

Importing required libraries

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

➤ Reading and importing Dataset which is Breast cancer detection

```
df=pd.read_csv("breast-cancer.csv.xls")
```

df

| | id | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | smoothn |
|-----|----------|-----------|-------------|--------------|----------------|-----------|---------|
| 0 | 842302 | M | 17.99 | 10.38 | 122.80 | 1001.0 | |
| 1 | 842517 | M | 20.57 | 17.77 | 132.90 | 1326.0 | |
| 2 | 84300903 | M | 19.69 | 21.25 | 130.00 | 1203.0 | |
| 3 | 84348301 | M | 11.42 | 20.38 | 77.58 | 386.1 | |
| 4 | 84358402 | M | 20.29 | 14.34 | 135.10 | 1297.0 | |
| ... | ... | ... | ... | ... | ... | ... | |
| 564 | 926424 | M | 21.56 | 22.39 | 142.00 | 1479.0 | |
| 565 | 926682 | M | 20.13 | 28.25 | 131.20 | 1261.0 | |
| 566 | 926954 | M | 16.60 | 28.08 | 108.30 | 858.1 | |
| 567 | 927241 | M | 20.60 | 29.33 | 140.10 | 1265.0 | |
| 568 | 92751 | B | 7.76 | 24.54 | 47.92 | 181.0 | |

569 rows × 32 columns



➤ checking descriptive statistics

```
df.describe()
```

| | id | radius_mean | texture_mean | perimeter_mean | area_mean | smoothness_mean | compactness |
|-------|--------------|-------------|--------------|----------------|-------------|-----------------|-------------|
| count | 5.690000e+02 | 569.000000 | 569.000000 | 569.000000 | 569.000000 | 569.000000 | 569 |
| mean | 3.037183e+07 | 14.127292 | 19.289649 | 91.969033 | 654.889104 | 0.096360 | 0 |
| std | 1.250206e+08 | 3.524049 | 4.301036 | 24.298981 | 351.914129 | 0.014064 | 0 |
| min | 8.670000e+03 | 6.981000 | 9.710000 | 43.790000 | 143.500000 | 0.052630 | 0 |
| 25% | 8.692180e+05 | 11.700000 | 16.170000 | 75.170000 | 420.300000 | 0.086370 | 0 |
| 50% | 9.060240e+05 | 13.370000 | 18.840000 | 86.240000 | 551.100000 | 0.095870 | 0 |
| 75% | 8.813129e+06 | 15.780000 | 21.800000 | 104.100000 | 782.700000 | 0.105300 | 0 |
| max | 9.113205e+08 | 28.110000 | 39.280000 | 188.500000 | 2501.000000 | 0.163400 | 0 |

8 rows × 31 columns



▼ checking total number of rows and columns

```
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 32 columns):
 #   Column                                  Non-Null Count  Dtype  
---  --
 0   id                                      569 non-null    int64  
 1   diagnosis                              569 non-null    object  
 2   radius_mean                            569 non-null    float64 
 3   texture_mean                           569 non-null    float64 
 4   perimeter_mean                         569 non-null    float64 
 5   area_mean                              569 non-null    float64 
 6   smoothness_mean                        569 non-null    float64 
 7   compactness_mean                       569 non-null    float64 
 8   concavity_mean                         569 non-null    float64 
 9   concave points_mean                    569 non-null    float64 
10  symmetry_mean                          569 non-null    float64 
11  fractal_dimension_mean                 569 non-null    float64 
12  radius_se                              569 non-null    float64 
13  texture_se                             569 non-null    float64 
14  perimeter_se                           569 non-null    float64 
15  area_se                                569 non-null    float64 
16  smoothness_se                          569 non-null    float64 
17  compactness_se                         569 non-null    float64 
18  concavity_se                           569 non-null    float64 
19  concave points_se                      569 non-null    float64 
20  symmetry_se                            569 non-null    float64 
21  fractal_dimension_se                   569 non-null    float64 
22  radius_worst                           569 non-null    float64 
23  texture_worst                          569 non-null    float64 
24  perimeter_worst                        569 non-null    float64 
25  area_worst                             569 non-null    float64 
26  smoothness_worst                       569 non-null    float64 
27  compactness_worst                      569 non-null    float64 
28  concavity_worst                        569 non-null    float64 
29  concave points_worst                   569 non-null    float64 
30  symmetry_worst                         569 non-null    float64 
31  fractal_dimension_worst                 569 non-null    float64 
dtypes: float64(30), int64(1), object(1)
memory usage: 142.4+ KB
```

▼ Checking columns(features)

```
df.head()
```

| | id | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | smoothness_mean | compactn |
|---|----------|-----------|-------------|--------------|----------------|-----------|-----------------|----------|
| 0 | 842302 | M | 17.99 | 10.38 | 122.80 | 1001.0 | 0.11840 | |
| 1 | 842517 | M | 20.57 | 17.77 | 132.90 | 1326.0 | 0.08474 | |
| 2 | 84300903 | M | 19.69 | 21.25 | 130.00 | 1203.0 | 0.10960 | |
| 3 | 84348301 | M | 11.42 | 20.38 | 77.58 | 386.1 | 0.14250 | |
| 4 | 84358402 | M | 20.29 | 14.34 | 135.10 | 1297.0 | 0.10030 | |

5 rows × 32 columns



```
df["diagnosis"].value_counts()

B    357
M    212
Name: diagnosis, dtype: int64
```

^ according to description of dataset B means BENIGN not cancerous and M means MALIGNANT it is cancerous

▼ in this dataset ID columns in not required

```
df.drop("id",inplace=True,axis=1)
```

```
df
```

| | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | smoothness_mean | compactness_mean |
|-----|-----------|-------------|--------------|----------------|-----------|-----------------|------------------|
| 0 | M | 17.99 | 10.38 | 122.80 | 1001.0 | 0.11840 | 0.27760 |
| 1 | M | 20.57 | 17.77 | 132.90 | 1326.0 | 0.08474 | 0.07864 |
| 2 | M | 19.69 | 21.25 | 130.00 | 1203.0 | 0.10960 | 0.15990 |
| 3 | M | 11.42 | 20.38 | 77.58 | 386.1 | 0.14250 | 0.28390 |
| 4 | M | 20.29 | 14.34 | 135.10 | 1297.0 | 0.10030 | 0.13280 |
| ... | ... | ... | ... | ... | ... | ... | ... |
| 564 | M | 21.56 | 22.39 | 142.00 | 1479.0 | 0.11100 | 0.11590 |
| 565 | M | 20.13 | 28.25 | 131.20 | 1261.0 | 0.09780 | 0.10340 |
| 566 | M | 16.60 | 28.08 | 108.30 | 858.1 | 0.08455 | 0.10230 |
| 567 | M | 20.60 | 29.33 | 140.10 | 1265.0 | 0.11780 | 0.27700 |
| 568 | B | 7.76 | 24.54 | 47.92 | 181.0 | 0.05263 | 0.04362 |

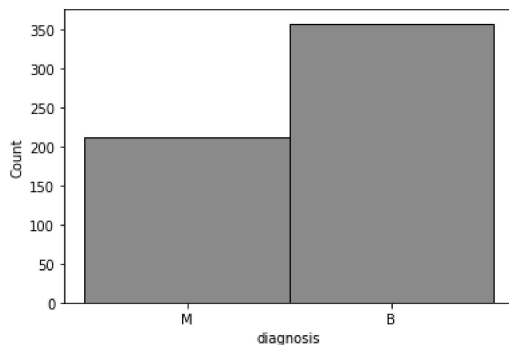
569 rows × 8 columns



▼ Visualization

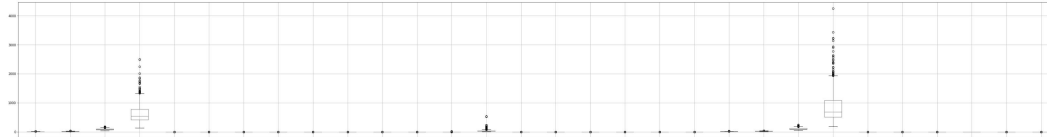
```
histplot=sns.histplot(df["diagnosis"],label="count")
B,M = df["diagnosis"].value_counts()
print("number of Benign(not cancerous): ",B)
print("number of Malignant(cancerous): ",M)
```

```
number of Benign(not cancerous): 357
number of Malignant(cancerous): 212
```



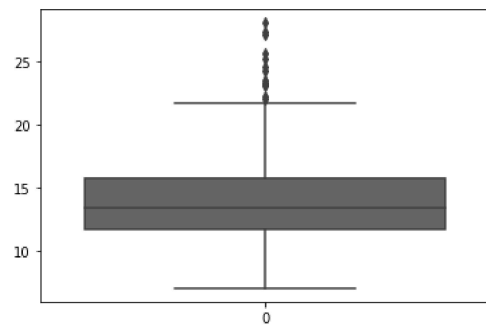
```
plt.figure(figsize=(60,8))
df.boxplot()
```

<Axes: >



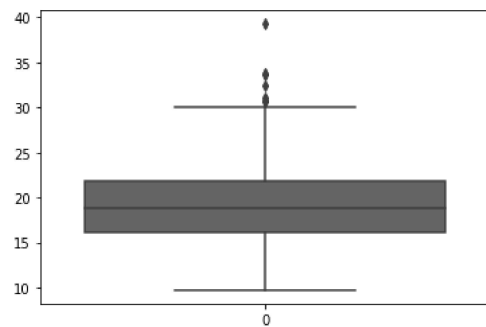
```
sns.boxplot(df["radius_mean"])
```

<Axes: >



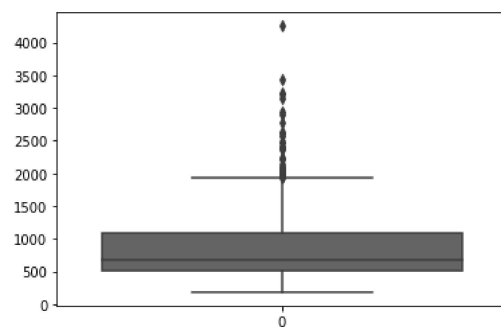
```
sns.boxplot(df["texture_mean"])
```

<Axes: >



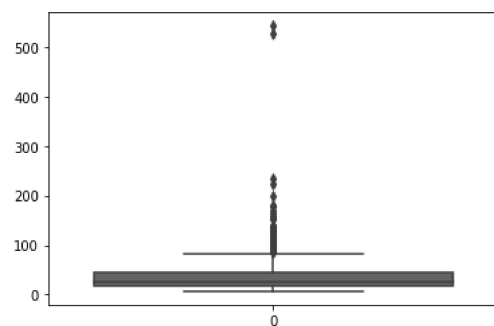
```
sns.boxplot(df["area_worst"])
```

<Axes: >



```
sns.boxplot(df["area_se"])
```

<Axes: >



```
#THERE ARE SOME OUTLIERS PRESENT IN BOXPLOTS
```

REMOVING OUTLIERS

```
df=df[(df["radius_mean"] < 23) & (df["texture_mean"] < 35) & (df["area_worst"] < 2300) & (df["area_se"] < 150)]
```

```
df
```

| | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | smoothness_mean | compactness_mean |
|-----|-----------|-------------|--------------|----------------|-----------|-----------------|------------------|
| 1 | M | 20.57 | 17.77 | 132.90 | 1326.0 | 0.08474 | 0.07864 |
| 2 | M | 19.69 | 21.25 | 130.00 | 1203.0 | 0.10960 | 0.15990 |
| 3 | M | 11.42 | 20.38 | 77.58 | 386.1 | 0.14250 | 0.28390 |
| 4 | M | 20.29 | 14.34 | 135.10 | 1297.0 | 0.10030 | 0.13280 |
| 5 | M | 12.45 | 15.70 | 82.57 | 477.1 | 0.12780 | 0.17000 |
| ... | ... | ... | ... | ... | ... | ... | ... |
| 563 | M | 20.92 | 25.09 | 143.00 | 1347.0 | 0.10990 | 0.22360 |
| 565 | M | 20.13 | 28.25 | 131.20 | 1261.0 | 0.09780 | 0.10340 |
| 566 | M | 16.60 | 28.08 | 108.30 | 858.1 | 0.08455 | 0.10230 |
| 567 | M | 20.60 | 29.33 | 140.10 | 1265.0 | 0.11780 | 0.27700 |
| 568 | B | 7.76 | 24.54 | 47.92 | 181.0 | 0.05263 | 0.04362 |

546 rows × 31 columns



so according to histplot there are 200 cancerous patients and 350 non cancerous so the count of non cancerous patients are greater than cancerous patients

splitting data into x(features) and y(label)

```
x=df.iloc[:,1:]
```

```
x
```

```
y=df.iloc[:, :1].values
```

 y

```
['M'],
['B'],
['B'],
['M'],
['B'],
['M'],
['M'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['B'],
['M'],
['B'],
['M'],
['M'],
['M'],
['B'], dtype=object)
```

```
# Encoding for label
```

converting categorical label into numeric

```
from sklearn.preprocessing import LabelEncoder
```

```
le=LabelEncoder()
```

```
y=le.fit_transform(y)
```

```
y
```

```
array([1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0,
       1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 1, 0, 0, 0, 0, 1, 0, 1, 1, 0, 0, 0,
       0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 0, 0, 0, 1, 0, 0, 1, 1, 1, 0, 0, 0, 1,
       0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 1, 1, 1, 0, 1, 0, 0, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 0, 0,
       1, 0, 0, 1, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0,
       0, 1, 1, 0, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 1, 0,
       1, 0, 1, 0, 0, 0, 1, 0, 0, 1, 1, 0, 1, 1, 1, 0, 1, 1, 0, 1, 0,
       1, 0, 0, 1, 0, 1, 1, 1, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 1,
       0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0,
       0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1,
       0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0,
       1, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0,
       0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 1,
       1, 0, 1, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0,
       1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0,
       1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 0,
       0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0,
       0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0,
       0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 1, 0, 0, 0, 0, 1, 1, 0, 1,
       0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 0, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 0])
```

1 = M (cancerous) , 0 = B (non cancerous)

▼ performing standardiation(scailing) for better results

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

```
sc=StandardScaler()
```

```
x=sc.fit_transform(x)
```

```
x
```

```
array([[ 2.21677361, -0.32770195,  2.06017605, ...,  1.22694486,
        -0.239839   ,  0.27945786],
       [ 1.92977419,  0.51032166,  1.92249255, ...,  2.14473916,
        1.15564227,  0.20033731],
       [-0.76736803,  0.30081576, -0.56625533, ...,  2.37821315,
        6.04710326,  4.89372479],
       ...,
       [ 0.92201489,  2.15506342,  0.89224019, ...,  0.51525173,
        -1.10008932, -0.31504521],
       [ 2.22655768,  2.45607765,  2.40201094, ...,  2.49897556,
        1.92210591,  2.20142806],
       [-1.96102468,  1.30259112, -1.97442516, ..., -1.76796287,
        -0.04418056, -0.74416434]])
```

▼ SPLITTING DATA INTO TRAINING AND TESTING

```
from sklearn.model_selection import train_test_split
```

```
xtrain,xtest,ytrain,ytest=train_test_split(x,y,test_size=0.3,random_state=1)
```

▼ BUILDING A FUNCTION FOR MODELS AND FOR ACCURACY

```
def mymodel(model):
    model.fit(xtrain,ytrain)
    ypred=model.predict(xtest)

    train=model.score(xtrain,ytrain)
    test=model.score(xtest,ytest)

    print(f"Training Accuracy(bias):- {train}\n Testing Accuracy(variance):- {test}")
    print(classification_report(ytest,ypred))

    return model
```

▼ IMPORTING ALL MODELS WHICH ARE REQUIRED

```
from sklearn.linear_model import LogisticRegression
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.naive_bayes import BernoulliNB
from sklearn.metrics import classification_report
```

FOR LOGISTIC REGRESSION

```
logreg = mymodel(LogisticRegression())
```

```
Training Accuracy(bias):- 0.9842931937172775
Testing Accuracy(variance):- 0.9939024390243902
      precision    recall  f1-score   support

      0         1.00      0.99      1.00       112
      1         0.98      1.00      0.99        52

   accuracy          0.99          1.00      0.99       164
  macro avg          0.99          1.00      0.99       164
weighted avg          0.99          0.99      0.99       164
```

FOR KNN ALGORITHM

```
k = mymodel(KNeighborsClassifier())
```

```
Training Accuracy(bias):- 0.9790575916230366
Testing Accuracy(variance):- 0.9817073170731707
      precision    recall  f1-score   support

      0         0.99      0.98      0.99       112
      1         0.96      0.98      0.97        52

   accuracy          0.98          0.98      0.98       164
  macro avg          0.98          0.98      0.98       164
weighted avg          0.98          0.98      0.98       164
```

FOR SUPPORT VECTOR MACHINE

```
s = mymodel(SVC())
```

```
Training Accuracy(bias):- 0.9869109947643979
Testing Accuracy(variance):- 0.975609756097561
      precision    recall  f1-score   support

      0         0.98      0.98      0.98       112
      1         0.96      0.96      0.96        52

   accuracy          0.97          0.97      0.98       164
  macro avg          0.97          0.97      0.97       164
weighted avg          0.98          0.98      0.98       164
```



```
#FOR DECISION TRESS
```

```
d = mymodel(DecisionTreeClassifier())

Training Accuracy(bias):- 1.0
Testing Accuracy(variance):- 0.9207317073170732
      precision    recall  f1-score   support

      0       0.96       0.92       0.94       112
      1       0.84       0.92       0.88       52

   accuracy       0.92       0.92       0.92       164
  macro avg       0.90       0.92       0.91       164
weighted avg       0.92       0.92       0.92       164
```

```
#FOR RANDOM FOREST
```

```
r = mymodel(RandomForestClassifier())

Training Accuracy(bias):- 1.0
Testing Accuracy(variance):- 0.9634146341463414
      precision    recall  f1-score   support

      0       0.98       0.96       0.97       112
      1       0.93       0.96       0.94       52

   accuracy       0.96       0.96       0.96       164
  macro avg       0.95       0.96       0.96       164
weighted avg       0.96       0.96       0.96       164
```

```
# FOR BAYES THEORM
```

```
n = mymodel(BernoulliNB())

Training Accuracy(bias):- 0.9240837696335078
Testing Accuracy(variance):- 0.9451219512195121
      precision    recall  f1-score   support

      0       0.99       0.93       0.96       112
      1       0.86       0.98       0.92       52

   accuracy       0.95       0.95       0.95       164
  macro avg       0.93       0.95       0.94       164
weighted avg       0.95       0.95       0.95       164
```

OBSERVING ALL THE MODELS ACCURACY OF LOGISTIC REGRESSION IS BEST FOR THIS DATASET BECAUSE OF IT IS A BINARY CLASSIFICATION DATASET

▼ WHAT IF WE USE ARTIFICIAL NEURAL NETWORK ON THIS DATASET LETS CHECK

```
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense

from tensorflow.keras.callbacks import EarlyStopping

early_stop= EarlyStopping(monitor="val_loss",mode="min",verbose=1,patience=10)

model=Sequential()
model.add(Dense(20,activation="relu"))
model.add(Dense(20,activation="relu"))
model.add(Dense(1,activation="sigmoid"))
```

```
model.compile(optimizer="sgd",loss="binary_crossentropy",metrics=["accuracy"])
```

```
model.fit(xtrain,ytrain,epochs=600,validation_data=(xtest,ytest),verbose=1,batch_size=30,callbacks=[early_stop])
```

```
Epoch 8/600
13/13 [=====] - 0s 4ms/step - loss: 0.0762 - accuracy: 0.9843 - val_loss: 0.0435 - val_accuracy: 0.9878
Epoch 88/600
13/13 [=====] - 0s 4ms/step - loss: 0.0758 - accuracy: 0.9843 - val_loss: 0.0432 - val_accuracy: 0.9878
Epoch 89/600
13/13 [=====] - 0s 4ms/step - loss: 0.0753 - accuracy: 0.9843 - val_loss: 0.0428 - val_accuracy: 0.9878
Epoch 90/600
13/13 [=====] - 0s 4ms/step - loss: 0.0748 - accuracy: 0.9843 - val_loss: 0.0425 - val_accuracy: 0.9878
Epoch 91/600
13/13 [=====] - 0s 4ms/step - loss: 0.0744 - accuracy: 0.9843 - val_loss: 0.0422 - val_accuracy: 0.9878
Epoch 92/600
13/13 [=====] - 0s 5ms/step - loss: 0.0740 - accuracy: 0.9843 - val_loss: 0.0419 - val_accuracy: 0.9878
Epoch 93/600
13/13 [=====] - 0s 4ms/step - loss: 0.0735 - accuracy: 0.9843 - val_loss: 0.0416 - val_accuracy: 0.9878
Epoch 94/600
13/13 [=====] - 0s 6ms/step - loss: 0.0732 - accuracy: 0.9843 - val_loss: 0.0413 - val_accuracy: 0.9878
Epoch 95/600
13/13 [=====] - 0s 4ms/step - loss: 0.0727 - accuracy: 0.9843 - val_loss: 0.0410 - val_accuracy: 0.9878
Epoch 96/600
13/13 [=====] - 0s 4ms/step - loss: 0.0723 - accuracy: 0.9869 - val_loss: 0.0407 - val_accuracy: 0.9878
Epoch 97/600
13/13 [=====] - 0s 5ms/step - loss: 0.0719 - accuracy: 0.9817 - val_loss: 0.0404 - val_accuracy: 0.9878
Epoch 98/600
13/13 [=====] - 0s 4ms/step - loss: 0.0715 - accuracy: 0.9843 - val_loss: 0.0401 - val_accuracy: 0.9878
Epoch 99/600
13/13 [=====] - 0s 4ms/step - loss: 0.0711 - accuracy: 0.9817 - val_loss: 0.0399 - val_accuracy: 0.9878
Epoch 100/600
13/13 [=====] - 0s 4ms/step - loss: 0.0707 - accuracy: 0.9869 - val_loss: 0.0396 - val_accuracy: 0.9878
Epoch 101/600
13/13 [=====] - 0s 4ms/step - loss: 0.0704 - accuracy: 0.9869 - val_loss: 0.0394 - val_accuracy: 0.9878
Epoch 102/600
13/13 [=====] - 0s 5ms/step - loss: 0.0700 - accuracy: 0.9843 - val_loss: 0.0392 - val_accuracy: 0.9878
Epoch 103/600
13/13 [=====] - 0s 4ms/step - loss: 0.0696 - accuracy: 0.9843 - val_loss: 0.0390 - val_accuracy: 0.9878
Epoch 104/600
13/13 [=====] - 0s 4ms/step - loss: 0.0692 - accuracy: 0.9843 - val_loss: 0.0388 - val_accuracy: 0.9878
Epoch 105/600
13/13 [=====] - 0s 4ms/step - loss: 0.0688 - accuracy: 0.9843 - val_loss: 0.0385 - val_accuracy: 0.9878
Epoch 106/600
13/13 [=====] - 0s 4ms/step - loss: 0.0685 - accuracy: 0.9843 - val_loss: 0.0383 - val_accuracy: 0.9878
Epoch 107/600
13/13 [=====] - 0s 4ms/step - loss: 0.0681 - accuracy: 0.9843 - val_loss: 0.0380 - val_accuracy: 0.9878
Epoch 108/600
13/13 [=====] - 0s 4ms/step - loss: 0.0679 - accuracy: 0.9843 - val_loss: 0.0379 - val_accuracy: 0.9878
Epoch 109/600
13/13 [=====] - 0s 4ms/step - loss: 0.0673 - accuracy: 0.9843 - val_loss: 0.0377 - val_accuracy: 0.9878
Epoch 110/600
13/13 [=====] - 0s 4ms/step - loss: 0.0670 - accuracy: 0.9843 - val_loss: 0.0374 - val_accuracy: 0.9878
Epoch 111/600
13/13 [=====] - 0s 5ms/step - loss: 0.0666 - accuracy: 0.9843 - val_loss: 0.0372 - val_accuracy: 0.9878
Epoch 112/600
13/13 [=====] - 0s 5ms/step - loss: 0.0662 - accuracy: 0.9843 - val_loss: 0.0370 - val_accuracy: 0.9878
Epoch 113/600
13/13 [=====] - 0s 4ms/step - loss: 0.0660 - accuracy: 0.9843 - val_loss: 0.0369 - val_accuracy: 0.9878
Epoch 114/600
13/13 [=====] - 0s 5ms/step - loss: 0.0656 - accuracy: 0.9843 - val_loss: 0.0366 - val_accuracy: 0.9878
Epoch 115/600
13/13 [=====] - 0s 4ms/step - loss: 0.0651 - accuracy: 0.9843 - val_loss: 0.0364 - val_accuracy: 0.9878
Epoch 116/600
```

```
ypred=model.predict(xtest)
```

```
6/6 [=====] - 0s 1ms/step
```

```
ypred=ypred>0.5
```

```
print(classification_report(ytest,ypred))
```

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 1.00 | 1.00 | 1.00 | 112 |
| 1 | 1.00 | 1.00 | 1.00 | 52 |
| accuracy | | | 1.00 | 164 |
| macro avg | 1.00 | 1.00 | 1.00 | 164 |
| weighted avg | 1.00 | 1.00 | 1.00 | 164 |

OUR ACCURACY OF ARTIFICIAL NEURAL NETWORK IS 100 PERCENT