In [1]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings("ignore")
```

In [32]:

```
df=pd.read_csv("Breast_Cancer.csv")
df.sample(10)
```

Out[32]:

	Sample code number	Clump Thickness	Uniformity of Cell Size	Uniformity of Cell Shape	Marginal Adhesion	Single Epithelial Cell Size	Bare Nuclei	Bland Chromatin	Norr Nucle
326	809912	10	3	3	1	2	10	7	
189	1212422	4	1	1	1	2	1	3	
32	1074610	2	1	1	2	2	1	3	
493	1297327	5	1	1	1	2	1	1	
564	824249	1	1	1	1	2	1	3	
637	1324681	4	1	1	1	2	1	2	
384	1196475	3	2	1	1	2	1	2	
93	1164066	1	1	1	1	2	1	3	
94	1165297	2	1	1	2	2	1	1	
41	1102573	5	6	5	6	10	1	3	
4									•

In [3]:

df.shape

Out[3]:

(683, 11)

In [4]:

df.columns

Out[4]:

In [5]:

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

Out[5]:

Sample code number 0 Clump Thickness 0 Uniformity of Cell Size 0 Uniformity of Cell Shape 0 Marginal Adhesion 0 Single Epithelial Cell Size 0 Bare Nuclei 0 Bland Chromatin 0 Normal Nucleoli 0 Mitoses 0 Class 0 dtype: int64

In [6]:

```
# Info of dataframe
df.info()
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 683 entries, 0 to 682
Data columns (total 11 columns):

#	Column	Non-Null Count	Dtype
0	Sample code number	683 non-null	int64
1	Clump Thickness	683 non-null	int64
2	Uniformity of Cell Size	683 non-null	int64
3	Uniformity of Cell Shape	683 non-null	int64
4	Marginal Adhesion	683 non-null	int64
5	Single Epithelial Cell Size	683 non-null	int64
6	Bare Nuclei	683 non-null	int64
7	Bland Chromatin	683 non-null	int64
8	Normal Nucleoli	683 non-null	int64
9	Mitoses	683 non-null	int64
10	Class	683 non-null	int64

dtypes: int64(11)
memory usage: 58.8 KB

In [7]:

```
df.Class.value_counts()
```

Out[7]:

2 4444 239

Name: Class, dtype: int64

In [8]:

```
df.corr()["Class"]
```

Out[8]:

Sample code number -0.084701 Clump Thickness 0.714790 Uniformity of Cell Size 0.820801 Uniformity of Cell Shape 0.821891 Marginal Adhesion 0.706294 Single Epithelial Cell Size 0.690958 Bare Nuclei 0.822696 Bland Chromatin 0.758228 Normal Nucleoli 0.718677 0.423448 Mitoses Class 1.000000

Name: Class, dtype: float64

In [9]:

```
df.drop("Sample code number",axis=1,inplace=True)
```

In [10]:

```
df.head()
```

Out[10]:

	Clump Thickness	Uniformity of Cell Size	Uniformity of Cell Shape	Marginal Adhesion	Single Epithelial Cell Size	Bare Nuclei	Bland Chromatin	Normal Nucleoli	Mitoses
0	5	1	1	1	2	1	3	1	1
1	5	4	4	5	7	10	3	2	1
2	3	1	1	1	2	2	3	1	1
3	6	8	8	1	3	4	3	7	1
4	4	1	1	3	2	1	3	1	1
4									•

In [11]:

```
def mymodel(obj):
    obj.fit(xtrain,ytrain)
    ypred=obj.predict(xtest)
    print(classification_report(ytest,ypred))
```

In [12]:

```
x=df.drop("Class",axis=1)
x.head()
```

Out[12]:

	Clump Thickness	Uniformity of Cell Size	Uniformity of Cell Shape	Marginal Adhesion	Single Epithelial Cell Size	Bare Nuclei	Bland Chromatin	Normal Nucleoli	Mitoses
0	5	1	1	1	2	1	3	1	1
1	5	4	4	5	7	10	3	2	1
2	3	1	1	1	2	2	3	1	1
3	6	8	8	1	3	4	3	7	1
4	4	1	1	3	2	1	3	1	1

→

In [13]:

```
y=df["Class"]
y.head()
```

Out[13]:

- 0 2
- 1 2
- 2 2
- 3 2
- 4 2

Name: Class, dtype: int64

To build the best model, we have to train and test the dataset with multiple Machine Learning algorithms then we can find the best ML model. So let's try.

First, we need to import the required packages.

In [14]:

```
from sklearn.model_selection import train_test_split
xtrain,xtest,ytrain,ytest=train_test_split(x,y,train_size=0.70,random_state=1)
```

In [15]:

```
from sklearn.linear_model import LogisticRegression
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier
```

In [16]:

```
from sklearn.metrics import classification_report
from sklearn.metrics import accuracy_score
```

In [17]:

```
lr=LogisticRegression()
knn=KNeighborsClassifier()
svc=SVC()
dt=DecisionTreeClassifier()
```

2

In [18]:

mymodel(lr)				
	precision	recall	f1-score	support

133

0.97

4	0.94	0.94	0.94	72
accuracy			0.96	205
macro avg	0.96	0.96	0.96	205
weighted avg	0.96	0.96	0.96	205

0.97

0.97

In [19]:

mymodel(knn)

	precision	recall	f1-score	support
2 4	0.98 0.93	0.96 0.97	0.97 0.95	133 72
accuracy macro avg weighted avg	0.96 0.97	0.97 0.97	0.97 0.96 0.97	205 205 205

In [20]:

mymodel(svc)

	precision	recall	f1-score	support
2 4	1.00 0.95	0.97 1.00	0.98 0.97	133 72
accuracy macro avg weighted avg	0.97 0.98	0.98 0.98	0.98 0.98 0.98	205 205 205

In [21]:

mym	odel(dt)					
-----	----------	--	--	--	--	--

support	f1-score	recall	precision	
133	0.97	0.98	0.97	2
72	0.95	0.94	0.96	4
205	0.97			accuracy
205	0.96	0.96	0.96	macro avg
205	0.97	0.97	0.97	weighted avg

In [22]:

dt1=DecisionTreeClassifier(max_depth=5)
mymodel(dt1)

	precision	recall	f1-score	support
2	0.98	0.97	0.97	133
4	0.95	0.96	0.95	72
accuracy			0.97	205
macro avg	0.96	0.96	0.96	205
weighted avg	0.97	0.97	0.97	205

In [23]:

```
for i in range(1,50):
    dt1.fit(xtrain,ytrain)
    ypred=dt1.predict(xtest)
    dt1=DecisionTreeClassifier(max_depth=i)
    print(f"{i}-----{accuracy_score(ytest,ypred)}")
```

```
1-----0.9658536585365853
2-----0.9219512195121952
3-----0.9512195121951219
4-----0.9463414634146341
5----0.9609756097560975
6-----0.9560975609756097
7-----0.9609756097560975
8-----0.9609756097560975
9-----0.9707317073170731
10-----0.9658536585365853
11-----0.9560975609756097
12-----0.9658536585365853
13-----0.9707317073170731
14-----0.9560975609756097
15-----0.9609756097560975
16-----0.9560975609756097
17-----0.9512195121951219
18-----0.9609756097560975
19-----0.9560975609756097
20-----0.9512195121951219
21-----0.9707317073170731
22----0.9560975609756097
23-----0.9658536585365853
24-----0.9658536585365853
25-----0.9560975609756097
26-----0.9560975609756097
27----0.9560975609756097
28-----0.9609756097560975
29-----0.9560975609756097
30-----0.9658536585365853
31----0.9560975609756097
32-----0.9560975609756097
33-----0.9609756097560975
34-----0.9609756097560975
35-----0.9512195121951219
36-----0.9609756097560975
37-----0.9609756097560975
38-----0.9609756097560975
39----0.9609756097560975
40-----0.9658536585365853
41-----0.9707317073170731
42----0.9609756097560975
43-----0.9560975609756097
44----0.9560975609756097
45-----0.9609756097560975
46-----0.9609756097560975
47-----0.9609756097560975
48-----0.9658536585365853
49----0.9609756097560975
```

In [24]:

dt3=DecisionTreeClassifier(max_depth=12)
mymodel(dt3)

	precision	recall	f1-score	support
2	0.97	0.97	0.97	133
4	0.94	0.94	0.94	72
accuracy			0.96	205
macro avg	0.96	0.96	0.96	205
weighted avg	0.96	0.96	0.96	205

In [25]:

dt4=DecisionTreeClassifier(min_samples_leaf=12)
mymodel(dt4)

	precision	recall	f1-score	support
2	0.98	0.95	0.97	133
4	0.92	0.96	0.94	72
accuracy			0.96	205
macro avg	0.95	0.96	0.95	205
weighted avg	0.96	0.96	0.96	205

