```
In [3]: import numpy as np
import pandas as pd

import seaborn as sns
import plotly.express as px
import matplotlib.pyplot as plt

import warnings
warnings.filterwarnings('ignore')

%matplotlib inline
```

In [4]: df=pd.read_csv("C:/Users/sindh/Downloads/archive (6)/WineQT.csv")

In [5]: df.head()

Out[5]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	рН	sulphates	alco
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	
4	_	_	_	_	_	_	_	_			

In [6]: print(df.columns) print(df.shape)

In [7]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1143 entries, 0 to 1142
Data columns (total 13 columns):

#	Column	Non-Null Count	Dtype
0	fixed acidity	1143 non-null	float64
1	volatile acidity	1143 non-null	float64
2	citric acid	1143 non-null	float64
3	residual sugar	1143 non-null	float64
4	chlorides	1143 non-null	float64
5	free sulfur dioxide	1143 non-null	float64
6	total sulfur dioxide	1143 non-null	float64
7	density	1143 non-null	float64
8	рН	1143 non-null	float64
9	sulphates	1143 non-null	float64
10	alcohol	1143 non-null	float64
11	quality	1143 non-null	int64
12	Id	1143 non-null	int64

dtypes: float64(11), int64(2)

memory usage: 116.2 KB

In [8]: df.describe().T

Out[8]:

	count	mean	std	min	25%	50%	75%	
fixed acidity	1143.0	8.311111	1.747595	4.60000	7.10000	7.90000	9.100000	15.9
volatile acidity	1143.0	0.531339	0.179633	0.12000	0.39250	0.52000	0.640000	1.5
citric acid	1143.0	0.268364	0.196686	0.00000	0.09000	0.25000	0.420000	1.0
residual sugar	1143.0	2.532152	1.355917	0.90000	1.90000	2.20000	2.600000	15.5
chlorides	1143.0	0.086933	0.047267	0.01200	0.07000	0.07900	0.090000	0.6
free sulfur dioxide	1143.0	15.615486	10.250486	1.00000	7.00000	13.00000	21.000000	68.0
total sulfur dioxide	1143.0	45.914698	32.782130	6.00000	21.00000	37.00000	61.000000	289.0
density	1143.0	0.996730	0.001925	0.99007	0.99557	0.99668	0.997845	1.0
рН	1143.0	3.311015	0.156664	2.74000	3.20500	3.31000	3.400000	4.0
sulphates	1143.0	0.657708	0.170399	0.33000	0.55000	0.62000	0.730000	2.0
alcohol	1143.0	10.442111	1.082196	8.40000	9.50000	10.20000	11.100000	14.9
quality	1143.0	5.657043	0.805824	3.00000	5.00000	6.00000	6.000000	8.0
ld	1143.0	804.969379	463.997116	0.00000	411.00000	794.00000	1209.500000	1597.0

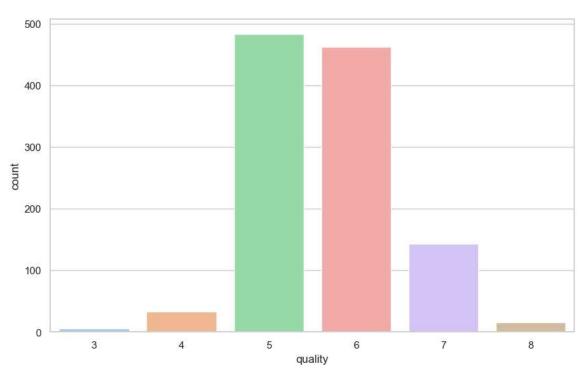
```
In [9]:
         df.nunique()
 Out[9]: fixed acidity
                                     91
         volatile acidity
                                    135
          citric acid
                                     77
          residual sugar
                                     80
          chlorides
                                    131
          free sulfur dioxide
                                     53
          total sulfur dioxide
                                    138
          density
                                     388
                                     87
          рΗ
          sulphates
                                     89
          alcohol
                                     61
          quality
                                      6
          Ιd
                                   1143
          dtype: int64
Out[10]: 0
```

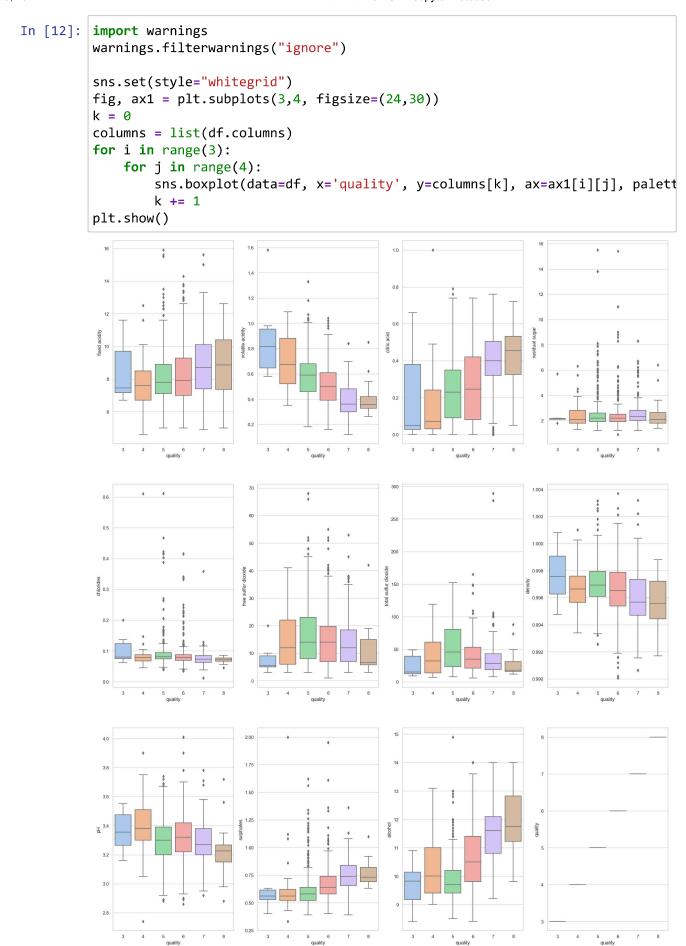
```
In [10]: df.duplicated().sum()
```

```
In [11]: | sns.set(style="whitegrid")
         print(df['quality'].value_counts())
         fig = plt.figure(figsize = (10,6))
         sns.countplot(data=df, x='quality', palette='pastel')
```

Name: count, dtype: int64

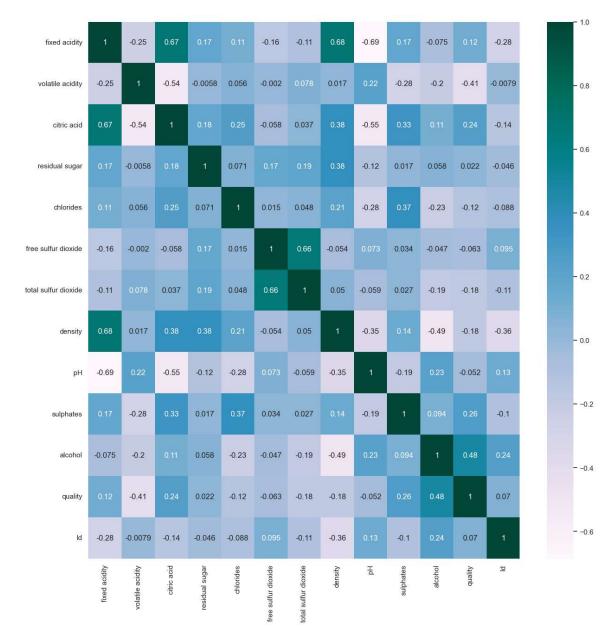
Out[11]: <Axes: xlabel='quality', ylabel='count'>





In [13]: plt.figure(figsize = (15,15))
sns.heatmap(df.corr(),annot=True, cmap= 'PuBuGn')

Out[13]: <Axes: >

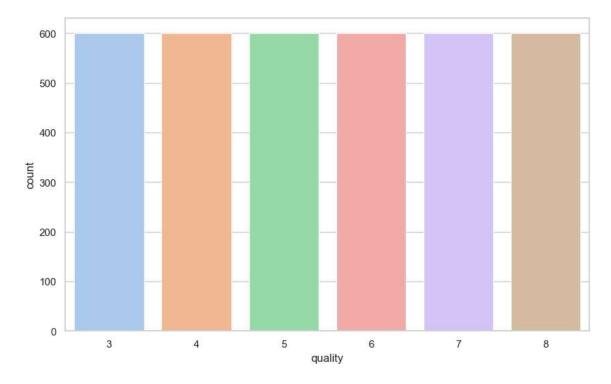


```
In [14]: | color = sns.color_palette("pastel")
            fig, ax1 = plt.subplots(3,4, figsize=(24,30))
            k = 0
            columns = list(df.columns)
            for i in range(3):
                 for j in range(4):
                            sns.distplot(df[columns[k]], ax = ax1[i][j], color = 'red')
            plt.show()
              0.25
                                                                                         0.5
              0.15
                                                                                         0.3
                                         0.00 0.25 0.50 0.75 1.00 1.25 1.50 1.75 volatile acidity
                                                                       0.2
                                                                                              2.5
                                                               0.010
                                               20 40
free sulfur dioxide
```

```
In [15]: | color = sns.color_palette("pastel")
           fig, ax1 = plt.subplots(3,4, figsize=(24,30))
           k = 0
           columns = list(df.columns)
           for i in range(3):
                for j in range(4):
                          sns.distplot(df[columns[k]], ax = ax1[i][j], color = 'green')
           plt.show()
             0.25
             0.15
                                                                                    0.3
                                                            0.0
                                       0.00 0.25 0.50 0.75 1.00 1.25 1.50 1.75 volatile acidity
                                                                0.0
                                                                   0.2
                                                                                        2.5
                                                           0.010
```

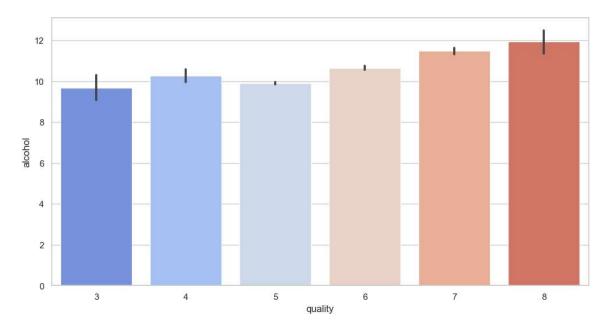
```
In [16]: def log transform(col):
             return np.log(col[0])
         df['residual sugar'] = df[['residual sugar']].apply(log_transform, axis=1)
         df['chlorides'] = df[['chlorides']].apply(log_transform, axis=1)
         df['free sulfur dioxide'] = df[['free sulfur dioxide']].apply(log_transform,
         df['total sulfur dioxide'] = df[['total sulfur dioxide']].apply(log_transfor
         df['sulphates'] = df[['sulphates']].apply(log transform, axis=1)
In [17]: | df.corr()['quality'].sort_values(ascending=False)
Out[17]: quality
                                  1.000000
         alcohol
                                  0.484866
         sulphates
                                  0.315097
         citric acid
                                  0.240821
         fixed acidity
                                 0.121970
         Ιd
                                 0.069708
         residual sugar
                                 0.031487
                                 -0.052453
         free sulfur dioxide
                                 -0.054185
         total sulfur dioxide
                                 -0.170128
         density
                                 -0.175208
         chlorides
                                 -0.175391
         volatile acidity
                                 -0.407394
         Name: quality, dtype: float64
In [18]: | df_3 = df[df.quality==3]
         df_4 = df[df.quality==4]
         df 5 = df[df.quality==5]
         df 6 = df[df.quality==6]
         df_7 = df[df.quality==7]
         df_8 = df[df.quality==8]
In [19]: | from sklearn.utils import resample
         df_3_upsampled = resample(df_3, replace=True, n_samples=600, random_state=12
         df 4 upsampled = resample(df 4, replace=True, n samples=600, random state=12
         df 7 upsampled = resample(df 7, replace=True, n samples=600, random state=12
         df_8_upsampled = resample(df_8, replace=True, n_samples=600, random_state=12
         df 5 downsampled = df[df.quality==5].sample(n=600, replace=True).reset index
         df 6 downsampled = df[df.quality==6].sample(n=600, replace=True).reset index
In [20]: Balanced df = pd.concat([df 3 upsampled, df 4 upsampled, df 7 upsampled,
                                   df_8_upsampled, df_5_downsampled, df_6_downsampled]
         Balanced df.quality.value counts()
Out[20]: quality
         3
              600
         4
              600
         7
              600
         8
              600
         5
              600
              600
         Name: count, dtype: int64
```

Out[24]: <Axes: xlabel='quality', ylabel='count'>



```
In [25]: plt.figure(figsize = (12,6))
sns.barplot(x='quality', y = 'alcohol', data = df, palette = 'coolwarm')
```

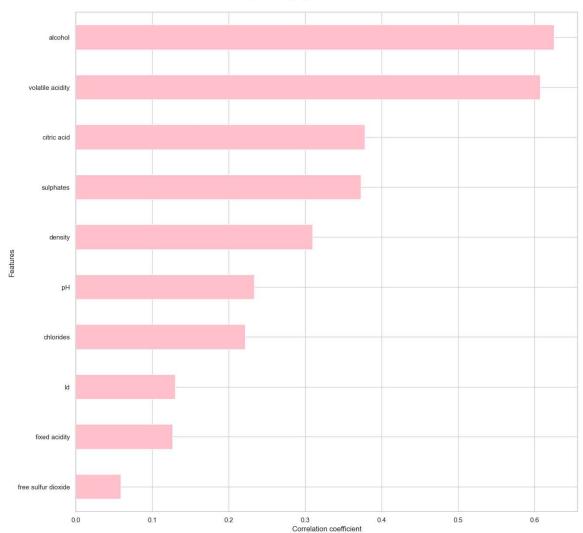
Out[25]: <Axes: xlabel='quality', ylabel='alcohol'>



```
In [26]: plt.figure(figsize=(15,15))
    Balanced_df.corr().quality.apply(lambda x: abs(x)).sort_values(ascending=Fal
    plt.title("Top 10 highly correlated features", size=20, pad=26)
    plt.xlabel("Correlation coefficient")
    plt.ylabel("Features")
```

Out[26]: Text(0, 0.5, 'Features')





```
In [28]: 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,random_
```

```
In [29]:
       from sklearn.neighbors import KNeighborsClassifier
        for n_neighbors in [5,10,15,20]:
           model = KNeighborsClassifier(n_neighbors)
           model.fit(X train, y train)
           scr = model.score(X_test, y_test)
           print("For n_neighbors = ", n_neighbors ," score is ",scr)
        For n_neighbors = 15 score is 0.7240740740740741
        For n_neighbors = 20 score is 0.6657407407407407
In [30]:
       KNN_Model = KNeighborsClassifier(n_neighbors=5, weights='distance')
        KNN_Model.fit(X_train, y_train)
        results = KNN_Model.fit(X_train, y_train)
In [31]: KNN_train_predictions = KNN_Model.predict(X_train)
In [32]: KNN_test_predictions = KNN_Model.predict(X_test)
```

```
In [33]: from sklearn.metrics import classification_report, confusion_matrix

print("\n Train Data: KNN_Confusion Matrix:\n ")
print(confusion_matrix(y_train, KNN_train_predictions))

print("\n Train Data: KNN_Classification Report:\n ")
print(classification_report(y_train, KNN_train_predictions))

print("\n \n Test Data: KNN_Confusion Matrix: \n ")
print(confusion_matrix(y_test, KNN_test_predictions))

print("\n Test Data: KNN_Classification Report:\n ")
print(classification_report(y_test, KNN_test_predictions))
```

Train Data: KNN_Confusion Matrix:

[[4	122	0	0	0	0	0]
[0	392	0	0	0	0]
[0	0	423	0	0	0]
[0	0	0	436	0	0]
[0	0	0	0	423	0]
[0	0	0	0	0	424]]

Train Data: KNN_Classification Report:

	precision	recall	f1-score	support
3	1.00	1.00	1.00	422
4	1.00	1.00	1.00	392
5	1.00	1.00	1.00	423
6	1.00	1.00	1.00	436
7	1.00	1.00	1.00	423
8	1.00	1.00	1.00	424
accuracy			1.00	2520
macro avg	1.00	1.00	1.00	2520
weighted avg	1.00	1.00	1.00	2520

Test Data: KNN_Confusion Matrix:

```
[[178      0      0      0      0      0]
[ 0 208      0      0      0      0]
[ 1 3 147 19 7 0]
[ 1 3 11 133 15 1]
[ 0 0 0 6 171 0]
[ 0 0 0 0 0 176]]
```

Test Data: KNN_Classification Report:

	precision	recall	f1-score	support
3	0.99	1.00	0.99	178
4	0.97	1.00	0.99	208
5	0.93	0.83	0.88	177
6	0.84	0.81	0.83	164
7	0.89	0.97	0.92	177
8	0.99	1.00	1.00	176
			0.04	1000
accuracy			0.94	1080
macro avg	0.94	0.93	0.93	1080
weighted avg	0.94	0.94	0.94	1080

In []: