

```
In [1]: import pandas as pd
```

```
In [2]: df = pd.read_csv('C:/Users/Raghavendra K/Downloads/winequality-red.csv')
```

```
In [3]: df.head()
```

Out[3]:

	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 [4]: df.shape
```

Out[4]: (1599, 12)

```
In [5]: df.tail()
```

Out[5]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
1594	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	10.5	5
1595	5.9	0.550	0.10	2.2	0.062	39.0	51.0	0.99512	3.52	0.76	11.2	6
1596	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99574	3.42	0.75	11.0	6
1597	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99547	3.57	0.71	10.2	5
1598	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6

```
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.isnull().sum()
```

Out[7]:

```
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 [8]: df['quality'].unique()
```

Out[8]: array([5, 6, 7, 4, 8, 3], dtype=int64)

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

	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 [10]:

df.columns

Out[10]:

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

In [11]:

import matplotlib.pyplot as plt
import seaborn as sns

In [12]:

plt.bar(df['quality'],df['fixed acidity'])
plt.xlabel('quality')
plt.ylabel('fixed acidity')
plt.show()

AttributeError

Traceback (most recent call last)

Input In [12], in <cell line: 2>():

1

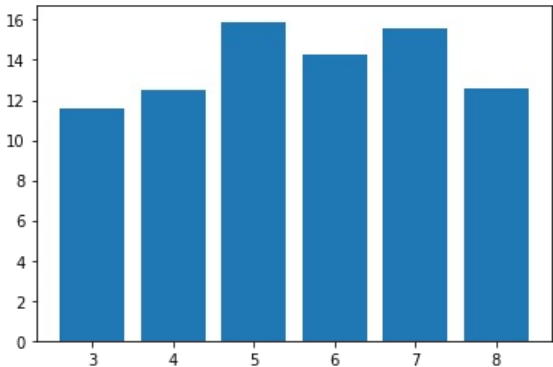
plt.bar(df['quality'],df['fixed acidity'])

----> 2 plt.xlabel('quality')

3 plt.ylabel('fixed acidity')

4 plt.show()

AttributeError: module 'matplotlib.pyplot' has no attribute 'xlabel'



In []:

plt.bar(df['quality'],df['volatile acidity'])
plt.xlabel('quality')
plt.ylabel('volatile acidity')
plt.show()

In []:

plt.bar(df['quality'],df['residual sugar'])
plt.xlabel('quality')
plt.ylabel('residual sugar')
plt.show()

In []:

plt.bar(df['quality'],df['chlorides'])
plt.xlabel('quality')
plt.ylabel('chlorides')
plt.show()

In []:

plt.bar(df['quality'],df['total sulfur dioxide'])
plt.xlabel('quality')
plt.ylabel('total sulfur dioxide')
plt.show()

In []:

plt.bar(df['quality'],df['alcohol'])
plt.xlabel('quality')
plt.ylabel('alcohol')
plt.show()

In []:

plt.figure(figsize=(10,5))
sns.heatmap(df.corr(),annot=True,fmt='0.1f')

```

In [ ]: #Binarization of target variable
df['quality'].unique()

In [ ]: df['quality']=[1 if x>=7 else 0 for x in df['quality']]

In [ ]: df['quality'].unique()

In [ ]: df['quality'].value_counts()

In [ ]: import seaborn as sns

In [ ]: sns.countplot(df['quality'])

In [ ]: #Handling Imbalanced dataset-smote techniques
from imblearn.over_sampling import SMOTE

In [ ]: from imblearn.over_sampling import SMOTE

In [ ]:

In [ ]: x_res,y_res = SMOTE().fit_resample(x,y)

In [ ]: y_res.value_counts()

In [ ]: x = df.drop('quality',axis=1)
y = df['quality']

In [ ]:

In [ ]: x

In [ ]: y

In [ ]: #Split the dataset into train and test
from sklearn.model_selection import train_test_split

In [ ]: x_train,x_test,y_train,y_test=train_test_split(x_res,y_res,test_size=0.20,random_state=42)

In [ ]: #FEATURE SCALING
from sklearn.preprocessing import StandardScaler

In [ ]: st = StandardScaler()
x_train = st.fit_transform(x_train)
x_test = st.transform(x_test)

In [ ]: x_train

```

APPLYING - PRINCIPAL COMPONENT ANALYSIS-PCA

```

In [ ]: from sklearn.decomposition import PCA

In [ ]: pca = PCA(n_components=0.90)

In [ ]: x_train = pca.fit_transform(x_train)
x_test = pca.transform(x_test)

In [ ]: pca.explained_variance_ratio_

In [ ]: sum(pca.explained_variance_ratio_)

In [ ]: #Logistic Regression
from sklearn.linear_model import LogisticRegression

In [ ]: log = LogisticRegression()
log.fit(x_train,y_train)

In [ ]: y_pred1 = log.predict(x_test)

In [ ]: from sklearn.metrics import accuracy_score

In [ ]: accuracy_score(y_test,y_pred1)

In [ ]: from sklearn.metrics import precision_score,recall_score,f1_score

```

```
In [ ]: precision_score(y_test,y_pred1)

In [ ]: recall_score(y_test,y_pred1)

In [ ]: f1_score(y_test,y_pred1)

In [ ]: #SUPPORT VECTOR CLASSIFIER-SVC
        from sklearn import svm

In [ ]: svm=svm.SVC()

In [ ]: svm.fit(x_train,y_train)

In [ ]: y_pred2=svm.predict(x_test)

In [ ]: accuracy_score(y_test,y_pred2)

In [ ]: precision_score(y_test,y_pred2)

In [ ]: f1_score(y_test,y_pred2)

In [ ]: #KNeighbors Classifier
        from sklearn.neighbors import KNeighborsClassifier

In [ ]: knn = KNeighborsClassifier()

In [ ]: knn.fit(x_train,y_train)

In [ ]: y_pred3 = knn.predict(x_test)

In [ ]: accuracy_score(y_test,y_pred3)

In [ ]: precision_score(y_test,y_pred3)

In [ ]: f1_score(y_test,y_pred3)

In [ ]: recall_score(y_test,y_pred3)

In [ ]: #Decision Tree Classifier
        from sklearn.tree import DecisionTreeClassifier

In [ ]: dt = DecisionTreeClassifier()

In [ ]: dt.fit(x_train,y_train)

In [ ]: y_pred4=dt.predict(x_test)

In [ ]: accuracy_score(y_test,y_pred4)

In [ ]: precision_score(y_test,y_pred4)

In [ ]: f1_score(y_test,y_pred4)

In [ ]: #Random Forest Classifier
        from sklearn.ensemble import RandomForestClassifier

In [ ]: from sklearn.ensemble import RandomForestClassifier

In [ ]: rf = RandomForestClassifier()
        rf.fit(x_train,y_train)

In [ ]: y_pred5=rf.predict(x_test)

In [ ]: accuracy_score(y_test,y_pred5)

In [ ]: precision_score(y_test,y_pred5)

In [ ]: f1_score(y_test,y_pred5)

In [ ]: recall_score(y_test,y_pred5)

In [ ]: #Gradient Boosting Classifier
        from sklearn.ensemble import GradientBoostingClassifier
```

```

In [ ]: gbc = GradientBoostingClassifier()
        gbc.fit(x_train,y_train)

In [ ]: y_pred6=gbc.predict(x_test)

In [ ]: accuracy_score(y_test,y_pred6)

In [ ]: precision_score(y_test,y_pred6)

In [ ]: f1_score(y_test,y_pred6)

In [ ]: import pandas as pd

In [ ]: final_data = pd.DataFrame({'Models':['LR', 'SVC', 'KNN', 'DT', 'RF', 'GBC'], 'ACC':[accuracy_score(y_test,y_pred1)*100,
                                                                                       accuracy_score(y_test,y_pred2)*100,
                                                                                       accuracy_score(y_test,y_pred3)*100,
                                                                                       accuracy_score(y_test,y_pred4)*100,
                                                                                       accuracy_score(y_test,y_pred5)*100,
                                                                                       accuracy_score(y_test,y_pred6)*100]})

```

final_data

```

In [ ]: import seaborn as sns

In [ ]: final_data

In [ ]:

In [ ]: sns.barplot(final_data['Models'],final_data['ACC'])

```

SAVE THE MODEL

```

In [ ]: x=df.drop('quality',axis=1)
        y=df['quality']

In [ ]: from imblearn.over_sampling import SMOTE
        x_res,y_res=SMOTE().fit_resample(x,y)

In [ ]: from sklearn.preprocessing import StandardScaler
        st = StandardScaler()
        x=st.fit_transform(x_res)

In [ ]: x=pca.fit_transform(x)

In [ ]: from sklearn.ensemble import RandomForestClassifier
        rf=RandomForestClassifier()
        rf.fit(x,y_res)

In [ ]: import joblib

In [ ]: joblib.dump(rf,'quality_prediction_model')

In [ ]: model = joblib.load('quality_prediction_model')

model = joblib.dump(rf,'quality_prediction_model')

```

PREDICTION ON NEW DATA

```

In [ ]: import pandas as pd
        new_data = pd.DataFrame({
            'fixed acidity':7.3,
            'volatile acidity':0.65,
            'citric acid':0.00,
            'residual sugar':1.2,
            'chlorides':0.065,
            'free sulfur dioxide':15.0,
            'total sulfur dioxide':21.0,
            'density':0.9946,
            'PH':3.39,
            'sulphate':0.47,
            'alcohol':10.0,
        },index=[0])

In [ ]: new_data

```

```
In [ ]: test = pca.transform(st.transform(new_data))
```

```
In [ ]: model.predict(test)
```

```
In [ ]: p = model.predict(test)
```

```
In [ ]: if p[0]==1:
        print("Good Quality Wine")
    else:
        print("Bad Quality Wine")
```

GUI

```
In [ ]: from tkinter import *

        from sklearn.preprocessing import StandardScaler
        import joblib
```

```
In [ ]: def show_entry_fields():
        p1=float(e1.get())
        p2=float(e2.get())
        p3=float(e3.get())
        p4=float(e4.get())
        p5=float(e5.get())
        p6=float(e6.get())
        p7=float(e7.get())
        p8=float(e8.get())
        p9=float(e9.get())
        p10=float(e10.get())
        p11=float(e11.get())
        model = joblib.load('quality_prediction_model')
        result=model.predict(pca.transform(st.transform([[p1,p2,p3,p4,p5,p6,p7,p8,p9,p10,p11]])))

        if p[0]==1:
            Label(master,text="quality_prediction_model").grid(row=31)
            Label(master, text="quality_prediction_model").grid(row=31)

master =Tk()
master.title("quality_prediction_model")
label = Label(master,text = "quality_prediction_model"
               ,bg = "black", fg = "white" ).\
        grid(row=0,columnspan=2)

Label(master, text='fixed acidity').grid(row=1)
Label(master, text='volatile acidity').grid(row=2)
Label(master, text='citric acid').grid(row=3)
Label(master, text='residual sugar').grid(row=4)
Label(master, text='chlorides').grid(row=5)
Label(master, text='free sulfur dioxide').grid(row=6)
Label(master, text='total sulfur dioxide').grid(row=7)
Label(master, text='density').grid(row=8)
Label(master, text='PH').grid(row=9)
Label(master, text='sulphate').grid(row=10)
Label(master, text='alcohol').grid(row=11)

e1=Entry(master)
e2=Entry(master)
e3=Entry(master)
e4=Entry(master)
e5=Entry(master)
e6=Entry(master)
e7=Entry(master)
e8=Entry(master)
e9=Entry(master)
e10=Entry(master)
e11=Entry(master)

e1.grid(row=1, column=1)
e2.grid(row=2, column=1)
e3.grid(row=3, column=1)
e4.grid(row=4, column=1)
e5.grid(row=5, column=1)
e6.grid(row=6, column=1)
e7.grid(row=7, column=1)
e8.grid(row=8, column=1)
e9.grid(row=9, column=1)
e10.grid(row=10, column=1)
e11.grid(row=11, column=1)

Button(master, text='predict', command=show_entry_fields).grid()

mainloop()
```

In []:

In []:

Loading [MathJax]/jax/output/CommonHTML/fonts/TeX/fontdata.js