In [26]:

```
for i in range(1,50):
    dt4.fit(xtrain,ytrain)
    ypred=dt4.predict(xtest)
    dt4=DecisionTreeClassifier(min_samples_leaf=i)
    print(f"{i}------{accuracy_score(ytest,ypred)}")
```

```
1-----0.9560975609756097
2-----0.9609756097560975
3-----0.9463414634146341
4----0.9609756097560975
5-----0.9512195121951219
6-----0.9512195121951219
7-----0.9512195121951219
8-----0.9463414634146341
9-----0.9560975609756097
10-----0.9560975609756097
11-----0.9560975609756097
12-----0.9560975609756097
13-----0.9560975609756097
14-----0.9560975609756097
15-----0.9365853658536586
16-----0.9512195121951219
17-----0.9512195121951219
18-----0.9512195121951219
19-----0.9512195121951219
20-----0.9512195121951219
21-----0.9512195121951219
22-----0.9512195121951219
23-----0.9512195121951219
24-----0.9512195121951219
25-----0.9512195121951219
26-----0.9512195121951219
27-----0.9512195121951219
28-----0.9512195121951219
29-----0.9512195121951219
30-----0.9512195121951219
31-----0.9512195121951219
32-----0.9512195121951219
33-----0.9414634146341463
34-----0.9414634146341463
35-----0.9414634146341463
36-----0.9414634146341463
37-----0.9414634146341463
38-----0.9414634146341463
39-----0.9414634146341463
40-----0.9414634146341463
41-----0.9414634146341463
42----0.9414634146341463
43-----0.9414634146341463
44-----0.9219512195121952
45----0.9219512195121952
46-----0.9219512195121952
47-----0.9219512195121952
48-----0.9219512195121952
49-----0.9219512195121952
```

In [27]:

dt5=DecisionTreeClassifier(min_samples_leaf=4)
mymodel(dt5)

	precision	recall	f1-score	support
2	0.98	0.94	0.96	133
4	0.90	0.97	0.93	72
accuracy			0.95	205
macro avg	0.94	0.96	0.95	205
weighted avg	0.95	0.95	0.95	205

In [28]:

dt6=DecisionTreeClassifier(min_samples_split=24)
mymodel(dt6)

	precision	recall	f1-score	support
2	0.98	0.94	0.96	133
4	0.90	0.96	0.93	72
accuracy			0.95	205
macro avg	0.94	0.95	0.94	205
weighted avg	0.95	0.95	0.95	205

In [29]:

```
for i in range(3,50):
    dt6.fit(xtrain,ytrain)
    ypred=dt6.predict(xtest)
    dt6=DecisionTreeClassifier(min_samples_split=i)
    print(f"{i}----{accuracy_score(ytest,ypred)}")
3----0.9463414634146341
```

```
3----0.9463414634146341
4----0.9560975609756097
5----0.9658536585365853
6----0.9609756097560975
7----0.9609756097560975
8----0.9560975609756097
9----0.9463414634146341
10----0.9560975609756097
11----0.9512195121951219
12----0.9512195121951219
13----0.9512195121951219
14----0.9512195121951219
15----0.9463414634146341
16----0.9463414634146341
17----0.9463414634146341
18----0.9463414634146341
19----0.9463414634146341
20----0.9463414634146341
21----0.9463414634146341
22----0.9463414634146341
23----0.9463414634146341
24----0.9463414634146341
25----0.9463414634146341
26----0.9463414634146341
27----0.9463414634146341
28----0.9463414634146341
29----0.9463414634146341
30----0.9463414634146341
31----0.9463414634146341
32----0.9463414634146341
33----0.9414634146341463
34----0.9414634146341463
35----0.9414634146341463
36----0.9414634146341463
37----0.9414634146341463
38----0.9414634146341463
39----0.9414634146341463
40----0.9414634146341463
41----0.9414634146341463
42----0.9414634146341463
43----0.9414634146341463
44----0.9414634146341463
45----0.9414634146341463
46----0.9414634146341463
47----0.9414634146341463
48----0.9414634146341463
49----0.9414634146341463
```

In [30]:

dt7=DecisionTreeClassifier(max_depth=12,min_samples_leaf=4,min_samples_split=4)
mymodel(dt7)

	precision	recall	f1-score	support
2 4	0.98 0.90	0.94 0.97	0.96 0.93	133 72
accuracy macro avg weighted avg	0.94 0.95	0.96 0.95	0.95 0.95 0.95	205 205 205

In [31]:

dt8=DecisionTreeClassifier(max_depth=12)
mymodel(dt8)

	precision	recall	f1-score	support
2 4	0.97 0.94	0.97 0.94	0.97 0.94	133 72
accuracy macro avg weighted avg	0.96 0.96	0.96 0.96	0.96 0.96 0.96	205 205 205

I have completed the Machine learning Project successfully with 97% accuracy which is great for 'Breast Cancer Detection using Machine learning' project.

To get more accuracy, we trained all supervised classification algorithms but you can try out a few of them which are always popular. After training all algorithms, we found that Logistic Regression and Random Forest are given high accuracy.

In []: