

In [1]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.naive_bayes import MultinomialNB
from sklearn.neighbors import KNeighborsClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
from sklearn.ensemble import AdaBoostClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import warnings
warnings.filterwarnings('ignore')
```

In [2]:

```
wine=pd.read_csv('wine.csv')
```

In [3]:

```
wine.head()
```

Out[3]:

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins	Color intensity	Hue	OD280/OD315 of diluted wines
0	1	14.23	1.71	2.43	15.6	127	2.80	3.06	0.28	2.29	5.64	1.04	3
1	1	13.20	1.78	2.14	11.2	100	2.65	2.76	0.26	1.28	4.38	1.05	3
2	1	13.16	2.36	2.67	18.6	101	2.80	3.24	0.30	2.81	5.68	1.03	3
3	1	14.37	1.95	2.50	16.8	113	3.85	3.49	0.24	2.18	7.80	0.86	3
4	1	13.24	2.59	2.87	21.0	118	2.80	2.69	0.39	1.82	4.32	1.04	2

In [4]:

```
wine.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 178 entries, 0 to 177
Data columns (total 14 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   Class                                178 non-null    int64
1   Alcohol                             178 non-null    float64
2   Malic acid                           178 non-null    float64
3   Ash                                  178 non-null    float64
4   Alcalinity of ash                    178 non-null    float64
5   Magnesium                            178 non-null    int64
6   Total phenols                        178 non-null    float64
7   Flavanoids                           178 non-null    float64
8   Nonflavanoid phenols                 178 non-null    float64
9   Proanthocyanins                      178 non-null    float64
10  Color intensity                       178 non-null    float64
11  Hue                                   178 non-null    float64
12  OD280/OD315 of diluted wines         178 non-null    float64
13  Proline                              178 non-null    int64
dtypes: float64(11), int64(3)
memory usage: 19.6 KB
```

In [5]:

```
wine.shape
```

Out[5]:

(178, 14)

In [6]:

```
wine.dtypes
```

Out[6]:

```
Class                int64
Alcohol              float64
Malic acid           float64
Ash                  float64
Alcalinity of ash    float64
Magnesium            int64
Total phenols        float64
Flavanoids           float64
Nonflavanoid phenols float64
Proanthocyanins       float64
Color intensity      float64
Hue                  float64
OD280/OD315 of diluted wines float64
Proline              int64
dtype: object
```

In [7]:

```
wine.columns
```

Out[7]:

```
Index(['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of ash',
      'Magnesium', 'Total phenols', 'Flavanoids', 'Nonflavanoid phenols',
      'Proanthocyanins', 'Color intensity', 'Hue',
      'OD280/OD315 of diluted wines', 'Proline'],
      dtype='object')
```

In [8]:

```
wine.describe()
```

Out[8]:

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanin
count	178.000000	178.000000	178.000000	178.000000	178.000000	178.000000	178.000000	178.000000	178.000000	178.000000
mean	1.938202	13.000618	2.336348	2.366517	19.494944	99.741573	2.295112	2.029270	0.361854	1.59089
std	0.775035	0.811827	1.117146	0.274344	3.339564	14.282484	0.625851	0.998859	0.124453	0.57235
min	1.000000	11.030000	0.740000	1.360000	10.600000	70.000000	0.980000	0.340000	0.130000	0.41000
25%	1.000000	12.362500	1.602500	2.210000	17.200000	88.000000	1.742500	1.205000	0.270000	1.25000
50%	2.000000	13.050000	1.865000	2.360000	19.500000	98.000000	2.355000	2.135000	0.340000	1.55500
75%	3.000000	13.677500	3.082500	2.557500	21.500000	107.000000	2.800000	2.875000	0.437500	1.95000
max	3.000000	14.830000	5.800000	3.230000	30.000000	162.000000	3.880000	5.080000	0.660000	3.58000

In [9]:

```
#to understand it has two or more unique variables and is it regression or classification
wine.Class.unique()
```

Out[9]:

```
array([1, 2, 3], dtype=int64)
```

In [10]:

```
# identifying null values
wine.isnull().sum()
```

Out[10]:

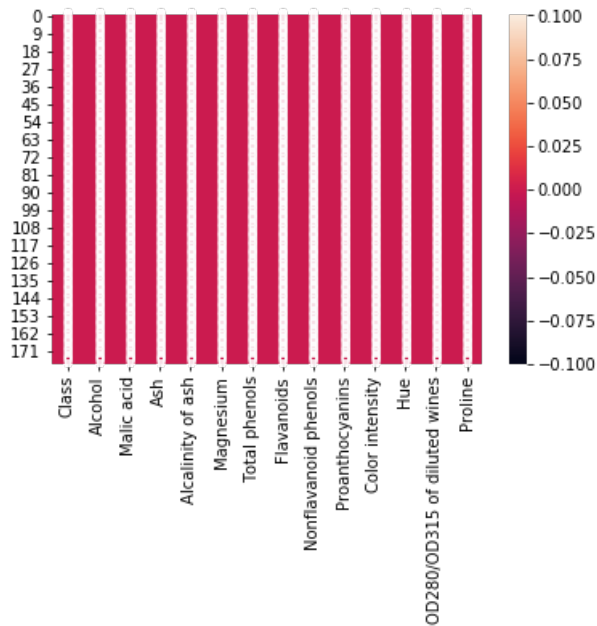
```
Class          0
Alcohol        0
Malic acid     0
Ash            0
Alcalinity of ash  0
Magnesium      0
Total phenols  0
Flavanoids     0
Nonflavanoid phenols  0
Proanthocyanins 0
Color intensity 0
Hue            0
OD280/OD315 of diluted wines 0
Proline        0
dtype: int64
```

In [11]:

```
sns.heatmap(wine.isnull(),annot=True)
```

Out[11]:

<matplotlib.axes._subplots.AxesSubplot at 0x22f607450c8>



In [12]:

```
wine.corr()
```

Out[12]:

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins
Class	1.000000	0.328222	0.437776	0.049643	0.517859	-0.209179	0.719163	-0.847498	0.489109	-0.499130
Alcohol	0.328222	1.000000	0.094397	0.211545	0.310235	0.270798	0.289101	0.236815	-0.155929	0.136698
Malic acid	0.437776	0.094397	1.000000	0.164045	0.288500	-0.054575	0.335167	-0.411007	0.292977	-0.220746
Ash	0.049643	0.211545	0.164045	1.000000	0.443367	0.286587	0.128980	0.115077	0.186230	0.009652

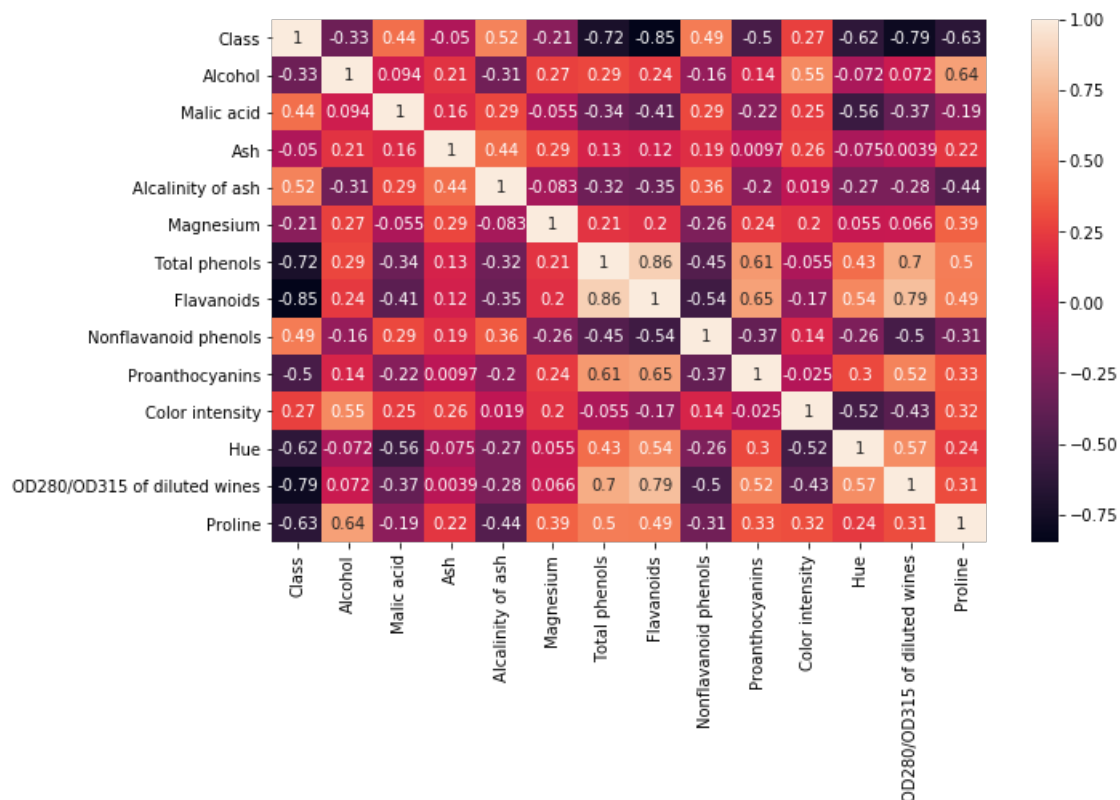
	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins
Alcalinity of ash	0.517893	0.310235	0.288569	0.443369	1.009000	0.068608	0.351178	0.351178	0.361922	0.197022
Magnesium	0.209179	0.270798	0.054575	0.286587	0.083333	1.000000	0.214401	0.195784	-0.256294	0.236441
Total phenols	0.719163	0.289101	0.335167	0.128980	0.321113	0.214401	1.000000	0.864564	-0.449935	0.612413
Flavanoids	0.847498	0.236815	0.411007	0.115077	0.351370	0.195784	0.864564	1.000000	-0.537900	0.652692
Nonflavanoid phenols	0.489109	0.155929	0.292977	0.186230	0.361922	-0.256294	0.449935	-0.537900	1.000000	-0.365845
Proanthocyanins	0.499130	0.136698	0.220746	0.009652	0.197327	0.236441	0.612413	0.652692	-0.365845	1.000000
Color intensity	0.265668	0.546364	0.248985	0.258887	0.018732	0.199950	0.055136	-0.172379	0.139057	-0.025250
Hue	0.617369	0.071747	0.561296	0.074667	0.273955	0.055398	0.433681	0.543479	-0.262640	0.295544
OD280/OD315 of diluted wines	0.788230	0.072343	0.368710	0.003911	0.276769	0.066004	0.699949	0.787194	-0.503270	0.519067
Proline	0.633717	0.643720	0.192011	0.223626	0.440597	0.393351	0.498115	0.494193	-0.311385	0.330417

In [13]:

```
plt.figure(figsize=(10,6))
sns.heatmap(wine.corr(),annot=True)
```

Out[13]:

<matplotlib.axes._subplots.AxesSubplot at 0x22f623b8b48>



In [14]:

```
wine.skew()
```

Out[14]:

```
Class          0.107431
Alcohol        -0.051482
Malic acid     1.039651
```

```

Ash -0.176699
Alcalinity of ash 0.213047
Magnesium 1.098191
Total phenols 0.086639
Flavanoids 0.025344
Nonflavanoid phenols 0.450151
Proanthocyanins 0.517137
Color intensity 0.868585
Hue 0.021091
OD280/OD315 of diluted wines -0.307285
Proline 0.767822
dtype: float64

```

In [15]:

```

for col in wine.columns:
    if wine.skew().loc[col]>0.55:
        wine[col]=np.log1p(wine[col])

```

In [16]:

```
wine.skew()
```

Out[16]:

```

Class 0.107431
Alcohol -0.051482
Malic acid 0.529222
Ash -0.176699
Alcalinity of ash 0.213047
Magnesium 0.605723
Total phenols 0.086639
Flavanoids 0.025344
Nonflavanoid phenols 0.450151
Proanthocyanins 0.517137
Color intensity 0.097222
Hue 0.021091
OD280/OD315 of diluted wines -0.307285
Proline 0.087930
dtype: float64

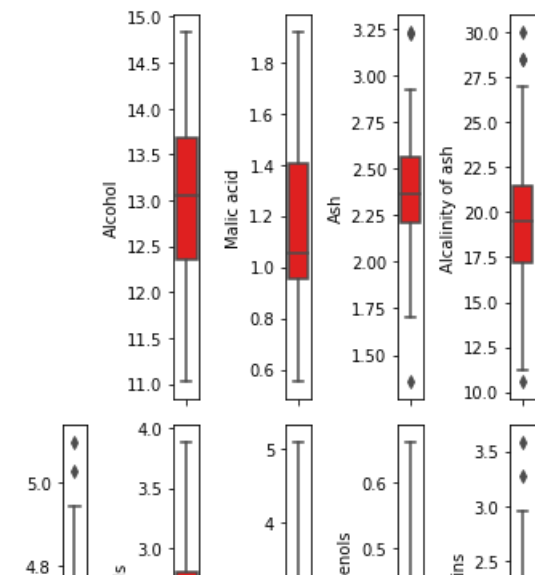
```

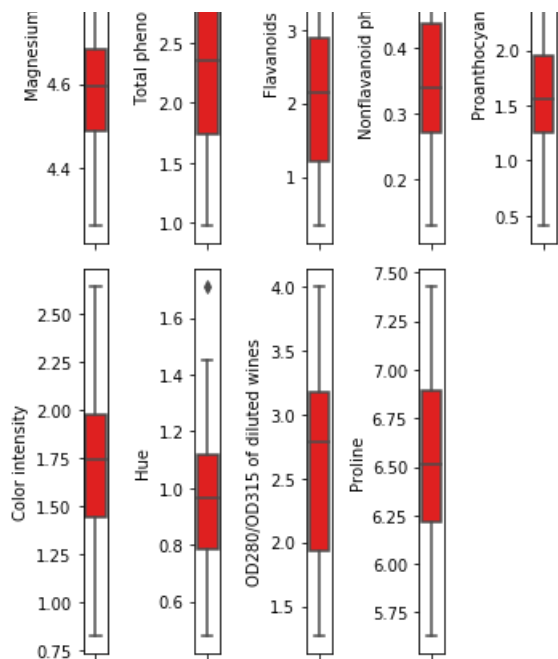
In [17]:

```

col=wine.columns.values
ncol=5
nrow=7
plt.figure(figsize=(ncol,5*ncol))
for i in range(1,len(col)):
    plt.subplot(nrow,ncol,i+1)
    sns.boxplot(wine[col[i]],color='red',orient='v')
    plt.tight_layout()

```





In [18]:

```
#Removing outliers
from scipy.stats import zscore
z_score=abs(zscore(wine))
print(wine.shape)
wine_df=wine.loc[(z_score<3).all(axis=1)]
print(wine_df.shape)
```

(178, 14)
(170, 14)

In [19]:

```
wine_df
```

Out[19]:

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavonoids	Nonflavanoid phenols	Proanthocyanins	Color intensity	Hue	OD280/OD315 of diluted wines
0	1	14.23	0.996949	2.43	15.6	4.852030	2.80	3.06	0.28	2.29	1.893112	1.04	
1	1	13.20	1.022451	2.14	11.2	4.615121	2.65	2.76	0.26	1.28	1.682688	1.05	
2	1	13.16	1.211941	2.67	18.6	4.624973	2.80	3.24	0.30	2.81	1.899118	1.03	
3	1	14.37	1.081805	2.50	16.8	4.736198	3.85	3.49	0.24	2.18	2.174752	0.86	
4	1	13.24	1.278152	2.87	21.0	4.779123	2.80	2.69	0.39	1.82	1.671473	1.04	
...	
173	3	13.71	1.894617	2.45	20.5	4.564348	1.68	0.61	0.52	1.06	2.163323	0.64	
174	3	13.40	1.591274	2.48	23.0	4.634729	1.80	0.75	0.43	1.41	2.116256	0.70	
175	3	13.27	1.663926	2.26	20.0	4.795791	1.59	0.69	0.43	1.35	2.415914	0.59	
176	3	13.17	1.278152	2.37	20.0	4.795791	1.65	0.68	0.53	1.46	2.332144	0.60	
177	3	14.13	1.629241	2.74	24.5	4.574711	2.05	0.76	0.56	1.35	2.322388	0.61	

170 rows × 14 columns

In [20]:

```
wine_df=pd.DataFrame(data=wine_df)
```

In [21]:

```
wine_df
```

Out [21]:

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins	Color intensity	Hue	OD280/OD310
0	1	14.23	0.996949	2.43	15.6	4.852030	2.80	3.06	0.28	2.29	1.893112	1.04	3.9
1	1	13.20	1.022451	2.14	11.2	4.615121	2.65	2.76	0.26	1.28	1.682688	1.05	3.4
2	1	13.16	1.211941	2.67	18.6	4.624973	2.80	3.24	0.30	2.81	1.899118	1.03	3.1
3	1	14.37	1.081805	2.50	16.8	4.736198	3.85	3.49	0.24	2.18	2.174752	0.86	3.4
4	1	13.24	1.278152	2.87	21.0	4.779123	2.80	2.69	0.39	1.82	1.671473	1.04	2.9
...
173	3	13.71	1.894617	2.45	20.5	4.564348	1.68	0.61	0.52	1.06	2.163323	0.64	1.7
174	3	13.40	1.591274	2.48	23.0	4.634729	1.80	0.75	0.43	1.41	2.116256	0.70	1.5
175	3	13.27	1.663926	2.26	20.0	4.795791	1.59	0.69	0.43	1.35	2.415914	0.59	1.5
176	3	13.17	1.278152	2.37	20.0	4.795791	1.65	0.68	0.53	1.46	2.332144	0.60	1.6
177	3	14.13	1.629241	2.74	24.5	4.574711	2.05	0.76	0.56	1.35	2.322388	0.61	1.6

170 rows × 14 columns

In [22]:

```
#x and y values allocation for training and testing
x=wine_df.iloc[:,1:-1]
```

In [23]:

```
x
```

Out [23]:

	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins	Color intensity	Hue	OD280/OD310 of dilute wine
0	14.23	0.996949	2.43	15.6	4.852030	2.80	3.06	0.28	2.29	1.893112	1.04	3.9
1	13.20	1.022451	2.14	11.2	4.615121	2.65	2.76	0.26	1.28	1.682688	1.05	3.4
2	13.16	1.211941	2.67	18.6	4.624973	2.80	3.24	0.30	2.81	1.899118	1.03	3.1
3	14.37	1.081805	2.50	16.8	4.736198	3.85	3.49	0.24	2.18	2.174752	0.86	3.4
4	13.24	1.278152	2.87	21.0	4.779123	2.80	2.69	0.39	1.82	1.671473	1.04	2.9
...
173	13.71	1.894617	2.45	20.5	4.564348	1.68	0.61	0.52	1.06	2.163323	0.64	1.7
174	13.40	1.591274	2.48	23.0	4.634729	1.80	0.75	0.43	1.41	2.116256	0.70	1.5
175	13.27	1.663926	2.26	20.0	4.795791	1.59	0.69	0.43	1.35	2.415914	0.59	1.5
176	13.17	1.278152	2.37	20.0	4.795791	1.65	0.68	0.53	1.46	2.332144	0.60	1.6
177	14.13	1.629241	2.74	24.5	4.574711	2.05	0.76	0.56	1.35	2.322388	0.61	1.6

170 rows × 12 columns

In [24]:

```
x.shape
```

Out [24]:

```
(170, 12)
```

In [25]:

```
y=wine_df.iloc[:,0]
```

In [26]:

```
y
```

Out[26]:

```
0      1
1      1
2      1
3      1
4      1
..
173    3
174    3
175    3
176    3
177    3
```

Name: Class, Length: 170, dtype: int64

In [27]:

```
y.shape
```

Out[27]:

```
(170,)
```

In [28]:

```
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=.22,random_state=42)
```

In [29]:

```
#we are using follwing models for prediction
#KNeighborsClassifier
#GaussianNB
#SVC
#DecisionTreeClassifier
#RandomForestClassifier
#AdaBoostClassifier
```

In [30]:

```
knn=KNeighborsClassifier()
knn.fit(x_train,y_train)
knn.score(x_train,y_train)
predknn=knn.predict(x_test)
print(accuracy_score(y_test,predknn))
print(confusion_matrix(y_test,predknn))
print(classification_report(y_test,predknn))
```

0.9736842105263158

```
[[16  0  0]
 [ 0 13  1]
 [ 0  0  8]]
```

	precision	recall	f1-score	support
1	1.00	1.00	1.00	16
2	1.00	0.93	0.96	14
3	0.89	1.00	0.94	8
accuracy			0.97	38
macro avg	0.96	0.98	0.97	38
weighted avg	0.98	0.97	0.97	38

In [31]:


```

from sklearn.model_selection import cross_val_score
knn_scores=cross_val_score(knn,x,y,cv=5)
print(knn_scores)
print(knn_scores.mean(),knn_scores.std())

```

```

[0.88235294 0.94117647 0.88235294 1.          0.94117647]
0.9294117647058824 0.04401949866792873

```

In [32]:

```

mnb=MultinomialNB()
mnb.fit(x_train,y_train)
mnb.score(x_train,y_train)
predmnb=mnb.predict(x_test)
print(accuracy_score(y_test,predmnb))
print(confusion_matrix(y_test,predmnb))
print(classification_report(y_test,predmnb))

```

```

0.8947368421052632
[[13  3  0]
 [ 1 13  0]
 [ 0  0  8]]

```

	precision	recall	f1-score	support
1	0.93	0.81	0.87	16
2	0.81	0.93	0.87	14
3	1.00	1.00	1.00	8
accuracy			0.89	38
macro avg	0.91	0.91	0.91	38
weighted avg	0.90	0.89	0.89	38

In [33]:

```

mnbscores=cross_val_score(mnb,x,y,cv=5)
print(mnbscores)
print(mnbscores.mean(),mnbscores.std())

```

```

[0.85294118 0.91176471 0.73529412 0.91176471 1.          ]
0.8823529411764707 0.08724939396583131

```

In [34]:

```

svc=SVC(kernel='rbf')
svc.fit(x_train,y_train)
svc.score(x_train,y_train)
predsvc=svc.predict(x_test)
print(accuracy_score(y_test,predsvc))
print(confusion_matrix(y_test,predsvc))
print(classification_report(y_test,predsvc))

```

```

0.8947368421052632
[[14  2  0]
 [ 0 12  2]
 [ 0  0  8]]

```

	precision	recall	f1-score	support
1	1.00	0.88	0.93	16
2	0.86	0.86	0.86	14
3	0.80	1.00	0.89	8
accuracy			0.89	38
macro avg	0.89	0.91	0.89	38
weighted avg	0.91	0.89	0.90	38

In [35]:

```

mnbscores=cross_val_score(mnb,x,y,cv=5)

```

```
svcscores=cross_val_score(svc,x,y,cv=5)
print(svcscores)
print(svcscores.mean(),svcscores.std())
```

```
[0.70588235 0.79411765 0.73529412 0.97058824 0.88235294]
0.8176470588235294 0.09737026680733438
```

In [36]:

```
dtc=DecisionTreeClassifier()
dtc.fit(x_train,y_train)
dtc.score(x_train,y_train)
preddtc=dtc.predict(x_test)
print(accuracy_score(y_test,preddtc))
print(confusion_matrix(y_test,preddtc))
print(classification_report(y_test,preddtc))
```

```
1.0
[[16  0  0]
 [ 0 14  0]
 [ 0  0  8]]
```

	precision	recall	f1-score	support
1	1.00	1.00	1.00	16
2	1.00	1.00	1.00	14
3	1.00	1.00	1.00	8
accuracy			1.00	38
macro avg	1.00	1.00	1.00	38
weighted avg	1.00	1.00	1.00	38

In [37]:

```
dtcscores=cross_val_score(dtc,x,y,cv=5)
print(dtcscores)
print(dtcscores.mean(),dtcscores.std())
```

```
[0.88235294 0.82352941 0.91176471 0.97058824 1.          ]
0.9176470588235294 0.06280634265900772
```

In [38]:

```
rf=RandomForestClassifier()
rf.fit(x_train,y_train)
rf.score(x_train,y_train)
predrf=rf.predict(x_test)
print(accuracy_score(y_test,predrf))
print(confusion_matrix(y_test,predrf))
print(classification_report(y_test,predrf))
```

```
0.9736842105263158
[[16  0  0]
 [ 0 13  1]
 [ 0  0  8]]
```

	precision	recall	f1-score	support
1	1.00	1.00	1.00	16
2	1.00	0.93	0.96	14
3	0.89	1.00	0.94	8
accuracy			0.97	38
macro avg	0.96	0.98	0.97	38
weighted avg	0.98	0.97	0.97	38

In [39]:

```
rfscores=cross_val_score(rf,x,y,cv=5)
print(rfscores)
print(rfscores.mean(),rfscores.std())
```

```
[0.94117647 0.97058824 1.          1.          1.          ]
0.9823529411764707 0.02352941176470589
```

In [40]:

```
ad=AdaBoostClassifier()
ad.fit(x_train,y_train)
ad.score(x_train,y_train)
predad=ad.predict(x_test)
print(accuracy_score(y_test,predad))
print(confusion_matrix(y_test,predad))
print(classification_report(y_test,predad))
```

```
0.5789473684210527
[[ 0 16  0]
 [ 0 14  0]
 [ 0  0  8]]
```

	precision	recall	f1-score	support
1	0.00	0.00	0.00	16
2	0.47	1.00	0.64	14
3	1.00	1.00	1.00	8
accuracy			0.58	38
macro avg	0.49	0.67	0.55	38
weighted avg	0.38	0.58	0.44	38

In [41]:

```
adscores=cross_val_score(ad,x,y,cv=5)
print(adscores)
print(adscores.mean(),adscores.std())
```

```
[0.70588235 1.          0.91176471 0.91176471 0.88235294]
0.8823529411764707 0.09665692191267634
```

In [42]:

```
#RandomForestClassifier and DecisionTreeClassifier are the best model among all models
import joblib
joblib.dump(dtc,'wine_df.pkl')
```

Out[42]:

```
['wine_df.pkl']
```

In []: