

IMPORTING THE LIBRARIES

```
In [20]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
```

READ THE DATASET

```
In [21]: df=pd.read_csv('winequality.csv')
```

DATA ANALYSIS

```
In [22]: df.head()
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	white	7.0	0.27	0.36	20.7	0.045	45.0	170.0	1.00100	3.00	0.45	8.8	6
1	white	8.3	0.30	0.34	1.6	0.049	14.0	132.0	0.99400	3.30	0.49	9.5	6
2	white	8.1	0.28	0.40	6.9	0.050	30.0	97.0	0.9961	3.26	0.44	10.1	6
3	white	7.2	0.23	0.32	8.5	0.058	47.0	186.0	0.9956	3.19	0.40	9.9	6
4	white	7.2	0.23	0.32	8.5	0.058	47.0	186.0	0.9956	3.19	0.40	9.9	6

```
In [23]: df.tail()
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
6482	red	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	10.5	5
6483	red	5.9	0.550	0.10	2.2	0.082	38.0	51.0	0.99612	3.52	0.58	11.2	6
6484	red	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99674	3.42	0.75	11.0	6
6485	red	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99647	3.57	0.71	10.2	5
6486	red	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6

```
In [24]: df.columns
```

```
In [24]: index(['type', 'fixed acidity', 'volatile acidity', 'citric acid',
'residual sugar', 'chlorides', 'free sulfur dioxide',
'total sulfur dioxide', 'density', 'pH', 'sulphates', 'alcohol',
'quality'],
dtype='object')
```

```
In [25]: df.shape
```

```
In [25]: (6497, 13)
```

```
In [26]: df.describe()
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
count	6497.000000	6497.000000	6494.000000	6496.000000	6496.000000	6497.000000	6497.000000	6497.000000	6496.000000	6497.000000	6497.000000	6497.000000
mean	7.265079	0.330691	0.318722	5.443226	0.050642	30.525319	115.744674	0.994687	3.218395	0.533215	10.148101	5.818378
std	1.296750	0.146449	0.148265	4.768125	0.030266	17.749400	66.621855	0.002999	0.160748	0.148814	1.192712	0.872265
min	3.600000	0.000000	0.000000	0.600000	0.000000	17.000000	6.000000	0.987110	2.720000	0.220000	8.000000	3.000000
25%	6.400000	0.230000	0.250000	1.800000	0.038000	17.000000	77.000000	0.992340	3.110000	0.430000	9.500000	5.000000
50%	7.000000	0.280000	0.310000	3.000000	0.047000	29.000000	118.000000	0.994880	3.210000	0.510000	10.300000	6.000000
75%	7.700000	0.400000	0.390000	8.100000	0.060000	41.000000	156.000000	0.996990	3.320000	0.600000	11.300000	6.000000
max	15.900000	1.800000	1.600000	65.800000	0.611000	289.000000	440.000000	1.038980	4.010000	2.600000	14.900000	9.000000

```
In [27]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 6497 entries, 0 to 6496
Data columns (total 13 columns):
#   Column                Non-Null Count  Dtype
---  -
0   type                  6497 non-null   object
1   fixed acidity          6497 non-null   float64
2   volatile acidity       6489 non-null   float64
3   citric acid            6494 non-null   float64
4   residual sugar         6495 non-null   float64
5   chlorides              6495 non-null   float64
6   free sulfur dioxide    6497 non-null   float64
7   total sulfur dioxide   6497 non-null   float64
8   density                6497 non-null   float64
9   pH                    6488 non-null   float64
10  sulphates              6493 non-null   float64
11  alcohol                6497 non-null   float64
12  quality                6497 non-null   int64
memory usage: 465.7+ KB
```

```
In [28]: df.isnull().sum()
```

```
type                  0
fixed acidity         0
volatile acidity      8
citric acid           3
residual sugar        2
chlorides             2
free sulfur dioxide   0
total sulfur dioxide  0
density              10
pH                   10
sulphates            10
alcohol              10
quality              10
dtype: int64
```

REPLACING THE NULL VALUES

```
In [29]: df.fillna(method='ffill',inplace=True)
```

```
In [30]: df.isnull().sum()
```

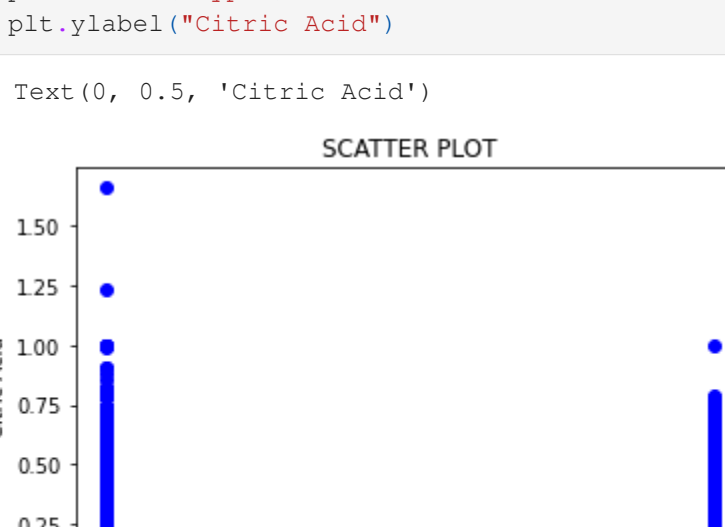
```
type                  0
fixed acidity         0
volatile acidity      0
citric acid           0
residual sugar        0
chlorides             0
free sulfur dioxide   0
total sulfur dioxide  0
density              0
pH                   0
sulphates            0
alcohol              0
quality              0
dtype: int64
```

DATA VISUALIZATION

MATPLOTLIB

```
In [35]: plt.bar(df['type'],df['fixed acidity'],color='b')
plt.title('BAR CHART')
plt.xlabel('Type')
plt.ylabel('Fixed Acidity')
```

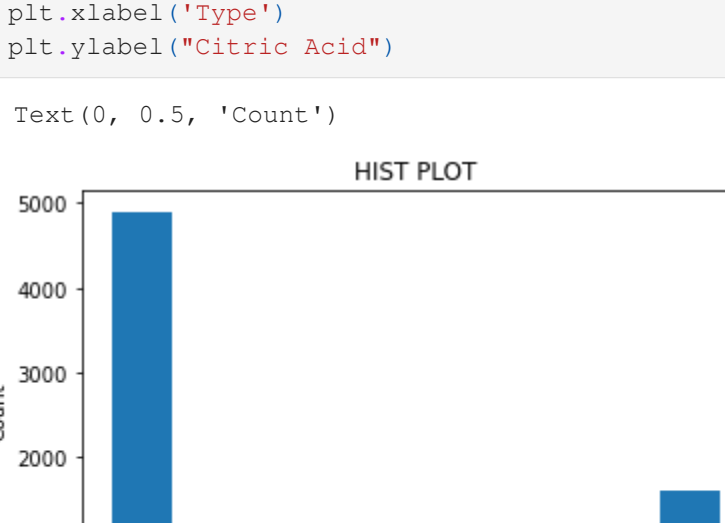
```
In [35]: Text(0, 0.5, 'Fixed Acidity')
```



```
In [36]: plt.scatter(df['type'],df['citric acid'],color='b')
```

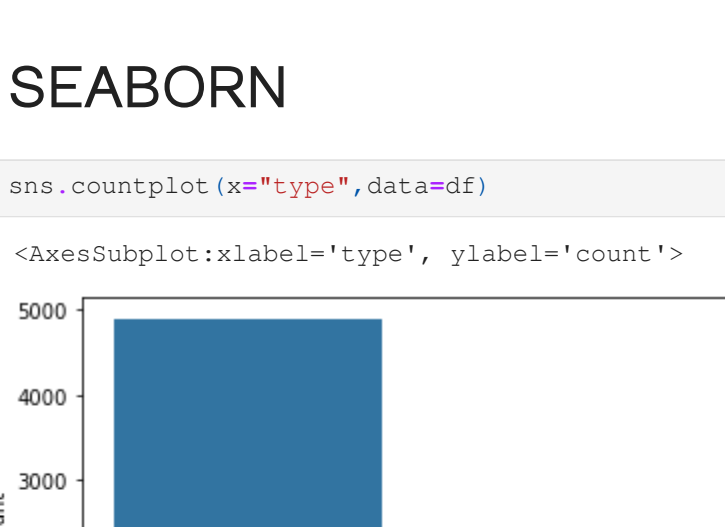
```
In [36]: plt.title('SCATTER PLOT')
plt.xlabel('Type')
plt.ylabel('Citric Acid')
```

```
In [36]: Text(0, 0.5, 'Citric Acid')
```



```
In [34]: plt.hist(df['type'])
plt.title('HIST PLOT')
plt.xlabel('Type')
plt.ylabel('Count')
```

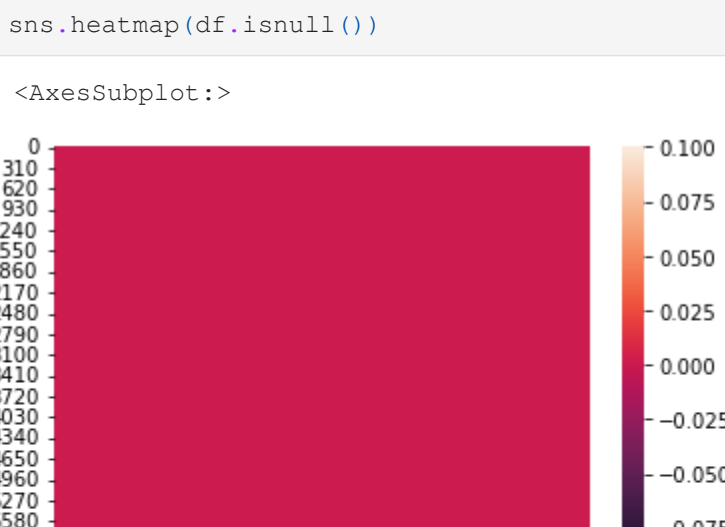
```
In [34]: Text(0, 0.5, 'Count')
```



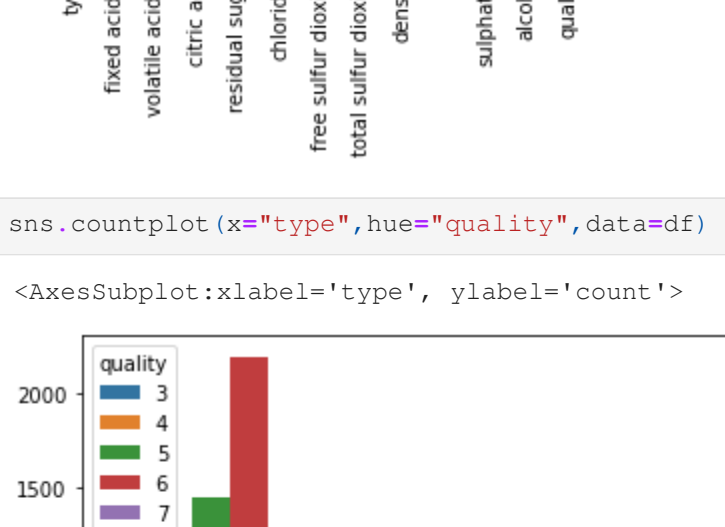
SEABORN

```
In [40]: sns.countplot(x='type',data=df)
```

```
In [40]: <seaborn.axisset.FacetGrid>
```

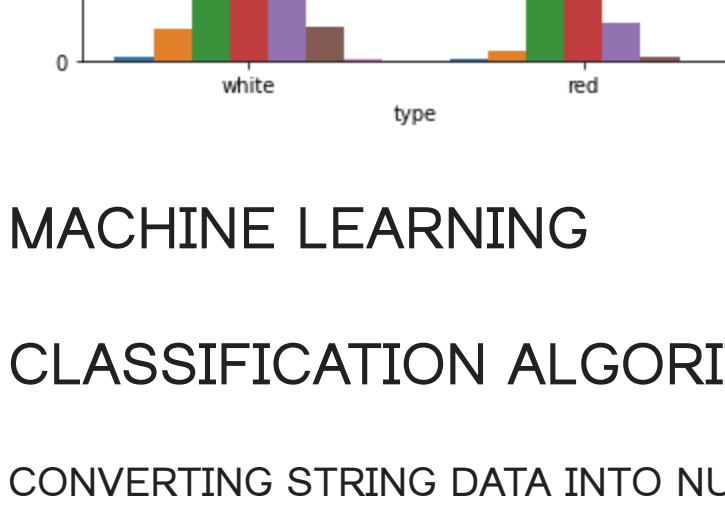


```
In [40]: sns.histplot(df.isnull(),
plt.title('HIST PLOT')
plt.xlabel('Type')
plt.ylabel('Count')
```



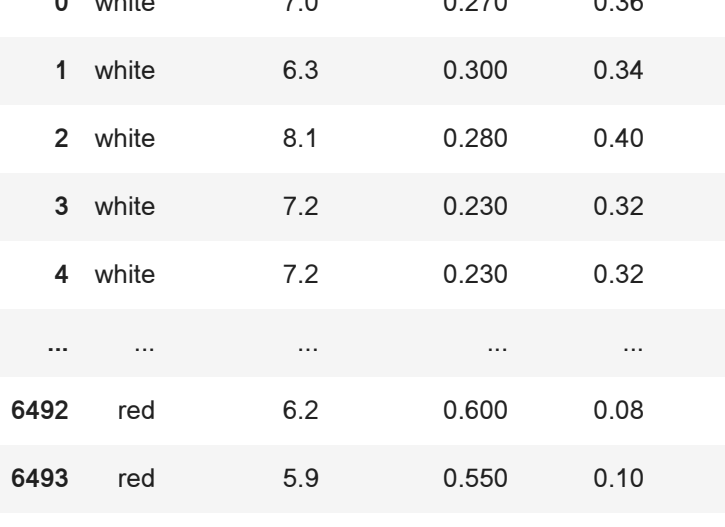
```
In [40]: sns.countplot(x='type',hue='quality',data=df)
```

```
In [40]: <seaborn.axisset.FacetGrid>
```



```
In [40]: sns.countplot(x='type',hue='quality',data=df)
```

```
In [40]: <seaborn.axisset.FacetGrid>
```



MACHINE LEARNING

CLASSIFICATION ALGORITHM

CONVERTING STRING DATA INTO NUMERIC VALUES

```
In [38]: df
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	white	7.0	0.270	0.36	20.7	0.045	45.0	170.0	1.00100	3.00	0.45	8.8	6
1	white	8.3	0.300	0.34	1.6	0.049	14.0	132.0	0.99400	3.30	0.49	9.5	6
2	white	8.1	0.280	0.40	6.9	0.050	30.0	97.0	0.99610	3.26	0.44	10.1	6
3	white	7.2	0.230	0.32	8.5	0.058	47.0	186.0	0.99560	3.19	0.40	9.9	6
4	white	7.2	0.230	0.32	8.5	0.058	47.0	186.0	0.99560	3.19	0.40	9.9	6

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
6482	red	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	10.5	5
6483	red	5.9	0.550	0.10	2.2	0.082	38.0	51.0	0.99612	3.52	0.58	11.2	6
6484	red	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99674	3.42	0.75	11.0	6
6485	red	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99647	3.57	0.71	10.2	5
6486	red	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6

```
6497 rows * 13 columns
```

```
In [39]: df
```

```
In [39]: df
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	7.0	0.270	0.36	20.7	0.045	45.0	170.0	1.00100	3.00	0.45	8.8	6
1	8.3	0.300	0.34	1.6	0.049	14.0	132.0	0.99400	3.30	0.49	9.5	6
2	8.1	0.280	0.40	6.9	0.050	30.0	97.0	0.99610	3.26	0.44	10.1	6
3	7.2	0.230	0.32	8.5	0.058	47.0	186.0	0.99560	3.19	0.40	9.9	6
4	7.2	0.230	0.32	8.5	0.058	47.0	186.0	0.99560	3.19	0.40	9.9	6

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
6482	red	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	10.5	5
6483	red	5.9	0.550	0.10	2.2	0.082	38.0	51.0	0.99612	3.52	0.58	11.2	6
6484	red	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99674	3.42	0.75	11.0	6
6485	red	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99647	3.57	0.71	10.2	5
6486	red	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6

```
6497 rows * 13 columns
```

```
In [39]: df
```

```
In [39]: df
```

6403	5.9	0.550	0.10	2.2	0.062	38.0	51.0	0.99512	3.52	0.58	11.2	6	0
6404	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99574	3.42	0.75	11.0	6	0
6405	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99547	3.57	0.71	10.2	5	0
6406	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6	0

6407 rows x 13 columns

FEATURE SCALING

```
from sklearn.preprocessing import StandardScaler
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
6482	red	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	10.5	5
6483	red	5.9	0.550	0.10	2.2	0.082	38.0	51.0	0.99612	3.52	0.58	11.2	6
6484	red	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99674	3.42	0.75	11.0	6
6485	red	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99647	3.57	0.71	10.2	5
6486	red	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	11.0	6

```
6497 rows * 13 columns
```

```
In [39]: df
```

```
In [39]: df
```

```
array([[]],
      [1],
      [1],
      [1],
      [0],
      [0],
      [0],
      [0], dtype=uint8)
```

TRAIN AND TEST

```
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_state=1)
```

LOGISTIC REGRESSION

```
from sklearn.linear_model import LogisticRegression
reg=LogisticRegression()

reg.fit(x_train,y_train)

Users\sharath\anaconda3\lib\site-packages\sklearn\utils\validation.py:955: DataConversionWarning: A column-vector y was passed when you need a 1D array, the column was ignored. Please use the keyword 'y=None' to ignore this warning.
  y = column_or_1d(y, warn=True)
Users\sharath\anaconda3\lib\site-packages\sklearn\linear_model\_logistic.py:144: ConvergenceWarning: lbfgs failed to converge (increase the number of iterations if you are sure the problem is convex)
  increase the number of iterations if you are sure the problem is convex)
  increase the number of iterations if you are sure the problem is convex)
```