

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [3]: df=pd.read_csv(r"C:\Users\Admin\Downloads\11_winequality-red - 11_winequality-r
df
```

Out[3]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcol
0	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.99780	3.51	0.56	
1	7.8	0.880	0.00	2.6	0.098	25.0	67.0	0.99680	3.20	0.68	
2	7.8	0.760	0.04	2.3	0.092	15.0	54.0	0.99700	3.26	0.65	
3	11.2	0.280	0.56	1.9	0.075	17.0	60.0	0.99800	3.16	0.58	
4	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.99780	3.51	0.56	
...	
1594	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	1
1595	5.9	0.550	0.10	2.2	0.062	39.0	51.0	0.99512	3.52	0.76	1
1596	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99574	3.42	0.75	1
1597	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99547	3.57	0.71	1
1598	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	1

1599 rows × 12 columns



```
In [4]: df.info()
```

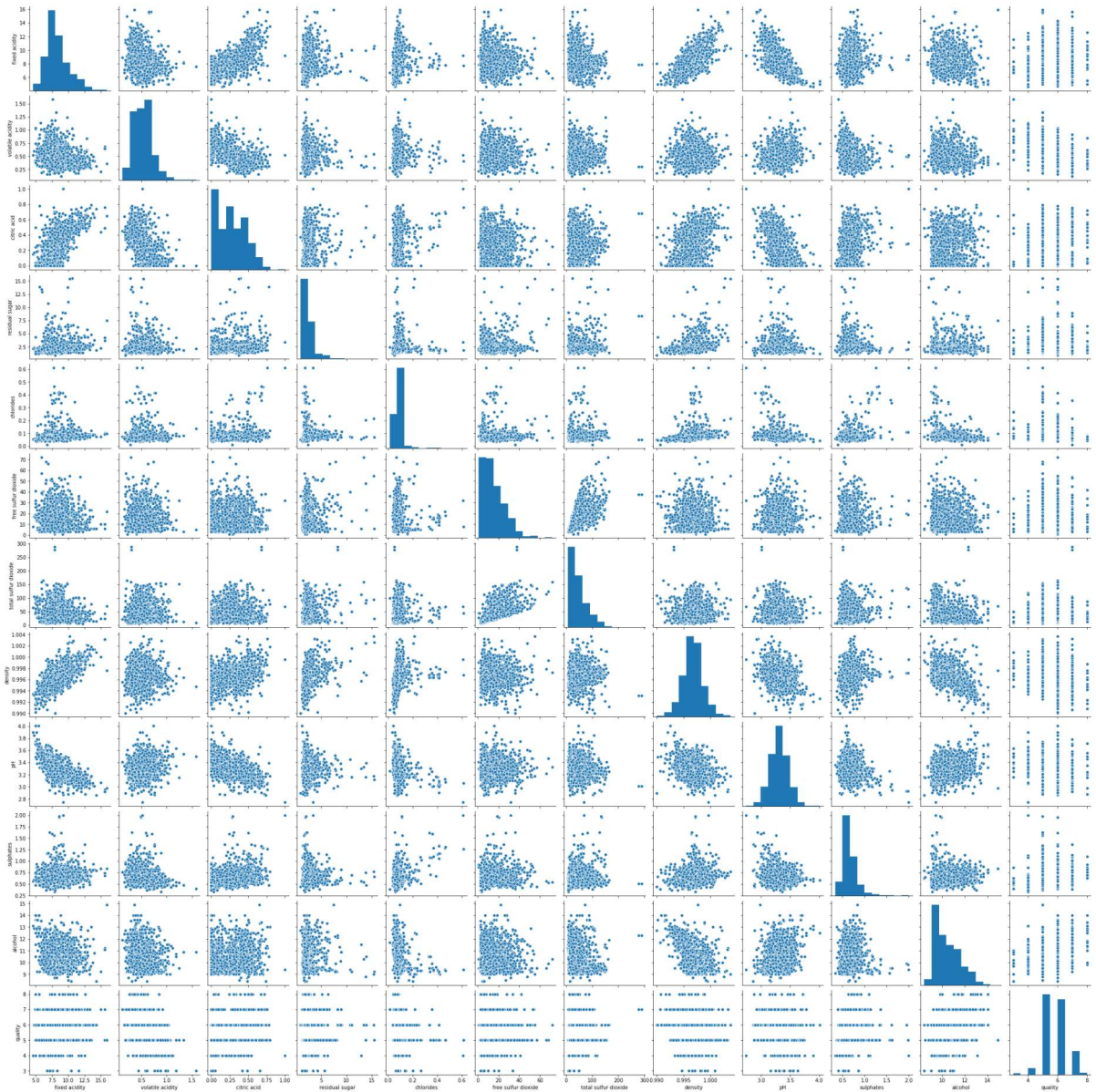
```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1599 entries, 0 to 1598
Data columns (total 12 columns):
#   Column              Non-Null Count  Dtype
---  -
0   fixed acidity        1599 non-null   float64
1   volatile acidity     1599 non-null   float64
2   citric acid          1599 non-null   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   int64
dtypes: float64(11), int64(1)
memory usage: 150.0 KB
```

```
In [5]: df.columns
```

```
Out[5]: Index(['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',  
              'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',  
              'pH', 'sulphates', 'alcohol', 'quality'],  
            dtype='object')
```

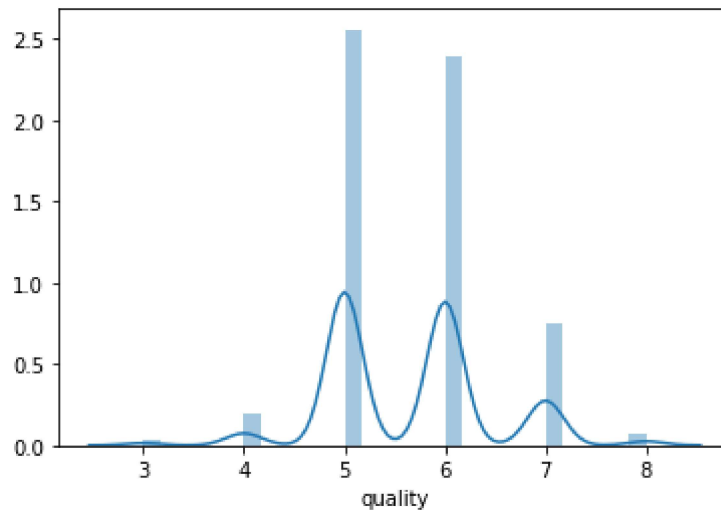
```
In [6]: sns.pairplot(df)
```

```
Out[6]: <seaborn.axisgrid.PairGrid at 0x26b207cf400>
```



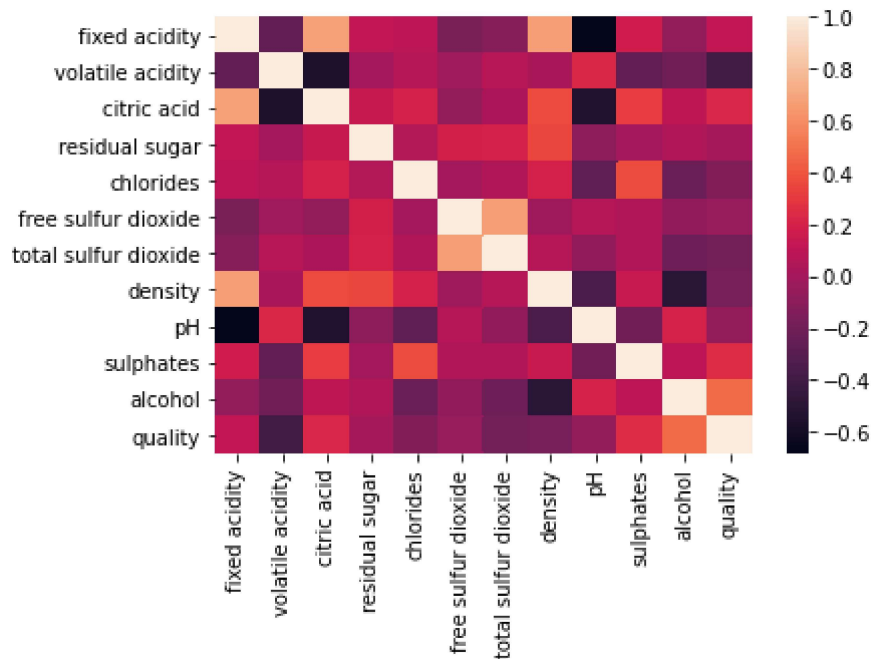
```
In [7]: sns.distplot(df['quality'])
```

```
Out[7]: <matplotlib.axes._subplots.AxesSubplot at 0x26b26ca3f70>
```



```
In [8]: sns.heatmap(df.corr())
```

```
Out[8]: <matplotlib.axes._subplots.AxesSubplot at 0x26b27fe3910>
```



```
In [9]: x=df[['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',
              'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',
              'pH', 'sulphates', 'alcohol']]
y=df['quality']
```

```
In [10]: from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.3)
```

```
In [11]: from sklearn.linear_model import LinearRegression  
lr= LinearRegression()  
lr.fit(x_train,y_train)
```

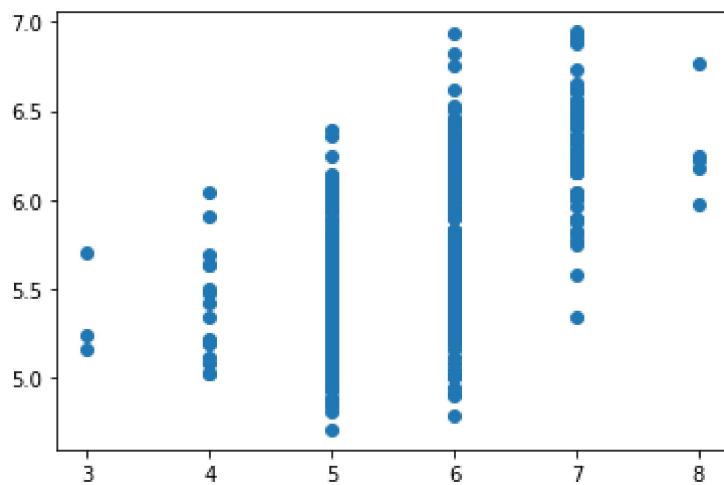
Out[11]: LinearRegression()

```
In [12]: print(lr.intercept_)
```

31.513398155038633

```
In [13]: prediction= lr.predict(x_test)  
plt.scatter(y_test,prediction)
```

Out[13]: <matplotlib.collections.PathCollection at 0x26b28434cd0>



```
In [14]: print(lr.score(x_test,y_test))
```

0.3441083808514308

```
In [ ]:
```