

D7

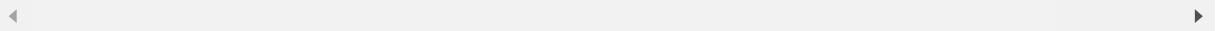
```
In [2]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

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

Out[3]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alco
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 [5]: `df.head(10)`

Out[5]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	9.8
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	9.8
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	9.8
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4
5	7.4	0.66	0.00	1.8	0.075	13.0	40.0	0.9978	3.51	0.56	9.4
6	7.9	0.60	0.06	1.6	0.069	15.0	59.0	0.9964	3.30	0.46	9.4
7	7.3	0.65	0.00	1.2	0.065	15.0	21.0	0.9946	3.39	0.47	10.0
8	7.8	0.58	0.02	2.0	0.073	9.0	18.0	0.9968	3.36	0.57	9.5
9	7.5	0.50	0.36	6.1	0.071	17.0	102.0	0.9978	3.35	0.80	10.5

In [6]: `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 [7]: `df.describe()`

Out[7]:

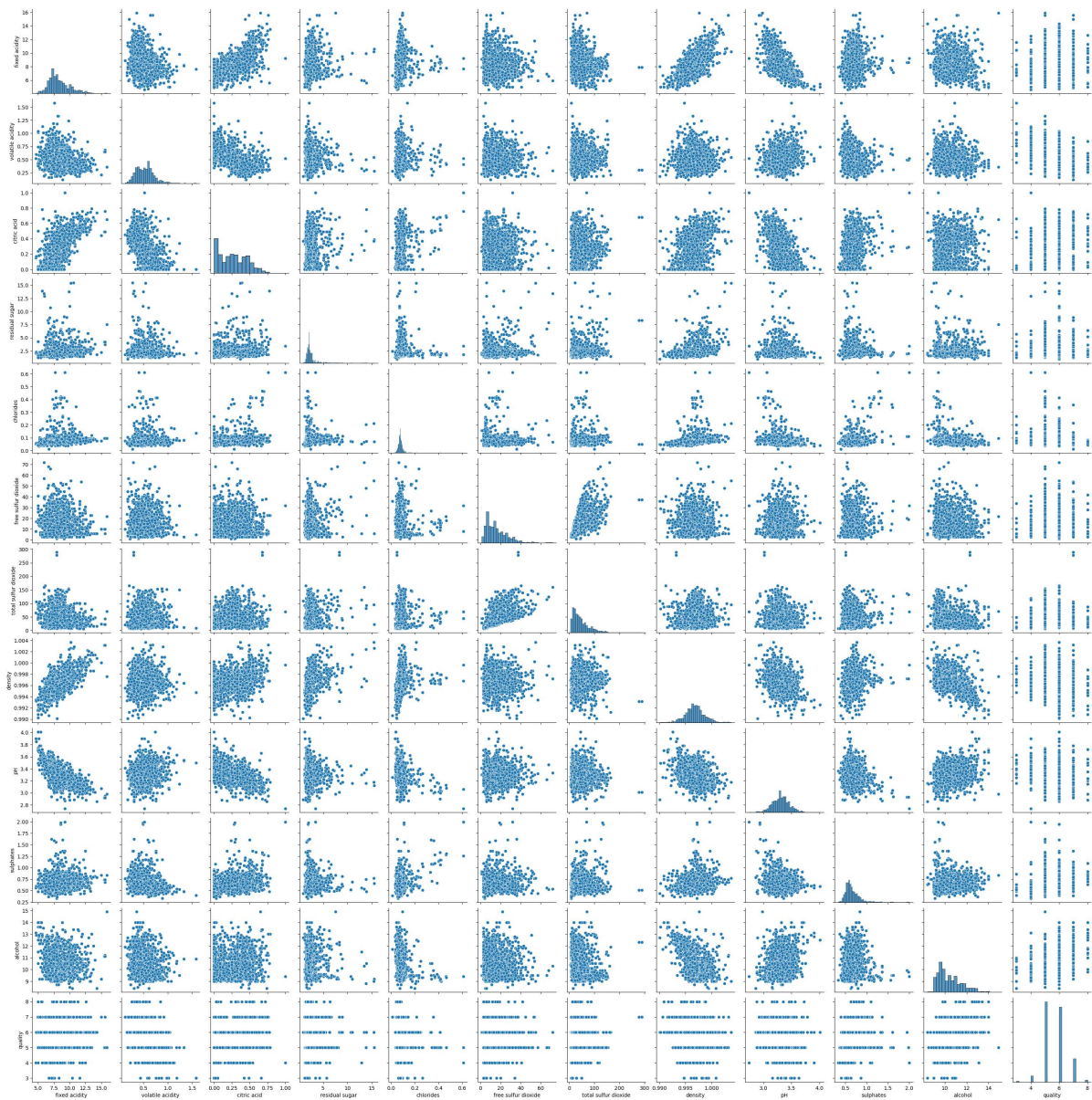
	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	
t	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	159
1	8.319637	0.527821	0.270976	2.538806	0.087467	15.874922	46.467792	
d	1.741096	0.179060	0.194801	1.409928	0.047065	10.460157	32.895324	
1	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	6.000000	
6	7.100000	0.390000	0.090000	1.900000	0.070000	7.000000	22.000000	
6	7.900000	0.520000	0.260000	2.200000	0.079000	14.000000	38.000000	
6	9.200000	0.640000	0.420000	2.600000	0.090000	21.000000	62.000000	
x	15.900000	1.580000	1.000000	15.500000	0.611000	72.000000	289.000000	

In [8]: `df.columns`

Out[8]: Index(['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density', 'pH', 'sulphates', 'alcohol', 'quality'], dtype='object')

```
In [9]: sns.pairplot(df)
```

```
Out[9]: <seaborn.axisgrid.PairGrid at 0x1663b050790>
```



```
In [10]: sns.distplot(df['quality'])
```

C:\Users\user\AppData\Local\Temp\ipykernel_8124\3043670886.py:1: UserWarning:

`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

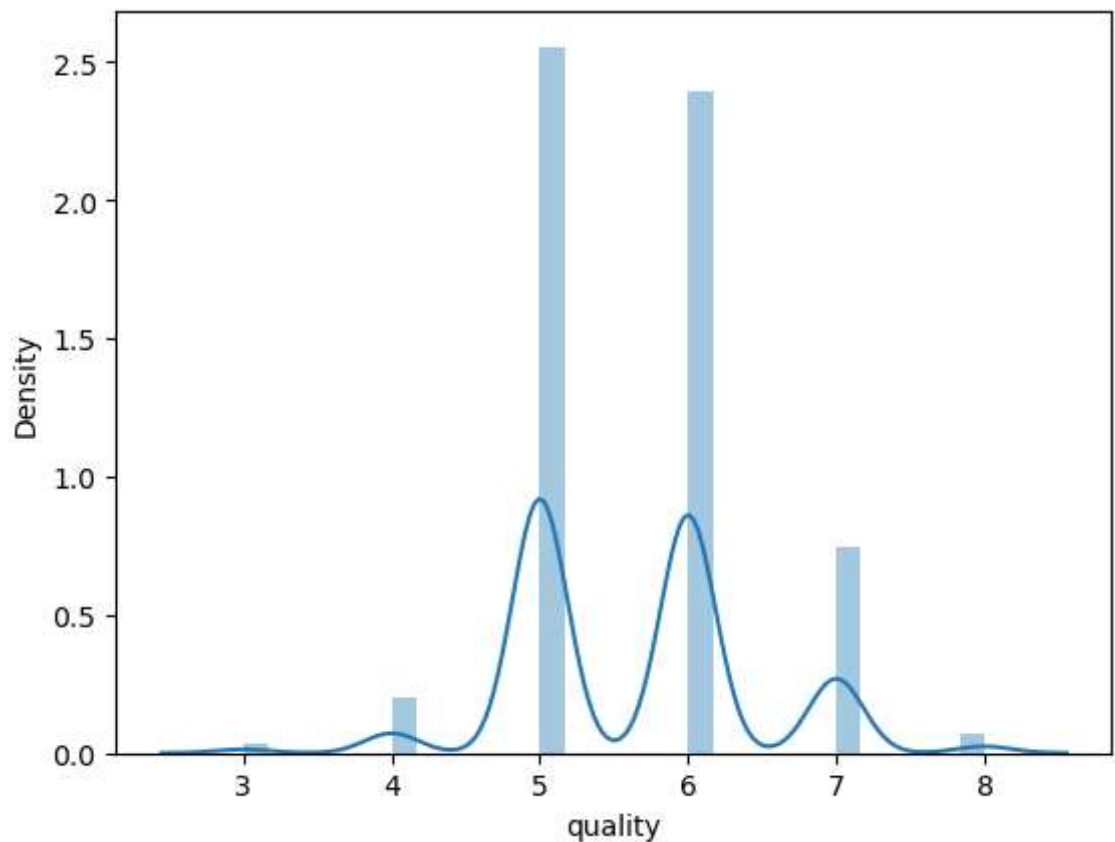
Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see

<https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751> (<https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>)

```
sns.distplot(df['quality'])
```

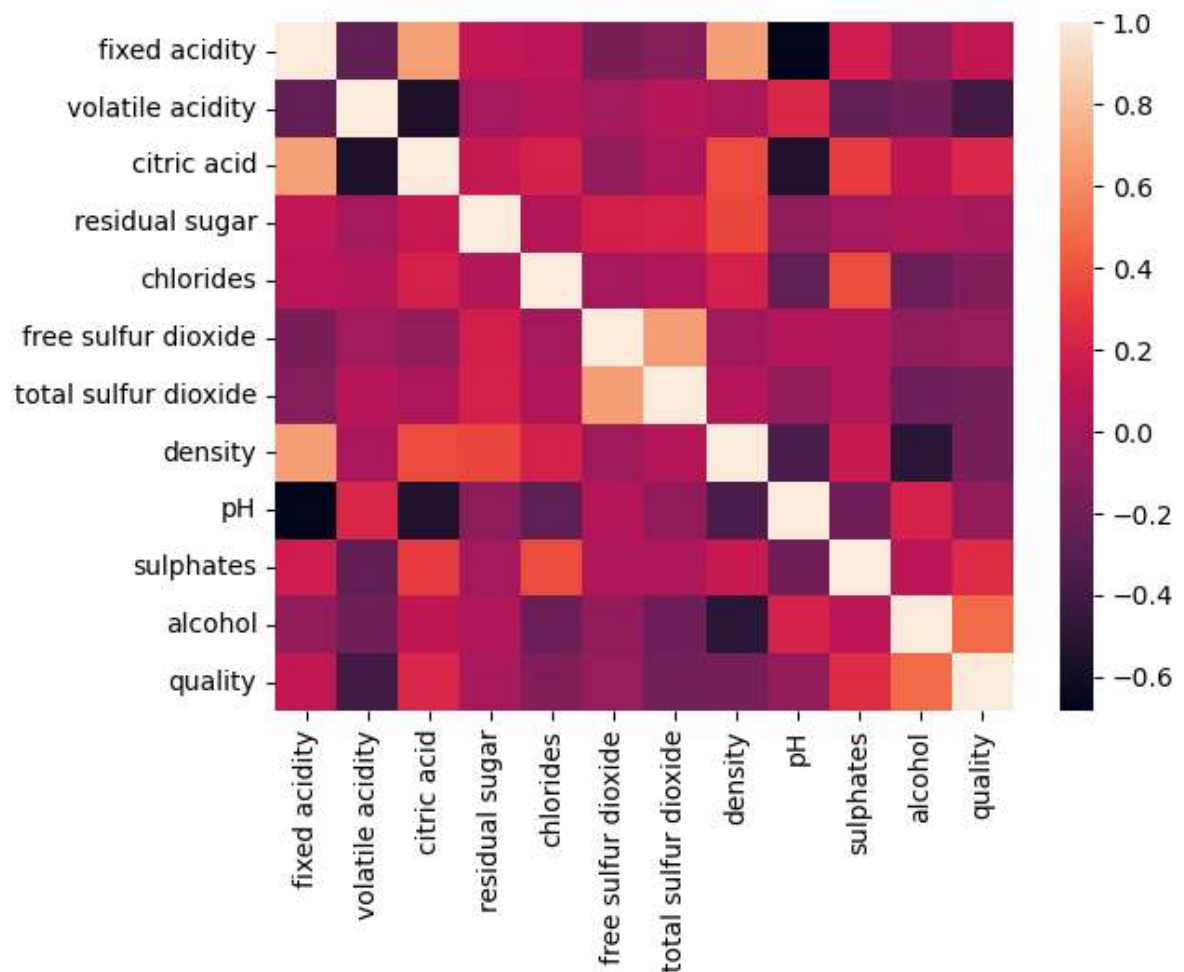
Out[10]: <Axes: xlabel='quality', ylabel='Density'>



```
In [12]: df1=df[['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',  
                'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',  
                'pH', 'sulphates', 'alcohol', 'quality']]
```

```
In [13]: sns.heatmap(df1.corr())
```

```
Out[13]: <Axes: >
```



```
In [14]: x=df1[['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',
               'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',
               'pH', 'sulphates', 'alcohol']]
y=df1["quality"]
```

```
In [15]: 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 [16]: from sklearn.linear_model import LinearRegression
lr=LinearRegression()
lr.fit(x_train,y_train)
```

```
Out[16]: LinearRegression
LinearRegression()
```

```
In [17]: print(lr.intercept_)

16.389277251021372
```

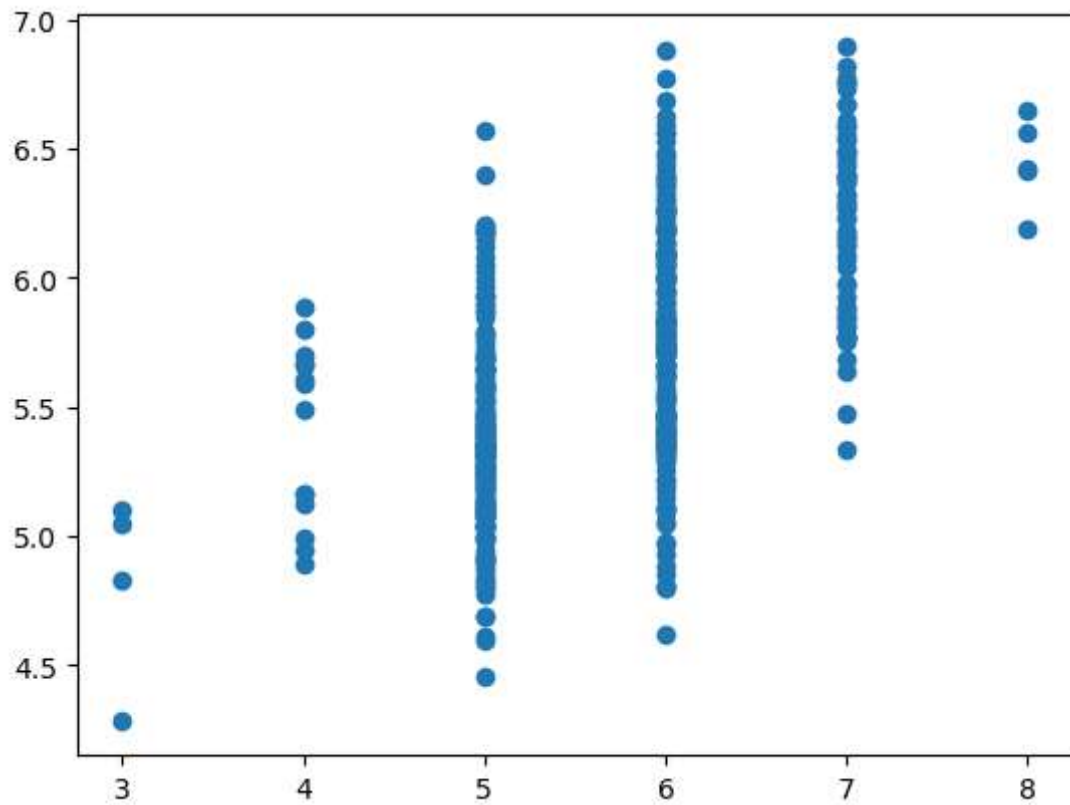
```
In [18]: coeff = pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])  
coeff
```

Out[18]:

	Co-efficient
fixed acidity	0.023438
volatile acidity	-1.215550
citric acid	-0.215720
residual sugar	-0.002547
chlorides	-1.426121
free sulfur dioxide	0.004208
total sulfur dioxide	-0.003038
density	-12.410726
pH	-0.354250
sulphates	0.791818
alcohol	0.285467

```
In [19]: prediction=lr.predict(x_test)  
plt.scatter(y_test,prediction)
```

Out[19]: <matplotlib.collections.PathCollection at 0x1664e18bd10>



In [20]: `print(lr.score(x_test,y_test))`

0.3589522506388996

In []: