

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score
```

```
In [2]: wine_dataset=pd.read_csv('winequality-red.csv')
```

```
In [3]: wine_dataset.shape
```

Out[3]: (1599, 12)

```
In [4]: wine_dataset.head()
```

Out[4]:

	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
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	9.8	5
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	9.8	5
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	9.8	6
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4	5

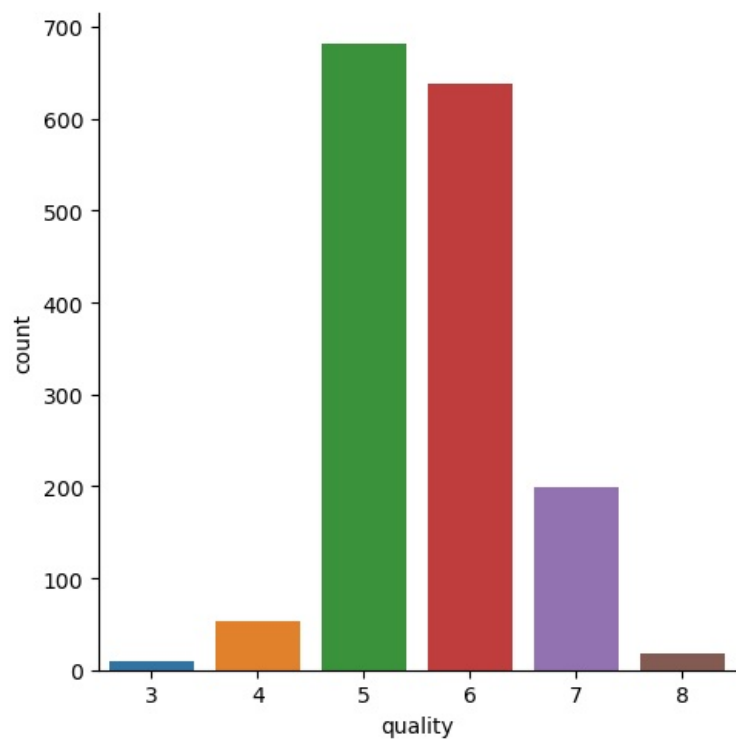
```
In [5]: wine_dataset.describe()
```

Out[5]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	
count	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1
mean	8.319637	0.527821	0.270976	2.538806	0.087467	15.874922	46.467792	0.996747	3.311113	0.658149	
std	1.741096	0.179060	0.194801	1.409928	0.047065	10.460157	32.895324	0.001887	0.154386	0.169507	
min	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	6.000000	0.990070	2.740000	0.330000	
25%	7.100000	0.390000	0.090000	1.900000	0.070000	7.000000	22.000000	0.995600	3.210000	0.550000	
50%	7.900000	0.520000	0.260000	2.200000	0.079000	14.000000	38.000000	0.996750	3.310000	0.620000	
75%	9.200000	0.640000	0.420000	2.600000	0.090000	21.000000	62.000000	0.997835	3.400000	0.730000	
max	15.900000	1.580000	1.000000	15.500000	0.611000	72.000000	289.000000	1.003690	4.010000	2.000000	

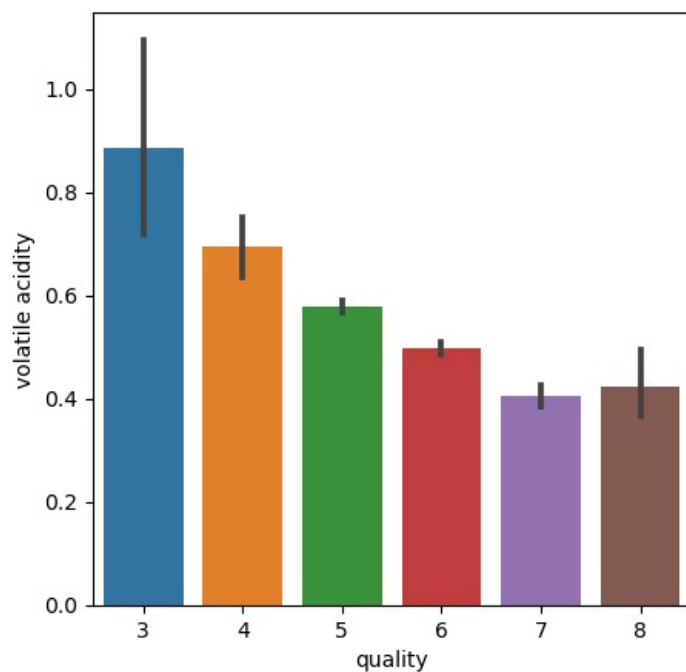
```
In [6]: sns.catplot(x='quality',data=wine_dataset,kind='count')
```

Out[6]: <seaborn.axisgrid.FacetGrid at 0x21a129ffa00>



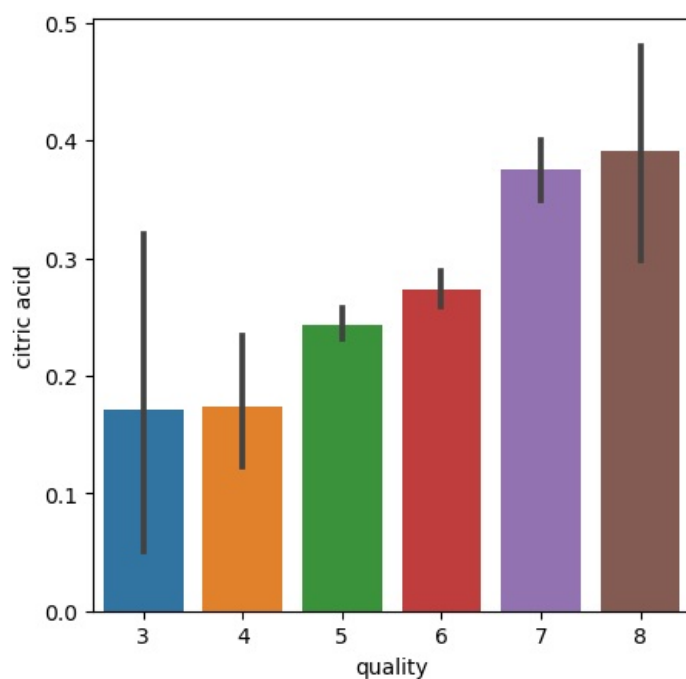
```
In [7]: plot=plt.figure(figsize=(5,5))
sns.barplot(x='quality',y='volatile acidity',data=wine_dataset)
```

Out[7]: <Axes: xlabel='quality', ylabel='volatile acidity'>



```
In [8]: plot=plt.figure(figsize=(5,5))
sns.barplot(x='quality',y='citric acid',data=wine_dataset)
```

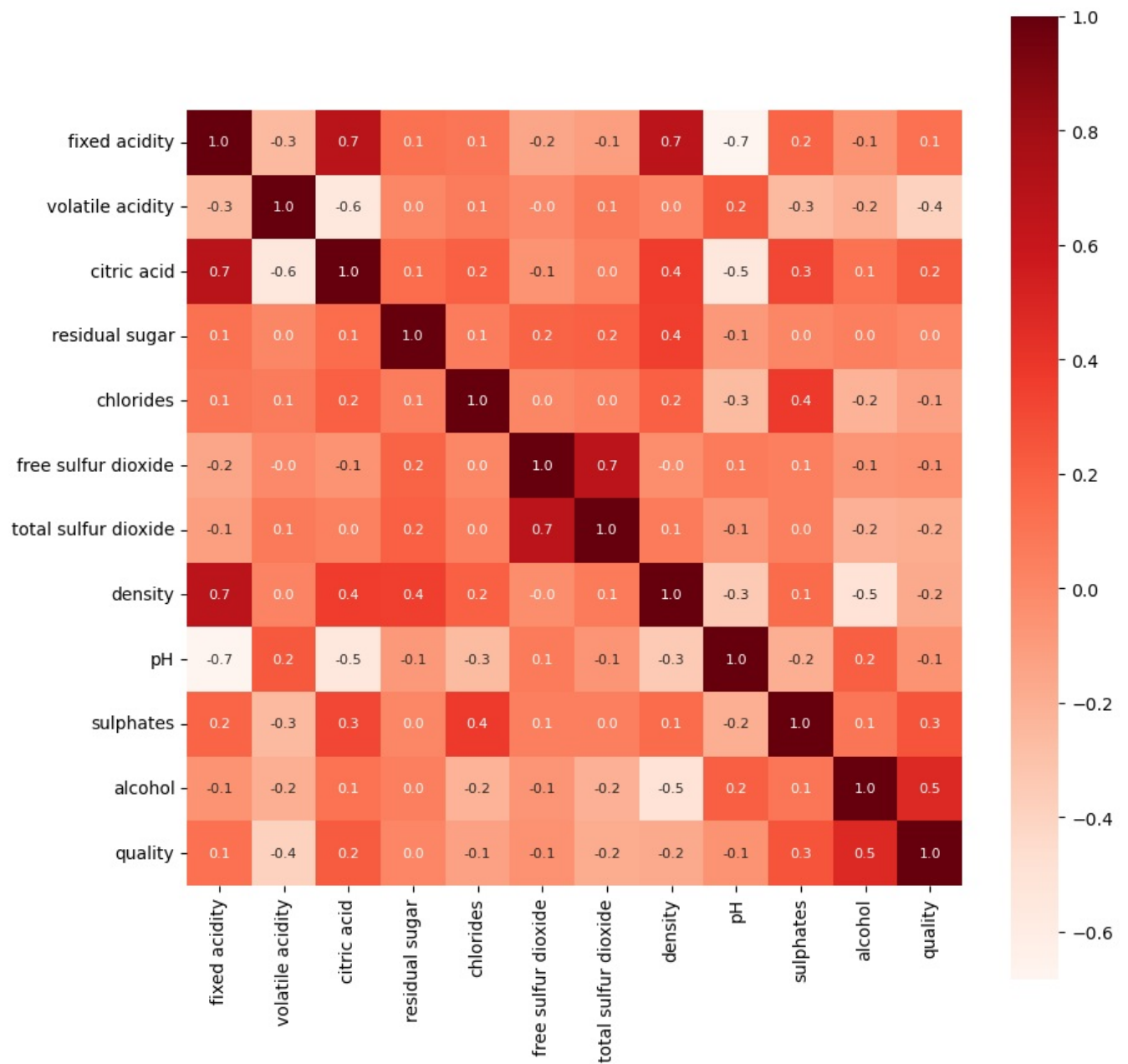
Out[8]: <Axes: xlabel='quality', ylabel='citric acid'>



```
In [9]: correlation=wine_dataset.corr()
```

```
In [13]: plt.figure(figsize=(10,10))
sns.heatmap(correlation,cbar=True, square=True, fmt='.1f',annot=True, annot_kws={'size':8},cmap='Reds')
```

Out[13]: <Axes: >



```
In [14]: x=wine_dataset.drop('quality',axis=1)
```

```
In [15]: print(x)
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
0	7.4	0.700	0.00	1.9	0.076	
1	7.8	0.880	0.00	2.6	0.098	
2	7.8	0.760	0.04	2.3	0.092	
3	11.2	0.280	0.56	1.9	0.075	
4	7.4	0.700	0.00	1.9	0.076	
...	...	...	...	...	...	
1594	6.2	0.600	0.08	2.0	0.090	
1595	5.9	0.550	0.10	2.2	0.062	
1596	6.3	0.510	0.13	2.3	0.076	
1597	5.9	0.645	0.12	2.0	0.075	
1598	6.0	0.310	0.47	3.6	0.067	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
0	11.0	34.0	0.99780	3.51	0.56	
1	25.0	67.0	0.99680	3.20	0.68	
2	15.0	54.0	0.99700	3.26	0.65	
3	17.0	60.0	0.99800	3.16	0.58	
4	11.0	34.0	0.99780	3.51	0.56	
...	...	...	...	...	...	
1594	32.0	44.0	0.99490	3.45	0.58	
1595	39.0	51.0	0.99512	3.52	0.76	
1596	29.0	40.0	0.99574	3.42	0.75	
1597	32.0	44.0	0.99547	3.57	0.71	
1598	18.0	42.0	0.99549	3.39	0.66	

	alcohol
0	9.4
1	9.8
2	9.8
3	9.8
4	9.4
...	...
1594	10.5
1595	11.2
1596	11.0
1597	10.2
1598	11.0

[1599 rows x 11 columns]

```
In [17]: y=wine_dataset['quality'].apply(lambda y_value: 1 if y_value>=7 else 0)
```

```
In [18]: print(y)
```

```
0      0
1      0
2      0
3      0
4      0
...
1594    0
1595    0
1596    0
1597    0
1598    0
Name: quality, Length: 1599, dtype: int64
```

```
In [20]: x_train, x_test,y_train, y_test=train_test_split(x,y,test_size=0.2,random_state=3)
```

```
In [22]: print(y.shape,y_train.shape,y_test.shape)
```

```
(1599,) (1279,) (320,)
```

```
In [23]: model=RandomForestClassifier()
```

```
In [24]: model.fit(x_train,y_train)
```

```
Out[24]: ▼ RandomForestClassifier
RandomForestClassifier()
```

```
In [25]: x_test_prediction=model.predict(x_test)
test_data_accuracy=accuracy_score(x_test_prediction,y_test)
```

```
In [26]: print('Accuracy:',test_data_accuracy)
```

```
Accuracy: 0.921875
```

```
In [29]: input_data= (7.3,0.65,0.0,1.2,0.065,15.0,21.0,0.9946,3.39,0.47,10.0)

input_data_as_numpy_array=np.asarray(input_data)

input_data_resaped=input_data_as_numpy_array.reshape(1,-1)
prediction=model.predict(input_data_resaped)
print(prediction)
if (prediction[0]==1):
```

```
    print("Good Quality Wine")
else:
    print("Bad Quality Wine")
```

```
[1]
Good Quality Wine
```

```
C:\ProgramData\anaconda3\lib\site-packages\sklearn\base.py:420: UserWarning: X does not have valid feature names, but RandomForestClassifier was fitted with feature names
  warnings.warn(
```

```
In [ ]:
```

```
In [ ]:
```

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