Task2-Wine Quality Prediction

```
In [2]: import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        import seaborn as sb
        from sklearn.model_selection import train_test_split
        from sklearn.preprocessing import MinMaxScaler
        from sklearn import metrics
        from sklearn.svm import SVC
        from sklearn.linear_model import LogisticRegression
        import warnings
        warnings.filterwarnings('ignore')
In [3]: | df = pd.read_csv("WineQT.csv")
        print(df.head())
           fixed acidity volatile acidity citric acid residual sugar chlori
        des \
                     7.4
                                      0.70
                                                    0.00
        0
                                                                     1.9
                                                                              0.
        076
                     7.8
        1
                                      0.88
                                                    0.00
                                                                     2.6
                                                                              0.
        098
        2
                     7.8
                                      0.76
                                                   0.04
                                                                     2.3
                                                                              0.
        092
                    11.2
                                      0.28
                                                    0.56
                                                                     1.9
                                                                              0.
        075
        4
                     7.4
                                      0.70
                                                    0.00
                                                                     1.9
                                                                              0.
        076
           free sulfur dioxide total sulfur dioxide density
                                                                 pH sulphates
        \
        0
                                                 34.0
                                                       0.9978 3.51
                          11.0
                                                                           0.56
        1
                          25.0
                                                 67.0
                                                       0.9968 3.20
                                                                           0.68
        2
                          15.0
                                                 54.0
                                                       0.9970 3.26
                                                                           0.65
        3
                          17.0
                                                 60.0
                                                       0.9980 3.16
                                                                           0.58
```

```
In [4]: 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 [5]: df.describe().T

Out	[5]	:

	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.!
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.
chlorides	1143.0	0.086933	0.047267	0.01200	0.07000	0.07900	0.090000	0.0
free sulfur dioxide	1143.0	15.615486	10.250486	1.00000	7.00000	13.00000	21.000000	68.(
total sulfur dioxide	1143.0	45.914698	32.782130	6.00000	21.00000	37.00000	61.000000	289.(
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
4								•

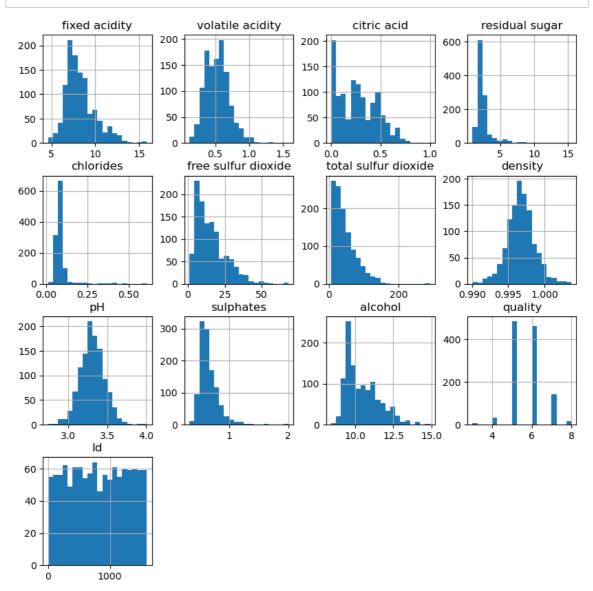
EDTA-Analysis

```
df.isnull().sum()
In [6]:
Out[6]: fixed acidity
                                 0
        volatile acidity
                                 0
        citric acid
                                 0
        residual sugar
                                 0
        chlorides
        free sulfur dioxide
                                 0
        total sulfur dioxide
                                 0
        density
                                 0
        рΗ
        sulphates
                                 0
        alcohol
                                 0
        quality
        Ιd
                                 0
        dtype: int64
In [7]: | for col in df.columns:
          if df[col].isnull().sum() > 0:
            df[col] = df[col].fillna(df[col].mean())
        df.isnull().sum().sum()
```

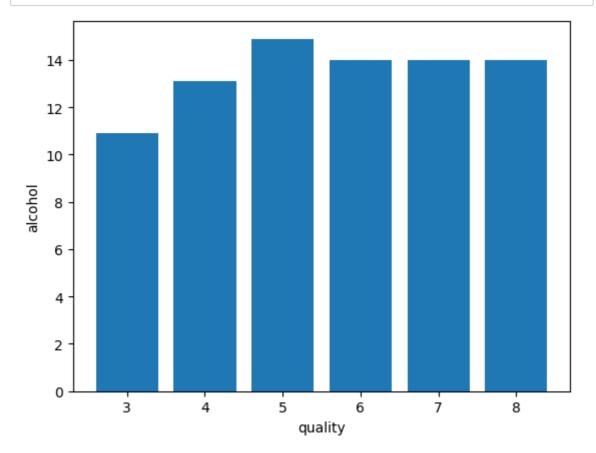
Out[7]: 0

Histogram

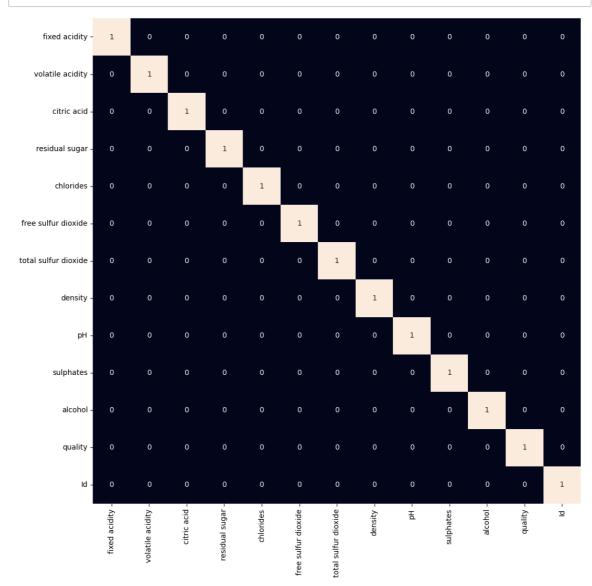
In [8]: df.hist(bins=20, figsize=(10, 10))
 plt.show()



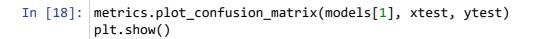
```
In [9]: plt.bar(df['quality'], df['alcohol'])
    plt.xlabel('quality')
    plt.ylabel('alcohol')
    plt.show()
```

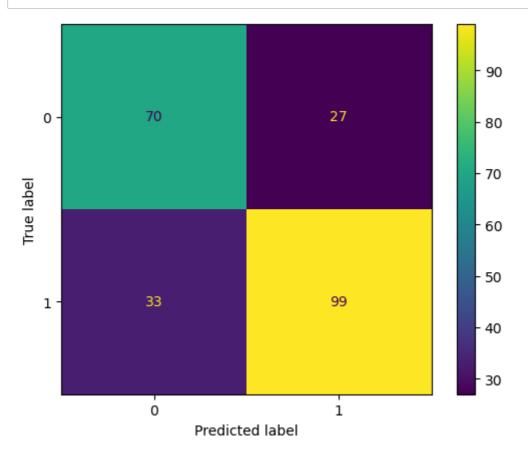


```
In [10]: plt.figure(figsize=(12, 12))
    sb.heatmap(df.corr() > 0.7, annot=True, cbar=False)
    plt.show()
```



```
#Normalizing the data
In [15]:
         norm = MinMaxScaler()
         xtrain = norm.fit_transform(xtrain)
         xtest = norm.transform(xtest)
In [17]: models = [LogisticRegression(),SVC(kernel='rbf')]
         for i in range(3):
             models[i].fit(xtrain, ytrain)
             print(f'{models[i]} : ')
             print('Training Accuracy : ', metrics.roc_auc_score(ytrain, models[i].p
             print('Validation Accuracy : ', metrics.roc_auc_score(
                 ytest, models[i].predict(xtest)))
             print()
         LogisticRegression() :
         Training Accuracy: 0.7546950559364851
         Validation Accuracy : 0.7255154639175256
         SVC():
         Training Accuracy : 0.7648213641284736
         Validation Accuracy: 0.7358247422680412
         IndexError
                                                   Traceback (most recent call las
         t)
         ~\AppData\Local\Temp\ipykernel_23416\723548108.py in <module>
               3 for i in range(3):
         ---> 4
                     models[i].fit(xtrain, ytrain)
               5
                     print(f'{models[i]} : ')
         IndexError: list index out of range
```





In [19]: print(metrics.classification_report(ytest, models[1].predict(xtest)))

	precision	recall	f1-score	support
0 1	0.68 0.79	0.72 0.75	0.70 0.77	97 132
accuracy macro avg weighted avg	0.73 0.74	0.74 0.74	0.74 0.73 0.74	229 229 229

In []: