

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

```
In [2]: sns.set_style("darkgrid")
plt.rcParams["font.size"] = 15
plt.rcParams['figure.figsize'] = (10,7)
plt.rcParams['figure.facecolor'] = '#FFE5B4'
```

```
In [3]: wine_dataset = pd.read_csv("winequality-red.csv")
```

```
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
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	

```
In [5]: wine_dataset.shape
```

Out[5]: (1599, 12)

```
In [6]: wine_dataset.head()
```

Out[6]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	

```
In [7]: wine_dataset.head()
```

Out[7]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	

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

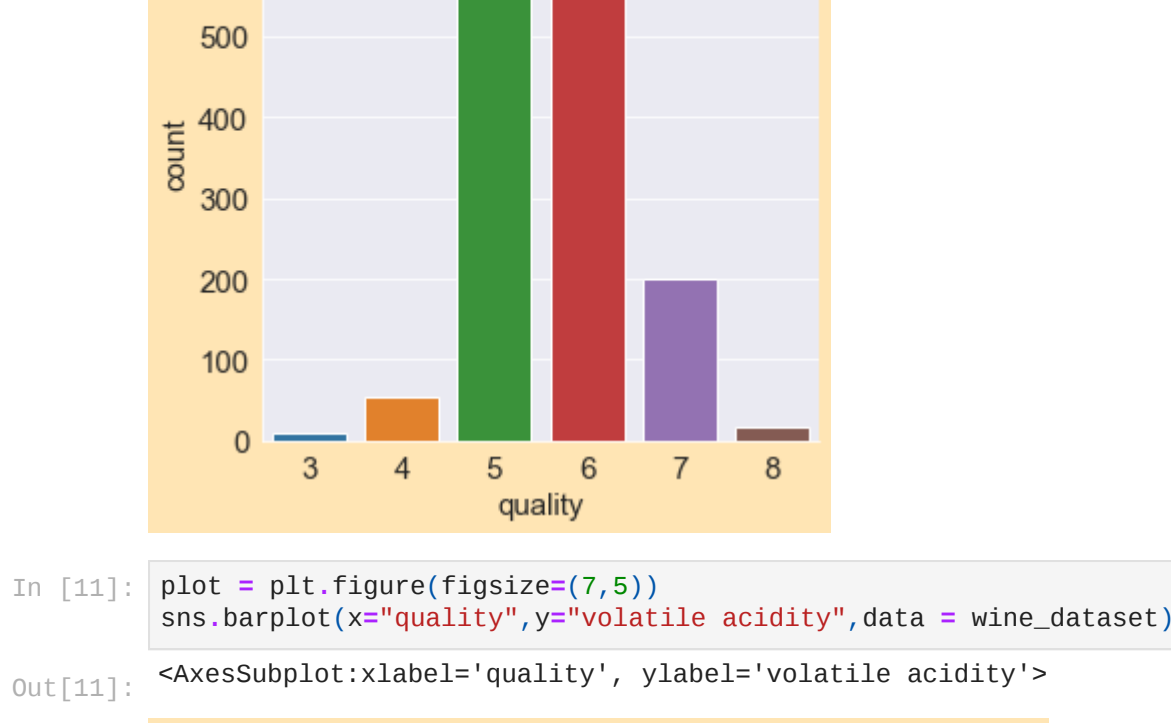
Out[8]: fixed acidity 0  
volatile acidity 0  
citric acid 0  
residual sugar 0  
chlorides 0  
free sulfur dioxide 0  
total sulfur dioxide 0  
density 0  
pH 0  
sulphates 0  
alcohol 0  
quality 0  
dtype: int64

```
In [9]: wine_dataset.describe()
```

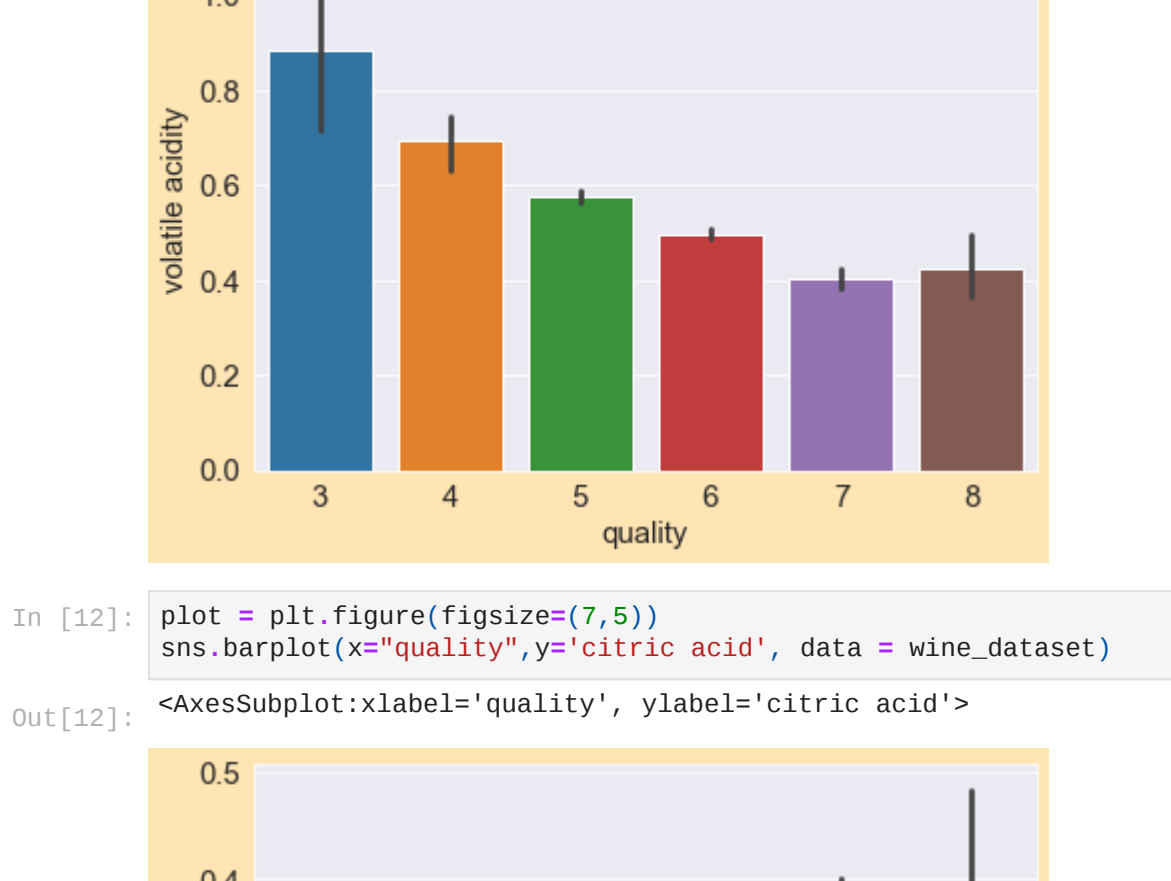
Out[9]:

	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
mean	8.319637	0.527821	0.270976	2.538806	0.087467	15.874000	44.159699	0.997145	3.429539	0.469708
std	1.741096	0.179060	0.194801	1.409928	0.047065	10.460000	22.010000	0.002561	0.080000	0.076000
min	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	1.000000	0.990000	2.940000	0.000000
25%	7.100000	0.390000	0.090000	1.900000	0.070000	7.000000	14.000000	0.996000	3.160000	0.000000
50%	7.900000	0.520000	0.260000	2.200000	0.079000	14.000000	34.000000	0.997000	3.200000	0.000000
75%	9.200000	0.640000	0.420000	2.600000	0.090000	21.000000	60.000000	0.998000	3.160000	0.000000
max	15.900000	1.580000	1.000000	15.500000	0.611000	72.000000	220.000000	1.000000	4.010000	0.900000

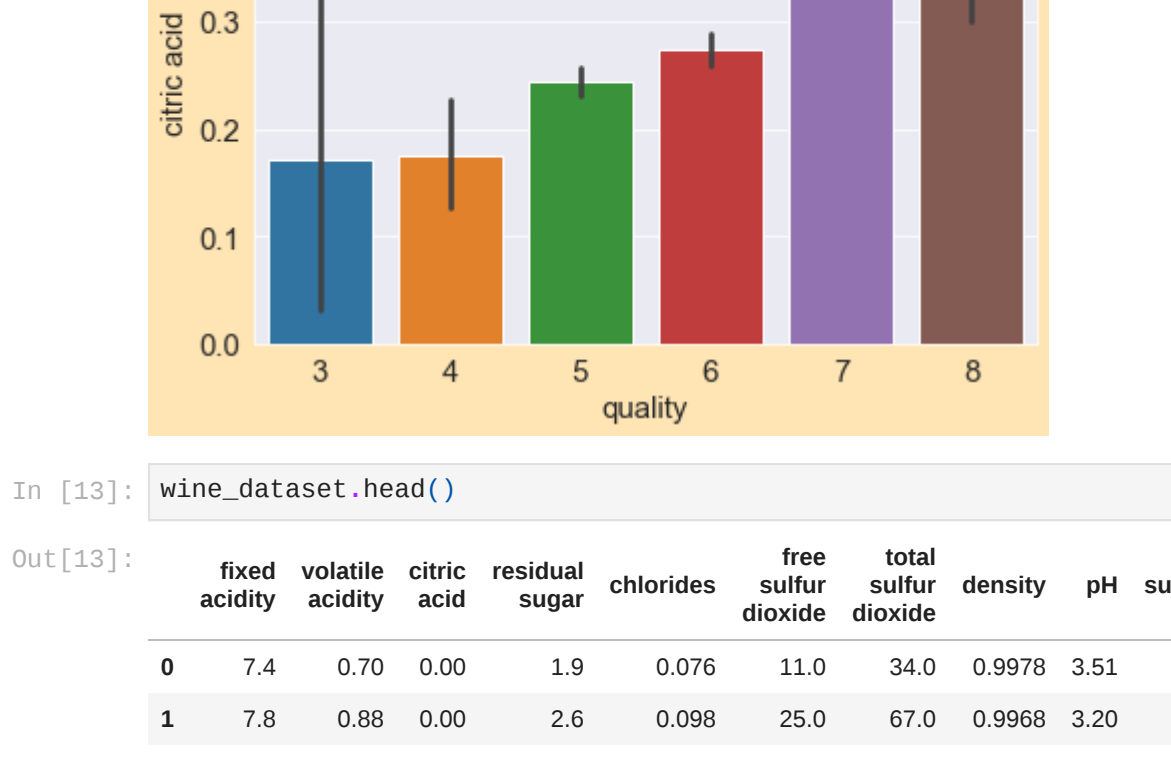
```
In [10]: plot=plt.figure(figsize=(7,5))
sns.catplot(x = "quality",data = wine_dataset, kind = "count")
```



```
In [11]: plot = plt.figure(figsize=(7,5))
sns.barplot(x="quality",y="volatile acidity",data = wine_dataset)
```



```
In [12]: plot = plt.figure(figsize=(7,5))
sns.barplot(x="quality",y='citric acid', data = wine_dataset)
```



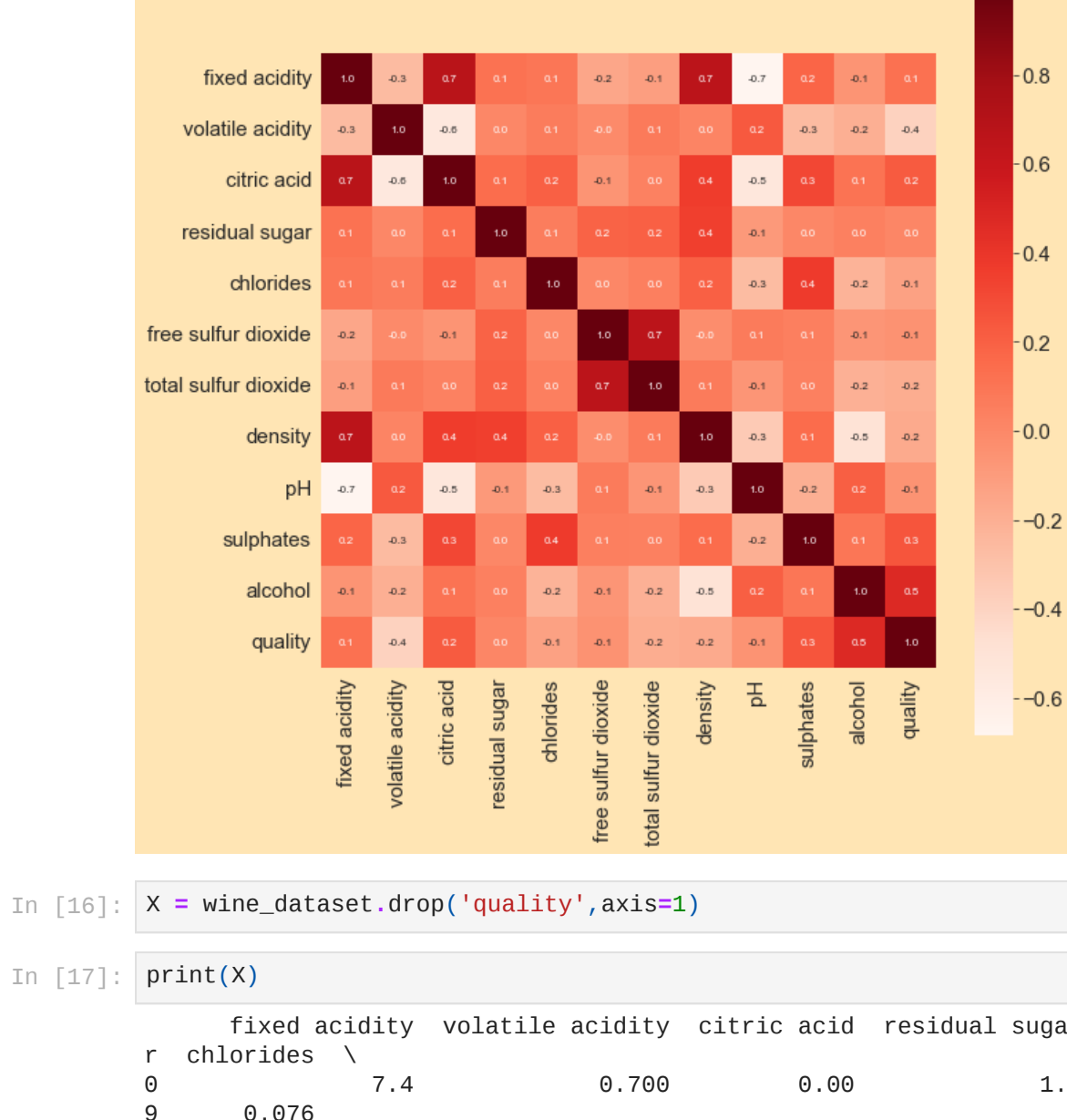
```
In [13]: wine_dataset.head()
```

Out[13]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	

```
In [14]: correlation = wine_dataset.corr()
```

```
In [15]: plt.figure(figsize=(10,10))
sns.heatmap(correlation,cbar=True,square=True,fmt=".1f",annot=True)
```



```
In [16]: X = wine_dataset.drop('quality',axis=1)
```

```
In [17]: print(X)
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
0	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
1	7.8	0.880	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	
2	7.8	0.760	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	
3	11.2	0.280	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	
4	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	
...	...	...	...	...	...	...	...	...	...	...
1594	6.2	0.600	0.08	2.6	0.090	25.0	67.0	0.9968	3.20	
1595	5.9	0.550	0.10	2.3	0.062	15.0	54.0	0.9970	3.26	
1596	6.3	0.510	0.13	2.3	0.076	15.0	54.0	0.9970	3.26	
1597	5.9	0.645	0.12	2.3	0.075	17.0	60.0	0.9980	3.16	
1598	6.0	0.310	0.47	2.3	0.067	15.0	54.0	0.9970	3.26	
...	...	...	...	...	...	...	...	...	...	...
1594	32.0	44.0	0.99490	3.45	0.58	39.0	51.0	0.99512	3.52	
1595	39.0	51.0	0.99512	3.52	0.76	29.0	40.0	0.99574	3.42	
1596	29.0	40.0	0.99574	3.42	0.75	32.0	44.0	0.99547	3.57	
1597	18.0	42.0	0.99549	3.39	0.66	...	...	...	...	
...	...	...	...	...	...	...	...	...	...	
1594	10.5	11.2	0.99512	3.52	0.58	39.0	51.0	0.99512	3.52	
1595	11.0	11.0	0.99512	3.52	0.76	29.0	40.0	0.99574	3.42	
1596	11.0	11.0	0.99574	3.42	0.75	32.0	44.0	0.99547	3.57	
1597	10.2	10.2	0.99549	3.39	0.66	...	...	...	...	
1598	11.0	11.0	0.99549	3.39	0.66	...	...	...	...	

[1599 rows x 11 columns]

```
In [18]: Y = wine_dataset["quality"].apply(lambda y_value: 1 if y_value>=7
```

```
In [19]: Y
```

Out[19]: 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=0.3
```

```
In [21]: print(Y.shape,y_train.shape,y_test.shape)
```

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

```
In [22]: model = RandomForestClassifier()
```

```
In [26]: model.fit(x_train,y_train)
```

Out[26]: RandomForestClassifier()

```
In [27]: #Accuracy score
```

```
In [33]: x_test_prediction = model.predict(x_test)
test_data_accuracy= accuracy_score(x_test_prediction,y_test)
```

```
In [35]: print("Accuracy: " , test_data_accuracy)
```

Accuracy: 0.9375

```
In [37]: input_data = (7.3,0.65,0.0,1.2,0.065,15.0,21.0,0.9946,3.39,0.047,
```

```
In [38]: input_data_as_numpy_array = np.asarray(input_data)
```

```
In [39]: input_data_reshaped = input_data_as_numpy_array.reshape(1,-1)
```

```
In [41]: prediction = model.predict(input_data_reshaped)
```

C:\Users\kisho\anaconda3\lib\site-packages\sklearn\base.py:450: UserWarning: X does not have valid feature names, but RandomForestClassifier was fitted with feature names  
warnings.warn(

```
In [42]: print(prediction)
```

[1]  
Good quality wine

```
In [ ]:
```