

```
pip install sklearn
```

Requirement already satisfied: sklearn in c:\users\nishant\anaconda3\lib\site-packages (0.0)Note: you may need to restart the kernel to use updated packages.

Requirement already satisfied: scikit-learn in c:\users\nishant\anaconda3\lib\site-packages (from sklearn) (0.24.2)

Requirement already satisfied: scipy>=0.19.1 in c:\users\nishant\anaconda3\lib\site-packages (from scikit-learn->sklearn) (1.7.1)

Requirement already satisfied: threadpoolctl>=2.0.0 in c:\users\nishant\anaconda3\lib\site-packages (from scikit-learn->sklearn) (2.2.0)

Requirement already satisfied: joblib>=0.11 in c:\users\nishant\anaconda3\lib\site-packages (from scikit-learn->sklearn) (1.1.0)

Requirement already satisfied: numpy>=1.13.3 in c:\users\nishant\anaconda3\lib\site-packages (from scikit-learn->sklearn) (1.20.3)

```
import numpy as np
import pandas as pd
import seaborn as sb
import matplotlib.pyplot as plt
import sklearn as sk
from sklearn.model_selection import train_test_split

data= pd.read_csv("C:\Users\Nishant\Downloads\QualityPrediction.csv")
data.head()
```

	fixed acidity	volatile acidity	citric acid	residual sugar
0	7.4	0.70	0.00	1.9
1	7.8	0.88	0.00	2.6
2	7.8	0.76	0.04	2.3
3	11.2	0.28	0.56	1.9
4	7.4	0.70	0.00	1.9

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
0	11.0	34.0	0.9978	3.51	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

4	11.0	34.0	0.9978	3.51	0.56
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	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5

data.describe()

	fixed acidity	volatile acidity	citric acid	residual sugar \
count	1599.000000	1599.000000	1599.000000	1599.000000
mean	8.319637	0.527821	0.270976	2.538806
std	1.741096	0.179060	0.194801	1.409928
min	4.600000	0.120000	0.000000	0.900000
25%	7.100000	0.390000	0.090000	1.900000
50%	7.900000	0.520000	0.260000	2.200000
75%	9.200000	0.640000	0.420000	2.600000
max	15.900000	1.580000	1.000000	15.500000

	chlorides	free sulfur dioxide	total sulfur dioxide
density \			
count	1599.000000	1599.000000	1599.000000
1599.000000			
mean	0.087467	15.874922	46.467792
0.996747			
std	0.047065	10.460157	32.895324
0.001887			
min	0.012000	1.000000	6.000000
0.990070			
25%	0.070000	7.000000	22.000000
0.995600			
50%	0.079000	14.000000	38.000000
0.996750			
75%	0.090000	21.000000	62.000000
0.997835			
max	0.611000	72.000000	289.000000
1.003690			

	pH	sulphates	alcohol	quality
count	1599.000000	1599.000000	1599.000000	1599.000000
mean	3.311113	0.658149	10.422983	5.636023
std	0.154386	0.169507	1.065668	0.807569
min	2.740000	0.330000	8.400000	3.000000
25%	3.210000	0.550000	9.500000	5.000000
50%	3.310000	0.620000	10.200000	6.000000

75%	3.400000	0.730000	11.100000	6.000000
max	4.010000	2.000000	14.900000	8.000000

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

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

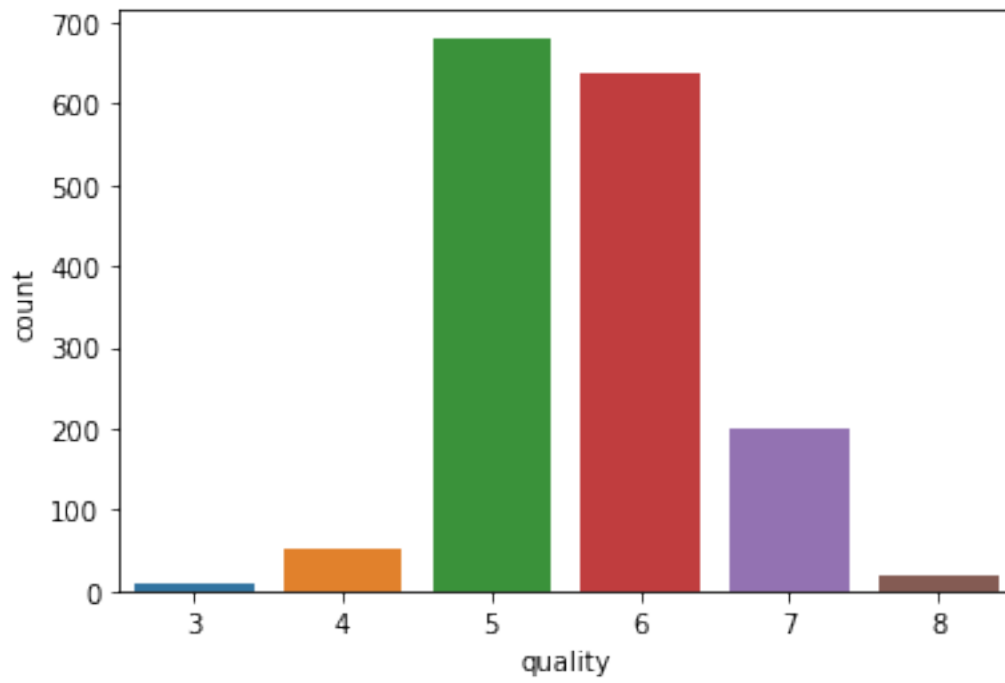
```
data['quality'].value_counts()
```

5	681
6	638
7	199
4	53
8	18
3	10

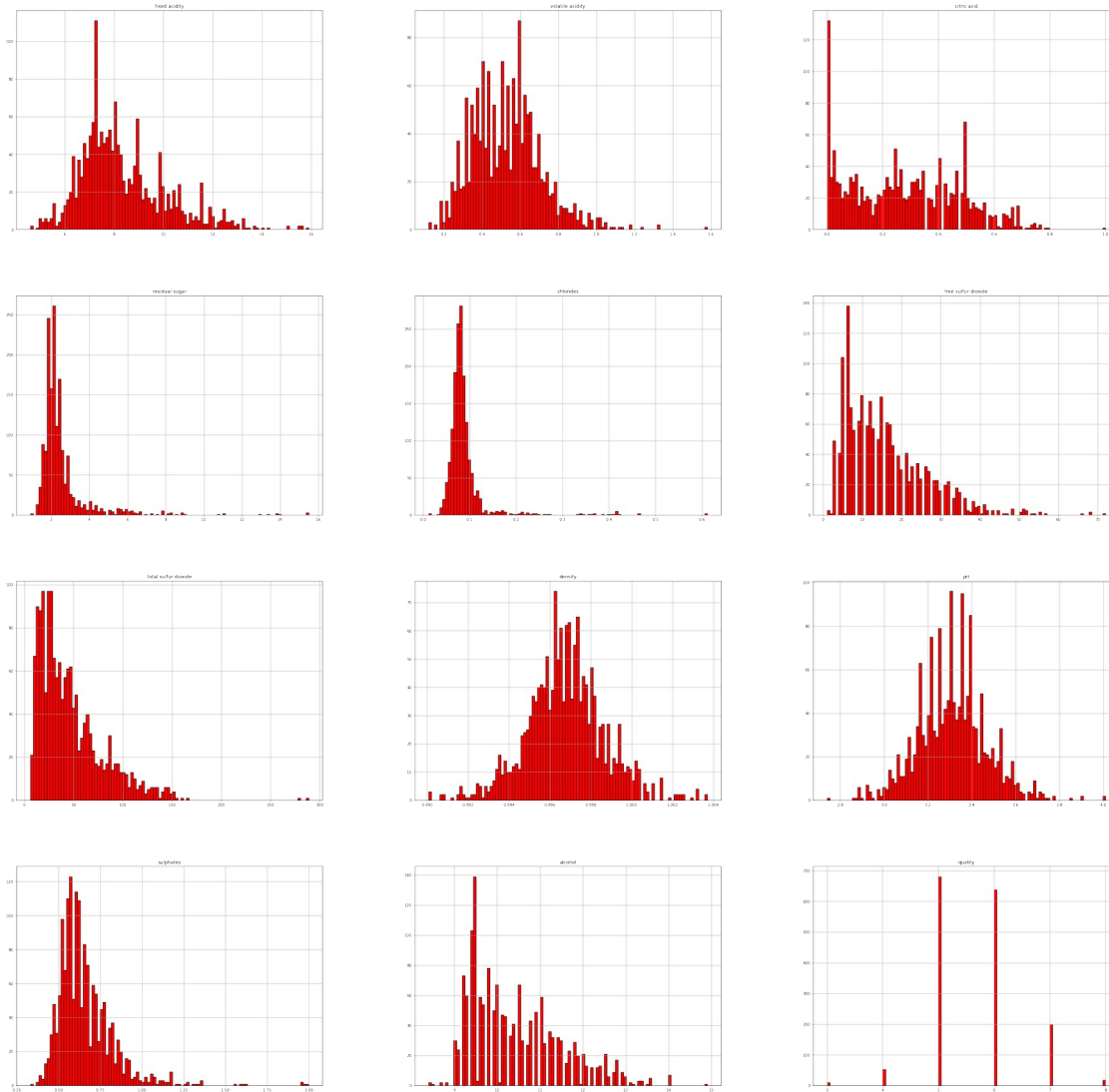
Name: quality, dtype: int64

```
sb.countplot(data['quality'])  
xlabel='quality'  
ylabel='count'  
plt.savefig('bar')
```

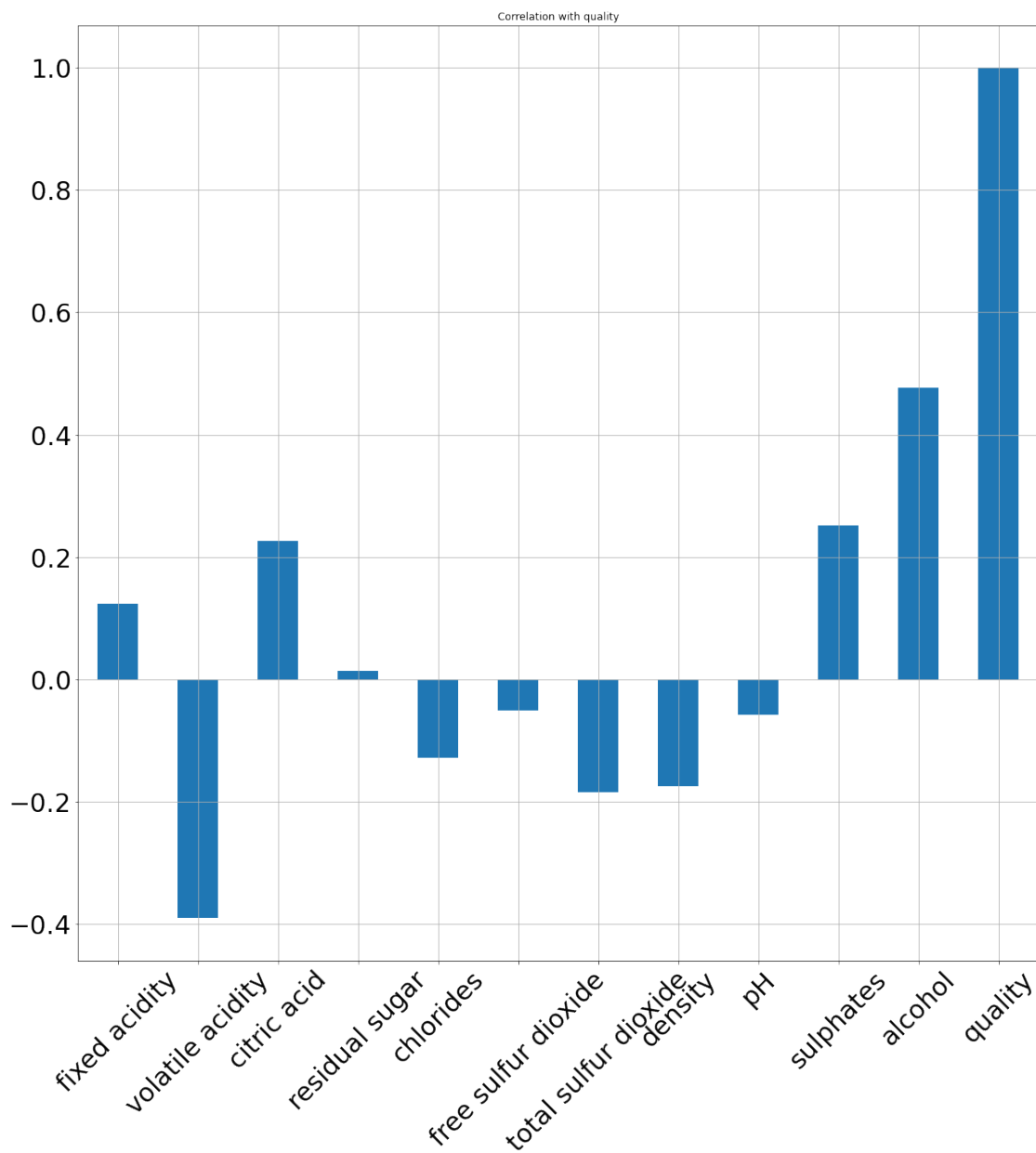
C:\Users\Nishant\anaconda3\lib\site-packages\seaborn\
_decorators.py:36: FutureWarning: Pass the following variable as a
keyword arg: x. From version 0.12, the only valid positional argument
will be `data`, and passing other arguments without an explicit
keyword will result in an error or misinterpretation.
warnings.warn(



```
data.hist(bins=100, figsize=(50,50),color='red',edgecolor='black')  
plt.show()
```



```
data.corrwith(data.quality).plot.bar(
    figsize=(20,20),title='Correlation with
quality',fontsize='30',rot='45',grid=True)
plt.savefig('same1')
```



```
sb.set(style='white')
corr=data.corr()
```

```
corr.head
```

```
<bound method NDFrame.head of
volatile acidity  citric acid \
fixed acidity    1.000000    -0.256131    0.671703
volatile acidity -0.256131    1.000000   -0.552496
citric acid      0.671703   -0.552496    1.000000
residual sugar   0.114777    0.001918    0.143577
chlorides        0.093705    0.061298    0.203823
free sulfur dioxide -0.153794 -0.010504   -0.060978
```

total sulfur dioxide	-0.113181	0.076470	0.035533
density	0.668047	0.022026	0.364947
pH	-0.682978	0.234937	-0.541904
sulphates	0.183006	-0.260987	0.312770
alcohol	-0.061668	-0.202288	0.109903
quality	0.124052	-0.390558	0.226373

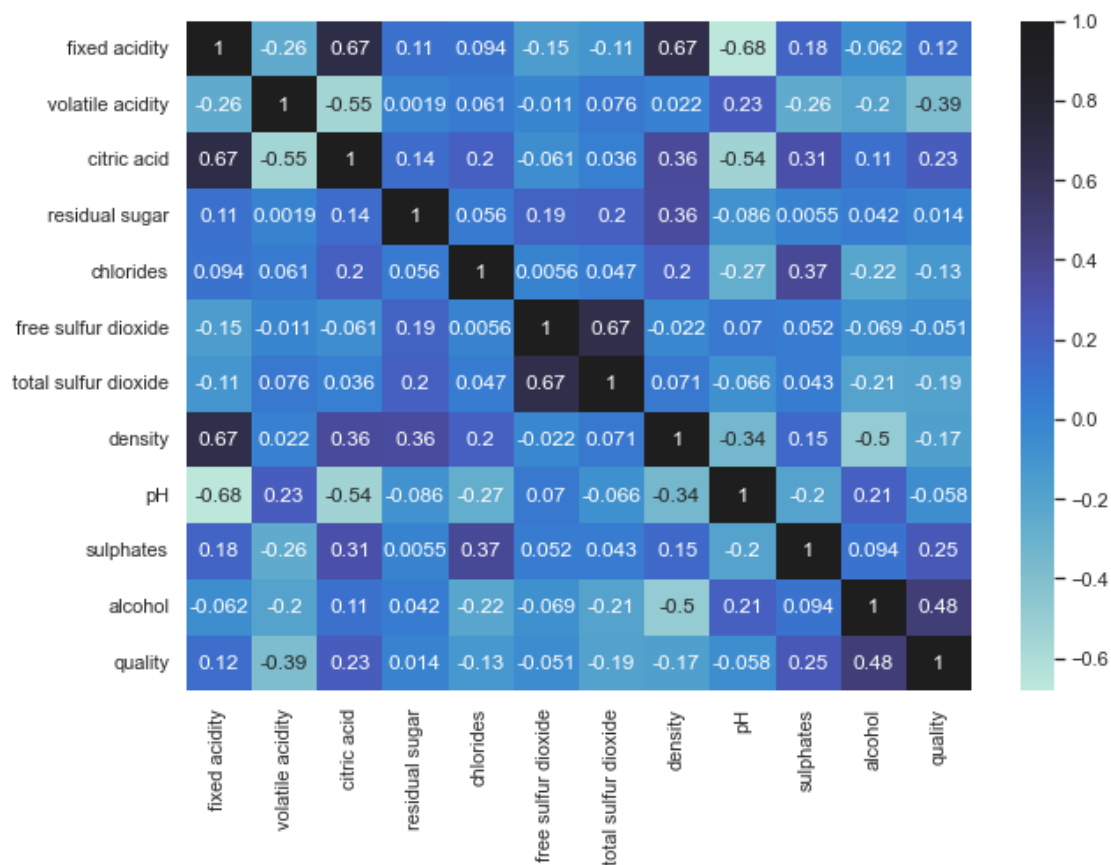
	residual sugar	chlorides	free sulfur
dioxide \			
fixed acidity	0.114777	0.093705	-0.153794
volatile acidity	0.001918	0.061298	-0.010504
citric acid	0.143577	0.203823	-0.060978
residual sugar	1.000000	0.055610	0.187049
chlorides	0.055610	1.000000	0.005562
free sulfur dioxide	0.187049	0.005562	1.000000
total sulfur dioxide	0.203028	0.047400	0.667666
density	0.355283	0.200632	-0.021946
pH	-0.085652	-0.265026	0.070377
sulphates	0.005527	0.371260	0.051658
alcohol	0.042075	-0.221141	-0.069408
quality	0.013732	-0.128907	-0.050656

	total sulfur dioxide	density	pH
sulphates \			
fixed acidity	-0.113181	0.668047	-0.682978
0.183006			
volatile acidity	0.076470	0.022026	0.234937
0.260987			-
citric acid	0.035533	0.364947	-0.541904
0.312770			
residual sugar	0.203028	0.355283	-0.085652
0.005527			
chlorides	0.047400	0.200632	-0.265026
0.371260			
free sulfur dioxide	0.667666	-0.021946	0.070377
0.051658			
total sulfur dioxide	1.000000	0.071269	-0.066495

0.042947				
density	0.071269	1.000000	-0.341699	
0.148506				
pH	-0.066495	-0.341699	1.000000	-
0.196648				
sulphates	0.042947	0.148506	-0.196648	
1.000000				
alcohol	-0.205654	-0.496180	0.205633	
0.093595				
quality	-0.185100	-0.174919	-0.057731	
0.251397				

	alcohol	quality
fixed acidity	-0.061668	0.124052
volatile acidity	-0.202288	-0.390558
citric acid	0.109903	0.226373
residual sugar	0.042075	0.013732
chlorides	-0.221141	-0.128907
free sulfur dioxide	-0.069408	-0.050656
total sulfur dioxide	-0.205654	-0.185100
density	-0.496180	-0.174919
pH	0.205633	-0.057731
sulphates	0.093595	0.251397
alcohol	1.000000	0.476166
quality	0.476166	1.000000

```
plt.figure(figsize=(10,7))
sb.heatmap(data.corr(), annot=True,center=1)
plt.title('Correlation of Data',size=20,color='white')
plt.savefig('same')
```

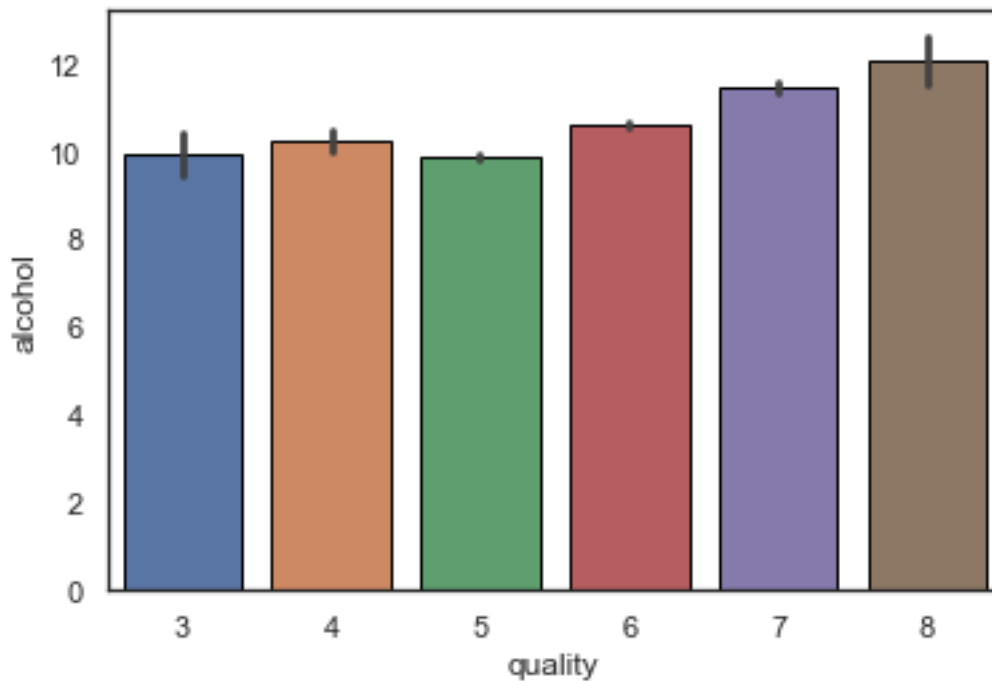
```
data.corr()['quality'].sort_values()
```

```
volatile acidity      -0.390558
total sulfur dioxide  -0.185100
density              -0.174919
chlorides            -0.128907
pH                  -0.057731
free sulfur dioxide  -0.050656
residual sugar        0.013732
fixed acidity         0.124052
citric acid           0.226373
sulphates             0.251397
alcohol              0.476166
quality               1.000000
Name: quality, dtype: float64
```

```
sb.barplot(data['quality'],data['alcohol'],edgecolor='black')
plt.savefig('sample')
```

C:\Users\Nishant\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an

explicit keyword will result in an error or misinterpretation.
 warnings.warn(



```
data['quality']= data.quality.apply(lambda x:1if x>=7 else 0)
```

```
data['quality'].value_counts()
```

```
0    1382
```

```
1     217
```

```
Name: quality, dtype: int64
```

```
x=data.drop('quality',axis=1)
```

```
y=data['quality']
```

```
x.head()
```

	fixed acidity	volatile acidity	citric acid	residual sugar
0	7.4	0.70	0.00	1.9
1	7.8	0.88	0.00	2.6
2	7.8	0.76	0.04	2.3
3	11.2	0.28	0.56	1.9
4	7.4	0.70	0.00	1.9

free sulfur dioxide	total sulfur dioxide	density	pH	sulphates
---------------------	----------------------	---------	----	-----------

\					
0	11.0	34.0	0.9978	3.51	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
4	11.0	34.0	0.9978	3.51	0.56

```

    alcohol
0      9.4
1      9.8
2      9.8
3      9.8
4      9.4

```

```
y.head()
```

```

0      0
1      0
2      0
3      0
4      0

```

```
Name: quality, dtype: int64
```

```

x_train,x_test,y_train,y_test =
train_test_split(x,y,test_size=0.3,random_state=0)

```

```

print('x_train',x_train.shape)
print('y_train',y_train.shape)
print('x_test',x_test.shape)
print('y_test',y_test.shape)

```

```

x_train (1119, 11)
y_train (1119,)
x_test (480, 11)
y_test (480,)

```

```

from sklearn.metrics import accuracy_score
from sklearn.metrics import classification_report
from sklearn import tree

```

```

decision_tree = tree.DecisionTreeClassifier()
decision_tree.fit(x_train,y_train)
decision_tree_pred = decision_tree.predict(x_test)
decision_tree_acc= accuracy_score(y_test,decision_tree_pred)
print("Test accuracy: {:.2f}%".format(decision_tree_acc*100))

```

Test accuracy: 88.333333%

```
print(classification_report(y_test, decision_tree_pred))
plt.savefig('CM1')
```

