995600	
50%	7.900000	0.520000	0.260000	2.200000	0.079000	14.000000	38.000000	0.996750	
75%	9.200000	0.640000	0.420000	2.600000	0.090000	21.000000	62.000000	0.997835	
max	15.900000	1.580000	1.000000	15.500000	0.611000	72.000000	289.000000	1.003690	

In [12]:

```
wine2.info()
```

```
<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	float64

```
dtypes: float64(12)
```

```
memory usage: 150.0 KB
```

In [13]:

```
wine2.isna().any()
```

Out[13]:

fixed acidity	False
volatile acidity	False
citric acid	False
residual sugar	False
chlorides	False
free sulfur dioxide	False
total sulfur dioxide	False
density	False
pH	False
sulphates	False
alcohol	False
quality	False

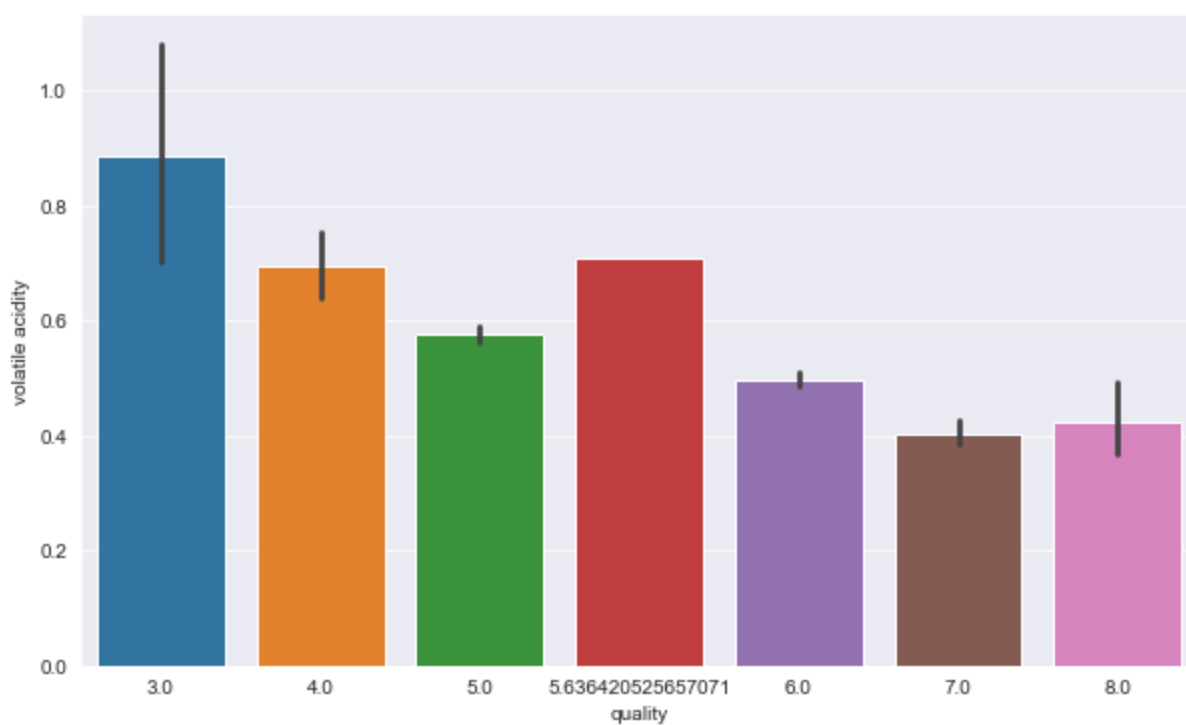
dtype: bool

In [14]:

```
fig = plt.figure(figsize = (10,6))
sns.barplot(x = 'quality', y = 'volatile acidity', data = wine2)
```

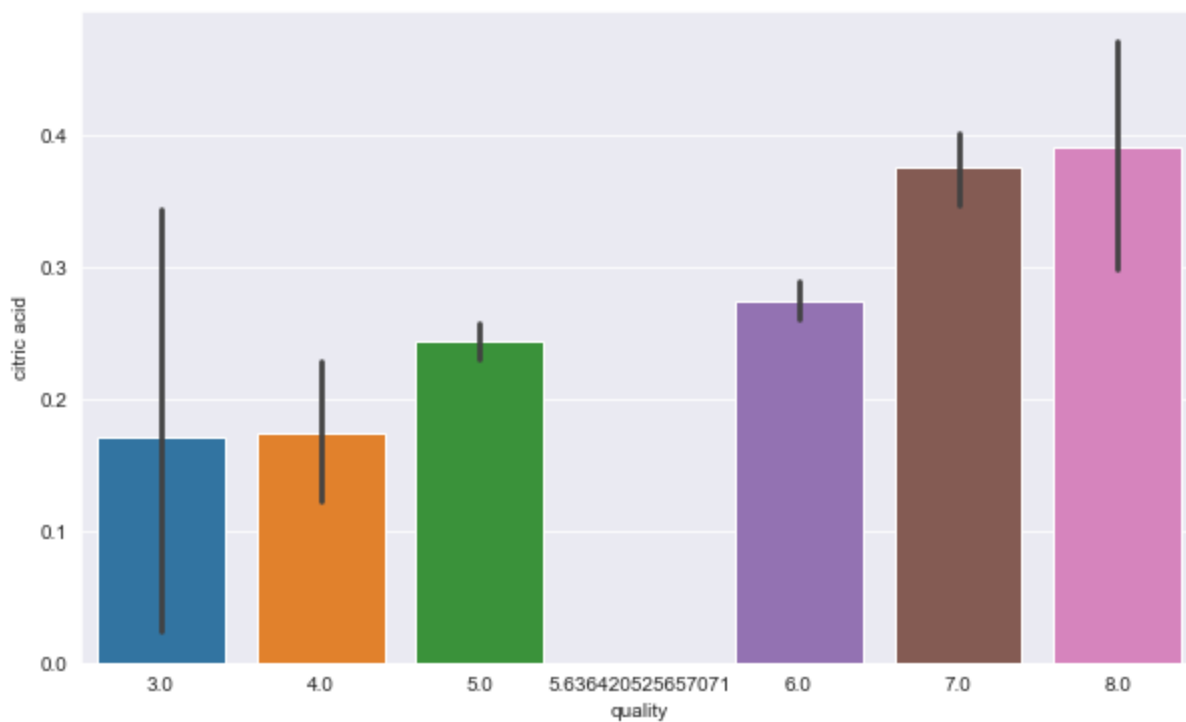
Out[14]:

```
<AxesSubplot:xlabel='quality', ylabel='volatile acidity'>
```



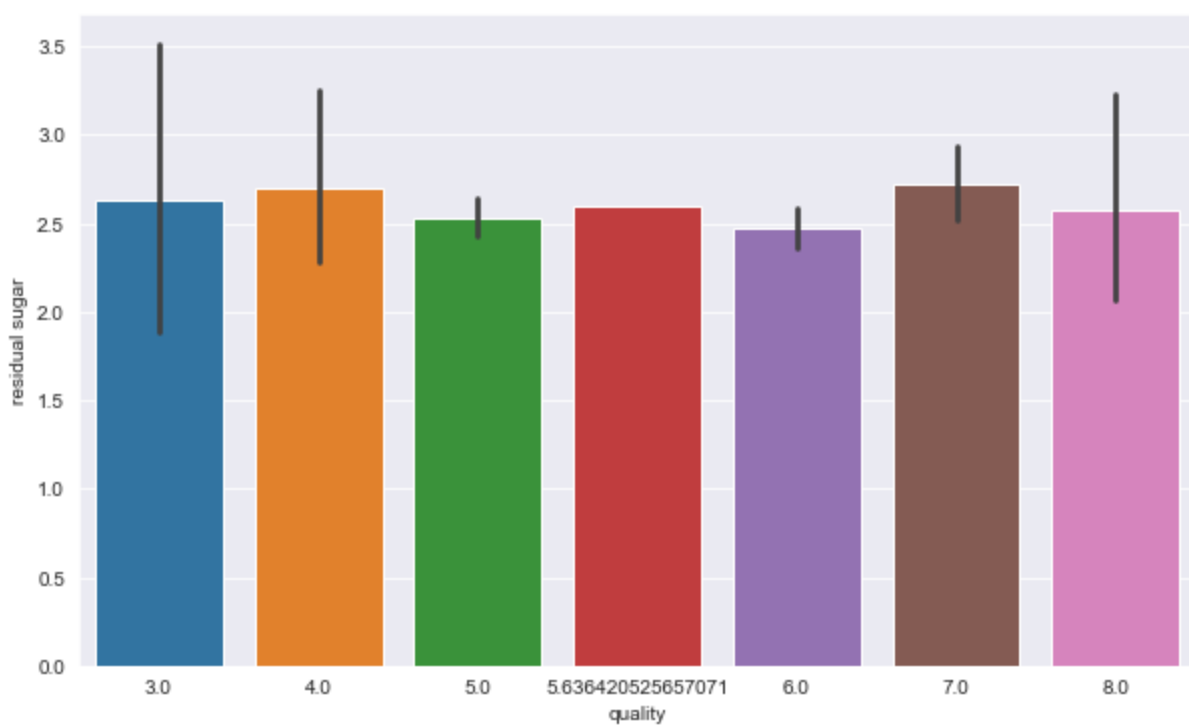
In [15]: `# composition of citric acid go higher as we go higher in the quality of the wine`  
`fig = plt.figure(figsize = (10,6))`  
`sns.barplot(x = 'quality', y = 'citric acid', data = wine2)`

Out[15]: <AxesSubplot:xlabel='quality', ylabel='citric acid'>



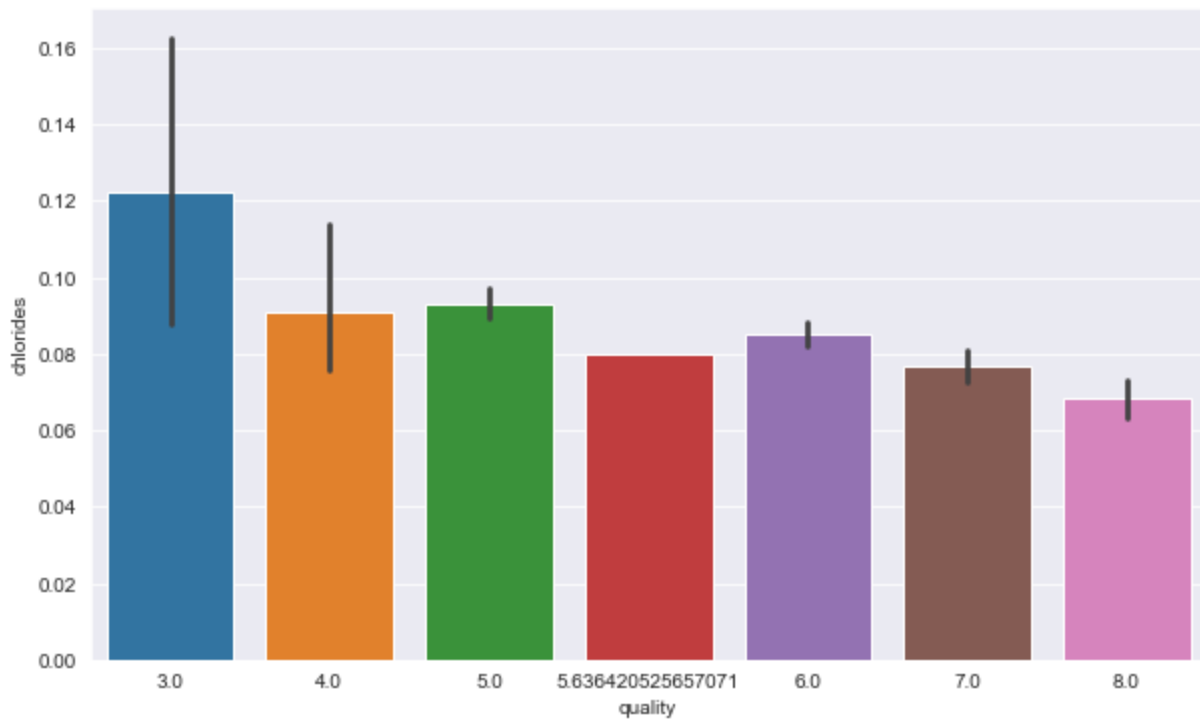
In [16]: `fig = plt.figure(figsize = (10,6))`  
`sns.barplot(x = 'quality', y = 'residual sugar', data = wine2)`

Out[16]: <AxesSubplot:xlabel='quality', ylabel='residual sugar'>

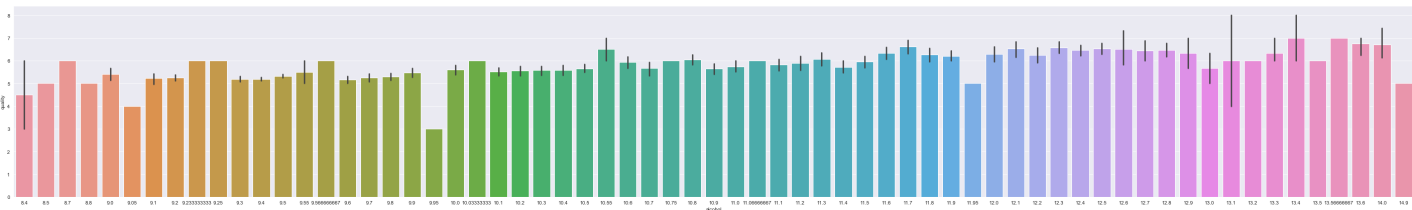


```
In [17]: fig = plt.figure(figsize = (10,6))
sns.barplot(x = 'quality', y = 'chlorides', data = wine2)
```

```
Out[17]: <AxesSubplot:xlabel='quality', ylabel='chlorides'>
```

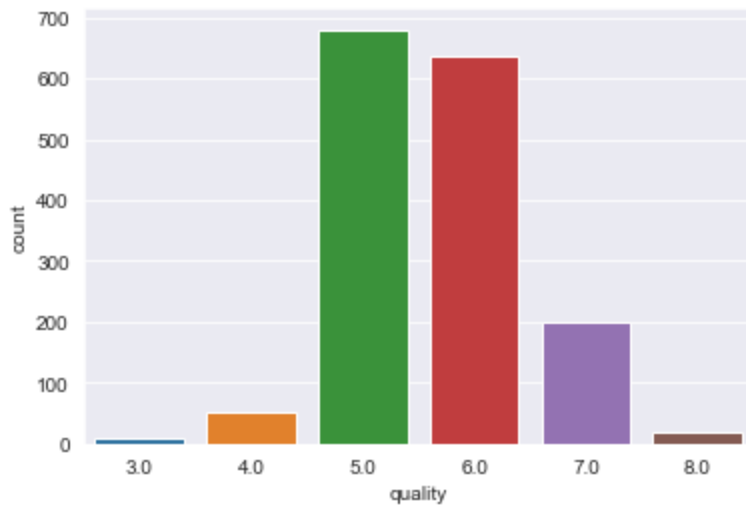


```
In [18]: fig = plt.figure(figsize = (50,7))
ax = sns.barplot(x = 'alcohol', y = 'quality', data = wine2)
```



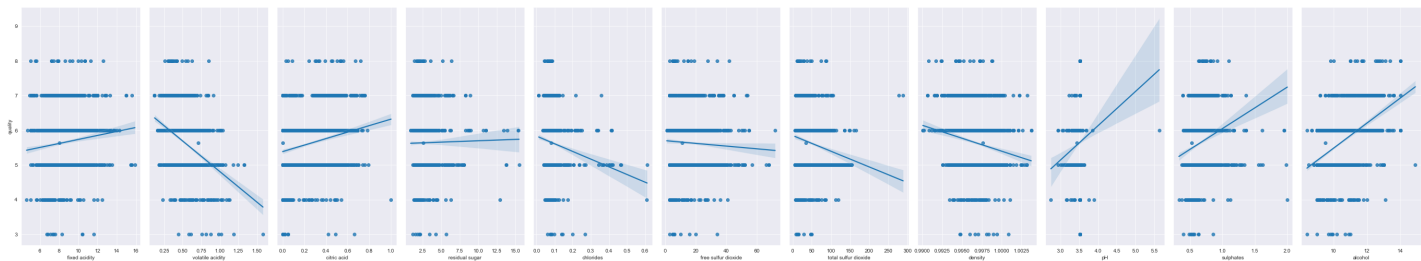
```
In [19]: sns.countplot(wine['quality'])
```

```
Out[19]: <AxesSubplot:xlabel='quality', ylabel='count'>
```



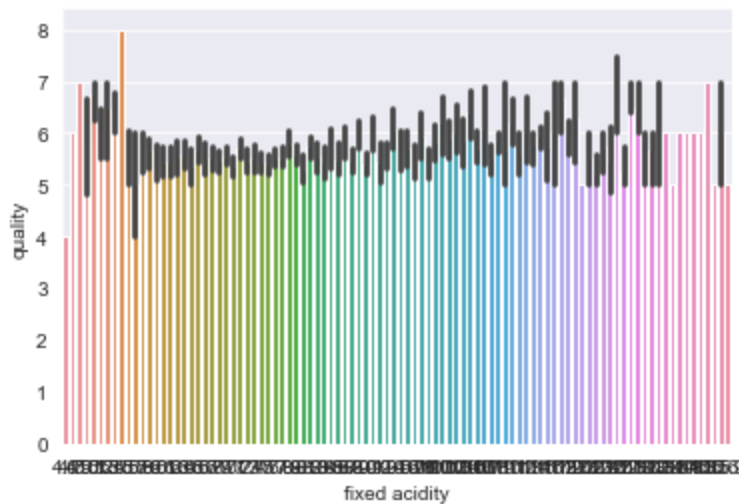
```
In [20]: features = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'chloric  
x = wine2[features]  
y = wine2['quality']  
sns.pairplot(wine2, x_vars=features , y_vars='quality', kind='reg', size=7, aspect=0.5)
```

```
Out[20]: <seaborn.axisgrid.PairGrid at 0x1ef5f693cd0>
```



```
In [21]: sns.barplot(x = 'fixed acidity', y = 'quality', data = wine2)
```

```
Out[21]: <AxesSubplot:xlabel='fixed acidity', ylabel='quality'>
```



```
In [22]: from sklearn.linear_model import LinearRegression  
from sklearn.model_selection import train_test_split  
  
x_train, x_test, y_train, y_test = train_test_split(x, y, random_state = 3)
```

```
In [23]: regressor = LinearRegression()  
         regressor.fit(x_train, y_train)
```

```
Out[23]: ▼ LinearRegression  
         LinearRegression()
```

```
In [24]: accuracy = regressor.score(x_test, y_test)  
         "Accuracy : {}".format(int(round(accuracy*100)))
```

```
Out[24]: 'Accuracy : 38%'
```