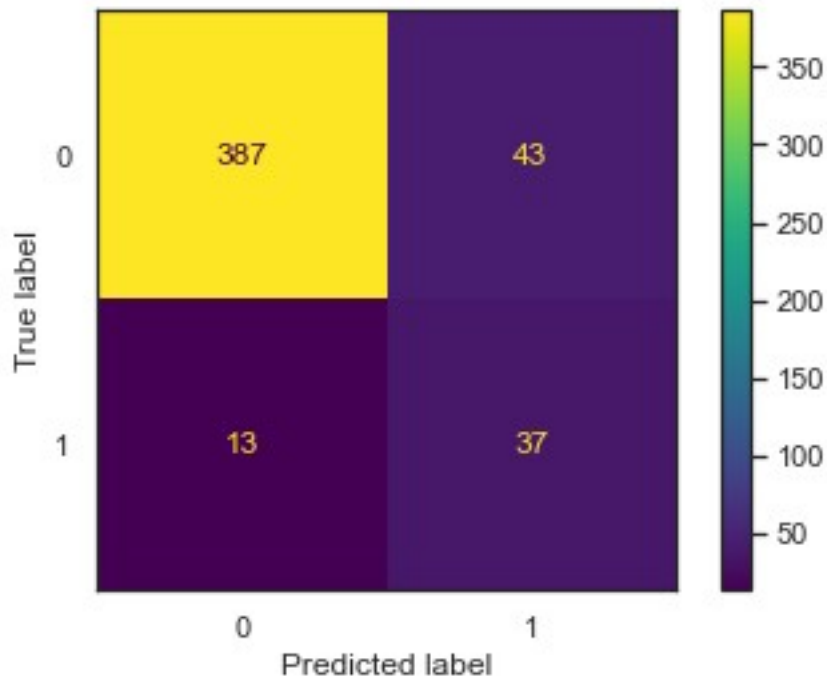
	precision	recall	f1-score	support
0	0.97	0.90	0.93	430
1	0.46	0.74	0.57	50
accuracy			0.88	480
macro avg	0.72	0.82	0.75	480
weighted avg	0.91	0.88	0.89	480

<Figure size 432x288 with 0 Axes>

```
from sklearn.metrics import confusion_matrix
import random
import itertools
from sklearn.metrics import ConfusionMatrixDisplay

cm = confusion_matrix(y_test, decision_tree_pred)
disp= ConfusionMatrixDisplay(confusion_matrix=cm)
disp.plot()
print('True Negative: ', cm[0][0])
print('False Negative: ', cm[1][0])
print('True Positive: ', cm[1][1])
print('False Positive: ', cm[0][1])
plt.savefig('CM')
```

True Negative: 387
False Negative: 13
True Positive: 37
False Positive: 43



```
from sklearn.ensemble import RandomForestClassifier
randforest=RandomForestClassifier()
randforest.fit(x_train, y_train)
randforest_pred=randforest.predict(x_test)
randforest_acc=accuracy_score(randforest_pred,y_test)
print('Test accuracy:{:.2f}%'.format(randforest_acc*100))
```

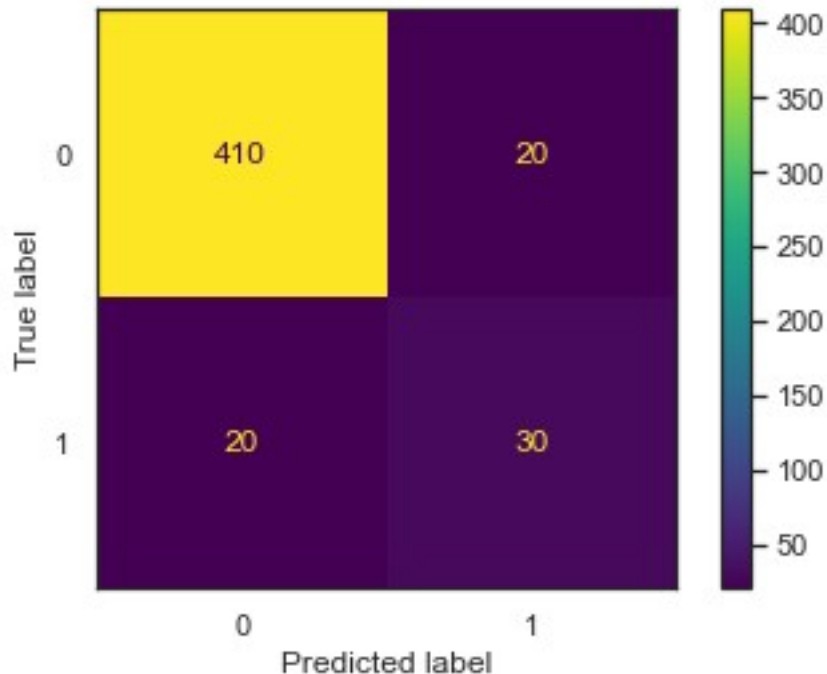
Test accuracy:91.67%

```
print(classification_report(y_test, randforest_pred))
```

	precision	recall	f1-score	support
0	0.95	0.95	0.95	430
1	0.60	0.60	0.60	50
accuracy			0.92	480
macro avg	0.78	0.78	0.78	480
weighted avg	0.92	0.92	0.92	480

```
cm = confusion_matrix(y_test,randforest_pred)
disp= ConfusionMatrixDisplay(confusion_matrix=cm)
disp.plot()
print('True Negative: ',cm[0][0])
print('False Negative: ',cm[1][0])
print('True Positive: ',cm[1][1])
print('False Positive: ',cm[0][1])
```

True Negative: 410
False Negative: 20
True Positive: 30
False Positive: 20

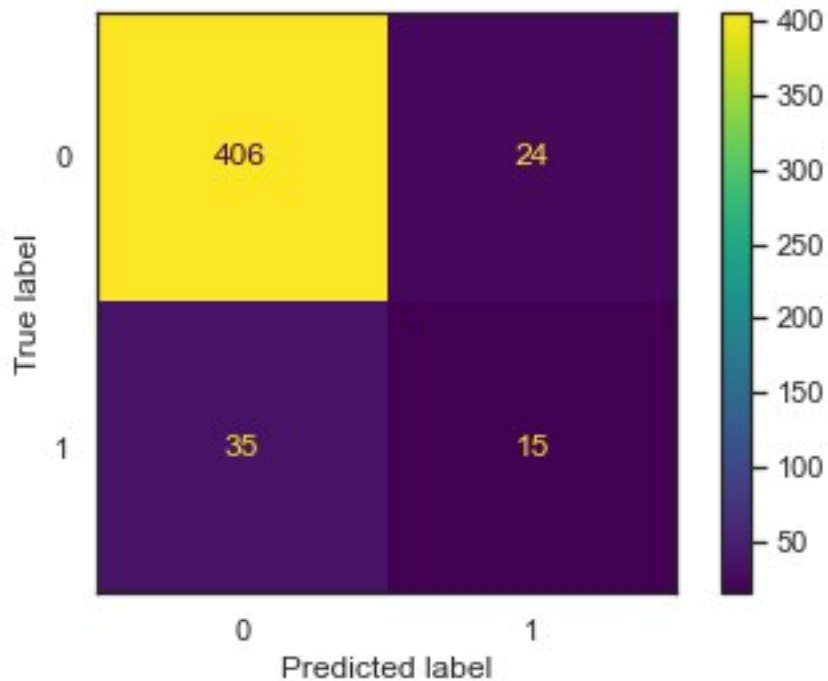


```
from sklearn import neighbors
from sklearn.metrics import mean_squared_error
from sklearn.neighbors import KNeighborsClassifier
kclassifier=KNeighborsClassifier()
kclassifier.fit(x_train, y_train)
kclassifier_pred=kclassifier.predict(x_test)
kclassifier_acc=accuracy_score(kclassifier_pred,y_test)
print('Test accuracy:{:.2f}%'.format(kclassifier_acc*100))
```

Test accuracy:87.71%

```
cm = confusion_matrix(y_test,kclassifier_pred)
disp= ConfusionMatrixDisplay(confusion_matrix=cm)
disp.plot()
print('True Negative: ',cm[0][0])
print('False Negative: ',cm[1][0])
print('True Positive: ',cm[1][1])
print('False Positive: ',cm[0][1])
```

True Negative: 406
False Negative: 35
True Positive: 15
False Positive: 24



```

error = []
for i in range(1, 100):
    knn = KNeighborsClassifier(n_neighbors=i)
    knn.fit(x_train,y_train)
    pred_i = knn.predict(x_test)
    error.append(np.mean(pred_i != y_test))

plt.figure(figsize=(12, 6))
plt.plot(range(1,100), error, color='red', linestyle='dashed',
marker='o',
        markerfacecolor='blue', markersize=10)
plt.title('Error Rate K Value')
plt.xlabel('K Value')
plt.ylabel('Mean Error')

Text(0, 0.5, 'Mean Error')

```

