

Breast Cancer Classification.

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import sklearn.datasets
```

Data Importing and Preprocessing

```
In [2]: columns_name = ['ID', 'DIAGNOSIS', 'RADIUS_MEAN', 'TEXTURE_MEAN', 'PERIMETER_MEAN',
                        'COMPACTNESS_MEAN', 'CONCATIVITY_MEAN', 'CONCAVE_POINTS_MEAN', 'SYMMETRIC_COEFF',
                        'PERIMETER_SE', 'AREA_SE', 'SMOOTHNESS_SE', 'COMPACTNESS_SE', 'CONCATIVITY_SE',
                        'RADIUS_WORST', 'TEXTURE_WORST', 'PERIMETER_WORST', 'AREA_WORST', 'CONCAVE_POINTS_WORST',
                        'SYMMETRIC_COEFF_WORST']
```

```
In [3]: Breast_Data = pd.read_csv("wdbc.data.csv", names= columns_name)
```

```
In [4]: Breast_Data.head()
```

```
Out[4]:
```

	ID	DIAGNOSIS	RADIUS_MEAN	TEXTURE_MEAN	PERIMETER_MEAN	AREA_MEAN	PERIMETER_SE	AREA_SE	SMOOTHNESS_SE	COMPACTNESS_SE	CONCATIVITY_SE	CONCAVE_POINTS_SE	SYMMETRIC_COEFF_SE
0	842302	M	17.99	10.38	122.80	1001.0	1.66	0.0066	0.0846	0.7571	0.1985	0.2738	0.0871
1	842517	M	20.57	17.77	132.90	1326.0	1.68	0.0067	0.0854	0.7602	0.1990	0.2740	0.0872
2	84300903	M	19.69	21.25	130.00	1203.0	1.67	0.0066	0.0849	0.7587	0.1988	0.2739	0.0871
3	84348301	M	11.42	20.38	77.58	386.1	1.65	0.0065	0.0841	0.7562	0.1980	0.2731	0.0869
4	84358402	M	20.29	14.34	135.10	1297.0	1.67	0.0066	0.0849	0.7587	0.1988	0.2739	0.0871

5 rows × 32 columns

```
In [5]: Breast_Data.tail()
```

```
Out[5]:
```

	ID	DIAGNOSIS	RADIUS_MEAN	TEXTURE_MEAN	PERIMETER_MEAN	AREA_MEAN	PERIMETER_SE	AREA_SE	SMOOTHNESS_SE	COMPACTNESS_SE	CONCATIVITY_SE	CONCAVE_POINTS_SE	SYMMETRIC_COEFF_SE
564	926424	M	21.56	22.39	142.00	1479.0	1.68	0.0067	0.0854	0.7602	0.1990	0.2740	0.0872
565	926682	M	20.13	28.25	131.20	1261.0	1.67	0.0066	0.0849	0.7587	0.1988	0.2739	0.0871
566	926954	M	16.60	28.08	108.30	858.1	1.66	0.0066	0.0846	0.7571	0.1985	0.2738	0.0871
567	927241	M	20.60	29.33	140.10	1265.0	1.68	0.0067	0.0854	0.7602	0.1990	0.2740	0.0872
568	92751	B	7.76	24.54	47.92	181.0	1.65	0.0065	0.0841	0.7562	0.1980	0.2731	0.0869

5 rows × 32 columns

Getting Rows and Columns of Dataset

```
In [6]: Breast_Data.shape
```

```
Out[6]: (569, 32)
```

Getting some information about the Dataset

In [7]: `Breast_Data.info`

```

Out[7]: <bound method DataFrame.info of
MEAN PERIMETER_MEAN AREA_MEAN \
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

SMOOTHNESS_MEAN COMPACTNESS_MEAN CONCATIVITY_MEAN CONCAVE_POINTS_MEAN
\
0      0.11840      0.27760      0.30010      0.14710
1      0.08474      0.07864      0.08690      0.07017
2      0.10960      0.15990      0.19740      0.12790
3      0.14250      0.28390      0.24140      0.10520
4      0.10030      0.13280      0.19800      0.10430
..      ...      ...      ...      ...
564      0.11100      0.11590      0.24390      0.13890
565      0.09780      0.10340      0.14400      0.09791
566      0.08455      0.10230      0.09251      0.05302
567      0.11780      0.27700      0.35140      0.15200
568      0.05263      0.04362      0.00000      0.00000

... RADIUS_WORST TEXTURE_WORST PERIMETER_WORST AREA_WORST \
0      ...      25.380      17.33      184.60      2019.0
1      ...      24.990      23.41      158.80      1956.0
2      ...      23.570      25.53      152.50      1709.0
3      ...      14.910      26.50      98.87      567.7
4      ...      22.540      16.67      152.20      1575.0
..      ...      ...      ...      ...
564      ...      25.450      26.40      166.10      2027.0
565      ...      23.690      38.25      155.00      1731.0
566      ...      18.980      34.12      126.70      1124.0
567      ...      25.740      39.42      184.60      1821.0
568      ...      9.456      30.37      59.16      268.6

SMOOTHNESS_WORST COMPACTNESS_WORST CONCATIVITY_WORST \
0      0.16220      0.66560      0.7119
1      0.12380      0.18660      0.2416
2      0.14440      0.42450      0.4504
3      0.20980      0.86630      0.6869
4      0.13740      0.20500      0.4000
..      ...      ...      ...
564      0.14100      0.21130      0.4107
565      0.11660      0.19220      0.3215
566      0.11390      0.30940      0.3403
567      0.16500      0.86810      0.9387
568      0.08996      0.06444      0.0000

CONCAVE_POINTS_WORST SYMMETRY_WORST FRACTAL_DIMENSION_WORST
0      0.2654      0.4601      0.11890
1      0.1860      0.2750      0.08902
2      0.2430      0.3613      0.08758
3      0.2575      0.6638      0.17300
4      0.1625      0.2364      0.07678

```

```

..          ...          ...          ...
564          0.2216          0.2060          0.07115
565          0.1628          0.2572          0.06637
566          0.1418          0.2218          0.07820
567          0.2650          0.4087          0.12400
568          0.0000          0.2871          0.07039

```

```
[569 rows x 32 columns]>
```

```
In [8]: Breast_Data.describe()
```

```
Out[8]:
```

	ID	RADIUS_MEAN	TEXTURE_MEAN	PERIMETER_MEAN	AREA_MEAN	SMOO
count	5.690000e+02	569.000000	569.000000	569.000000	569.000000	
mean	3.037183e+07	14.127292	19.289649	91.969033	654.889104	
std	1.250206e+08	3.524049	4.301036	24.298981	351.914129	
min	8.670000e+03	6.981000	9.710000	43.790000	143.500000	
25%	8.692180e+05	11.700000	16.170000	75.170000	420.300000	
50%	9.060240e+05	13.370000	18.840000	86.240000	551.100000	
75%	8.813129e+06	15.780000	21.800000	104.100000	782.700000	
max	9.113205e+08	28.110000	39.280000	188.500000	2501.000000	

```
8 rows x 31 columns
```

Getting Target points from Datasets B: Benign, M: Malignant

```
In [9]: Breast_Data['DIAGNOSIS'].value_counts()
```

```
Out[9]: B      357
        M      212
        Name: DIAGNOSIS, dtype: int64
```

Getting Mean of Malignant and Benign Data from dataset (GroupBy).

```
In [10]: Breast_Data_Grouped = Breast_Data.groupby('DIAGNOSIS')
print(Breast_Data_Grouped)

<pandas.core.groupby.generic.DataFrameGroupBy object at 0x7fe273106430>
```

```
In [11]: Breast_Data_Grouped.mean()
```

```
Out[11]:
```

	ID	RADIUS_MEAN	TEXTURE_MEAN	PERIMETER_MEAN	AREA_MEAN	SMOO
DIAGNOSIS						
B	2.654382e+07	12.146524	17.914762	78.075406	462.790196	
M	3.681805e+07	17.462830	21.604906	115.365377	978.376415	

```
2 rows x 31 columns
```

```
In [ ]:
```

Converting Diagnosis column to 0,1 for training the model.

```
In [12]: from sklearn.preprocessing import LabelEncoder
```

```
In [13]: lb_Diagnosis = LabelEncoder()
```

```
In [14]: Breast_Data['Label_Diagnosis'] = lb_Diagnosis.fit_transform(Breast_Data['DIAGNOSIS'])
```

Created Label_Diagnosis column where M: 1 and B:0

```
In [16]: Breast_Data['Label_Diagnosis'].info
```

```
Out[16]: <bound method Series.info of 0      1
1      1
2      1
3      1
4      1
..
564    1
565    1
566    1
567    1
568    0
Name: Label_Diagnosis, Length: 569, dtype: int64>
```

Training the Dataset and Splitting into Test and Training variables.

```
In [20]: from sklearn import preprocessing
from sklearn.model_selection import train_test_split
```

```
In [21]: X = Breast_Data.drop(columns=['DIAGNOSIS', 'Label_Diagnosis'], axis=1)
Y = Breast_Data['Label_Diagnosis']
```

```
In [23]: X.shape, Y.shape
```

```
Out[23]: ((569, 31), (569,))
```

```
In [24]: X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2, random_state=42)
```

```
In [26]: X_train.shape, X_test.shape, Y_train.shape, Y_test.shape
```

```
Out[26]: ((455, 31), (114, 31), (455,), (114,))
```

```
In [27]: Y_train
```

```
Out[27]: 560    0
         428    0
         198    1
         203    1
         41     1
         ..
         299    0
         534    0
         493    0
         527    0
         168    1
Name: Label_Diagnosis, Length: 455, dtype: int64
```

Scaling the Training and Tsting Dataframe "As we know that remaining columns is a Numerical Data so it would be beneficial for sklearn's regresstion model understand the data while training"

```
In [72]: Scaler = preprocessing.StandardScaler()
```

```
In [29]: X_trainS = Scaler.fit_transform(X_train)
         X_testS = Scaler.fit_transform(X_test)
```

```
In [30]: X_trainS.shape, X_testS.shape
```

```
Out[30]: ((455, 31), (114, 31))
```

Importing Tensorflow and keras to setup Neural network for classification

```
In [31]: import tensorflow as tf
         tf.random.set_seed(3)
```

2022-12-07 15:44:40.380632: I tensorflow/core/platform/cpu_feature_guard.cc:193] This TensorFlow binary is optimized with oneAPI Deep Neural Network Library (oneDNN) to use the following CPU instructions in performance-critical operations: AVX2 FMA
To enable them in other operations, rebuild TensorFlow with the appropriate compiler flags.

Making neural network layers using keras, Hidden Layer

```
In [32]: from tensorflow import keras
         MODEL = keras.Sequential([
             keras.layers.Flatten(input_shape=(31,)),
             keras.layers.Dense(20, activation='relu'),
             keras.layers.Dense(2, activation='sigmoid')
         ])
```

2022-12-07 15:44:48.893139: I tensorflow/core/platform/cpu_feature_guard.cc:193] This TensorFlow binary is optimized with oneAPI Deep Neural Network Library (oneDNN) to use the following CPU instructions in performance-critical operations: AVX2 FMA
To enable them in other operations, rebuild TensorFlow with the appropriate compiler flags.

```
In [33]: MODEL.compile(
         optimizer='adam',
```

```
loss='sparse_categorical_crossentropy',  
metrics=['accuracy']  
)
```

Training the Neural network

```
In [51]: trained_model = MODEL.fit(X_trainS, Y_train, epochs=25, validation_split=0.01)
```

```
Epoch 1/25
15/15 [=====] - 0s 7ms/step - loss: 0.0386 - accurac
y: 0.9889 - val_loss: 1.7687e-04 - val_accuracy: 1.0000
Epoch 2/25
15/15 [=====] - 0s 4ms/step - loss: 0.0383 - accurac
y: 0.9889 - val_loss: 1.7642e-04 - val_accuracy: 1.0000
Epoch 3/25
15/15 [=====] - 0s 4ms/step - loss: 0.0376 - accurac
y: 0.9889 - val_loss: 1.7509e-04 - val_accuracy: 1.0000
Epoch 4/25
15/15 [=====] - 0s 4ms/step - loss: 0.0370 - accurac
y: 0.9889 - val_loss: 1.5607e-04 - val_accuracy: 1.0000
Epoch 5/25
15/15 [=====] - 0s 4ms/step - loss: 0.0364 - accurac
y: 0.9911 - val_loss: 1.4801e-04 - val_accuracy: 1.0000
Epoch 6/25
15/15 [=====] - 0s 4ms/step - loss: 0.0359 - accurac
y: 0.9889 - val_loss: 1.4849e-04 - val_accuracy: 1.0000
Epoch 7/25
15/15 [=====] - 0s 4ms/step - loss: 0.0356 - accurac
y: 0.9889 - val_loss: 1.4262e-04 - val_accuracy: 1.0000
Epoch 8/25
15/15 [=====] - 0s 4ms/step - loss: 0.0352 - accurac
y: 0.9911 - val_loss: 1.3385e-04 - val_accuracy: 1.0000
Epoch 9/25
15/15 [=====] - 0s 4ms/step - loss: 0.0346 - accurac
y: 0.9889 - val_loss: 1.1884e-04 - val_accuracy: 1.0000
Epoch 10/25
15/15 [=====] - 0s 4ms/step - loss: 0.0341 - accurac
y: 0.9911 - val_loss: 1.2170e-04 - val_accuracy: 1.0000
Epoch 11/25
15/15 [=====] - 0s 4ms/step - loss: 0.0335 - accurac
y: 0.9911 - val_loss: 1.1555e-04 - val_accuracy: 1.0000
Epoch 12/25
15/15 [=====] - 0s 3ms/step - loss: 0.0330 - accurac
y: 0.9911 - val_loss: 1.1140e-04 - val_accuracy: 1.0000
Epoch 13/25
15/15 [=====] - 0s 4ms/step - loss: 0.0327 - accurac
y: 0.9911 - val_loss: 1.1400e-04 - val_accuracy: 1.0000
Epoch 14/25
15/15 [=====] - 0s 4ms/step - loss: 0.0321 - accurac
y: 0.9911 - val_loss: 1.0945e-04 - val_accuracy: 1.0000
Epoch 15/25
15/15 [=====] - 0s 4ms/step - loss: 0.0317 - accurac
y: 0.9911 - val_loss: 1.0046e-04 - val_accuracy: 1.0000
Epoch 16/25
15/15 [=====] - 0s 4ms/step - loss: 0.0313 - accurac
y: 0.9933 - val_loss: 9.8982e-05 - val_accuracy: 1.0000
Epoch 17/25
15/15 [=====] - 0s 3ms/step - loss: 0.0308 - accurac
y: 0.9933 - val_loss: 9.2450e-05 - val_accuracy: 1.0000
Epoch 18/25
15/15 [=====] - 0s 4ms/step - loss: 0.0303 - accurac
y: 0.9933 - val_loss: 8.5132e-05 - val_accuracy: 1.0000
Epoch 19/25
15/15 [=====] - 0s 5ms/step - loss: 0.0299 - accurac
y: 0.9933 - val_loss: 8.3368e-05 - val_accuracy: 1.0000
Epoch 20/25
15/15 [=====] - 0s 4ms/step - loss: 0.0296 - accurac
y: 0.9933 - val_loss: 8.2724e-05 - val_accuracy: 1.0000
```



```

Epoch 21/25
15/15 [=====] - 0s 5ms/step - loss: 0.0293 - accuracy: 0.9933 - val_loss: 7.5859e-05 - val_accuracy: 1.0000
Epoch 22/25
15/15 [=====] - 0s 4ms/step - loss: 0.0287 - accuracy: 0.9933 - val_loss: 7.6098e-05 - val_accuracy: 1.0000
Epoch 23/25
15/15 [=====] - 0s 4ms/step - loss: 0.0284 - accuracy: 0.9933 - val_loss: 7.2927e-05 - val_accuracy: 1.0000
Epoch 24/25
15/15 [=====] - 0s 4ms/step - loss: 0.0281 - accuracy: 0.9933 - val_loss: 6.8851e-05 - val_accuracy: 1.0000
Epoch 25/25
15/15 [=====] - 0s 4ms/step - loss: 0.0278 - accuracy: 0.9933 - val_loss: 6.6777e-05 - val_accuracy: 1.0000

```

In [61]: `trained_model.params`

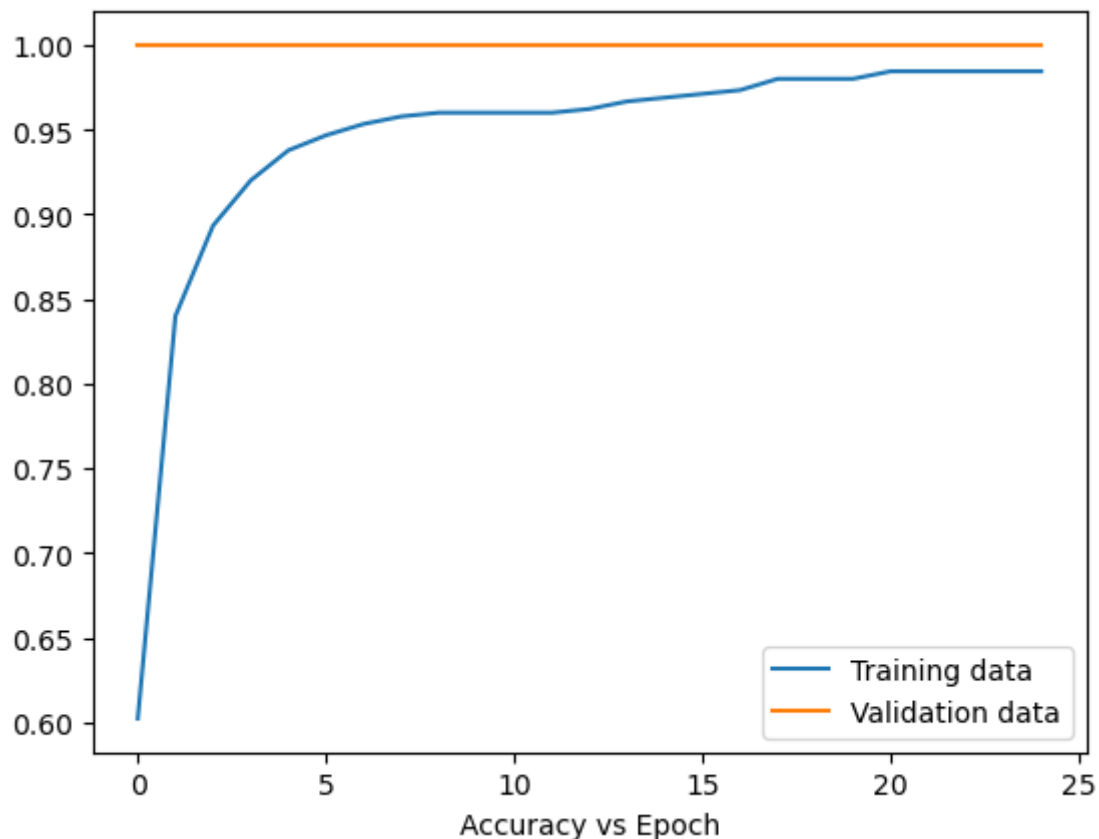
Out[61]: `{'verbose': 1, 'epochs': 25, 'steps': 15}`

```

In [195... %matplotlib inline
plt.plot(trained_model.history['accuracy'])
plt.plot(trained_model.history['val_accuracy'])
plt.xlabel('Accuracy vs Epoch')
plt.legend(['Training data', 'Validation data'], loc = 'lower right')

```

Out[195]: `<matplotlib.legend.Legend at 0x7fe24c28ea60>`



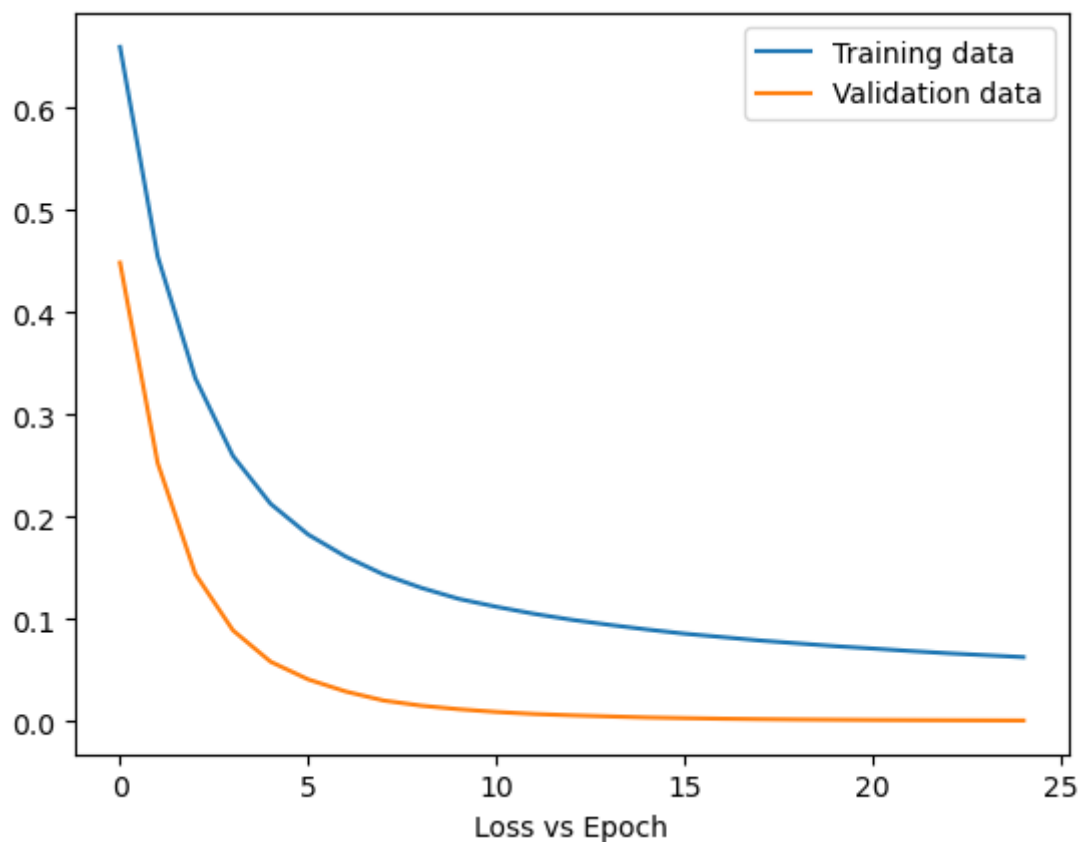
```

In [97]: %matplotlib inline
plt.plot(trained_model.history['loss'])
plt.plot(trained_model.history['val_loss'])

```

```
plt.xlabel('Loss vs Epoch')
plt.legend(['Training data', 'Validation data'], loc = 'upper right')
```

Out[97]: <matplotlib.legend.Legend at 0x7fe24ac2e730>



In [64]: `trained_model.history['accuracy']`

Out[64]:

```
[0.602222204208374,
0.8399999737739563,
0.8933333158493042,
0.9200000166893005,
0.9377777576446533,
0.9466666579246521,
0.95333331823349,
0.9577777981758118,
0.9599999785423279,
0.9599999785423279,
0.9599999785423279,
0.9599999785423279,
0.9599999785423279,
0.9622222185134888,
0.9666666388511658,
0.9688888788223267,
0.9711111187934875,
0.9733333587646484,
0.9800000190734863,
0.9800000190734863,
0.9800000190734863,
0.9844444394111633,
0.9844444394111633,
0.9844444394111633,
0.9844444394111633,
0.9844444394111633]
```

Finding The accuracy Of the model

```
In [66]: Loss, Accuracy = MODEL.evaluate(X_testS, Y_test)

4/4 [=====] - 0s 3ms/step - loss: 0.0879 - accuracy: 0.9737
```

```
In [67]: print(Loss, Accuracy)

0.08787020295858383 0.9736841917037964
```

```
In [68]: MODEL.predict(X_testS)

4/4 [=====] - 0s 2ms/step
```

```
Out[68]: array([[7.31051564e-01, 1.60803854e-01],
 [6.40998840e-01, 1.19231343e-01],
 [9.37755644e-01, 7.71941384e-04],
 [1.46365725e-10, 9.99996722e-01],
 [7.75570273e-01, 1.45885289e-01],
 [2.51738061e-06, 9.99234438e-01],
 [7.65389800e-01, 3.26402187e-02],
 [9.77818191e-01, 5.78860636e-04],
 [9.34239268e-01, 3.35949101e-03],
 [9.51056778e-01, 3.64095857e-03],
 [2.69510746e-01, 5.29248536e-01],
 [8.63291442e-01, 2.08233427e-02],
 [3.33902031e-01, 5.19329421e-02],
 [7.74546206e-01, 5.58350533e-02],
 [8.96992207e-01, 2.01532012e-03],
 [1.39508245e-03, 8.70443106e-01],
 [9.41922307e-01, 2.42094253e-03],
 [9.88540828e-01, 1.83954940e-03],
 [8.10406327e-01, 6.12178305e-03],
 [1.59763676e-05, 9.99054551e-01],
 [7.55457938e-01, 8.93481309e-04],
 [9.81762886e-01, 1.28782354e-03],
 [9.41038787e-01, 2.99099670e-03],
 [9.88616705e-01, 7.75113702e-04],
 [9.31395590e-01, 1.40412310e-02],
 [3.82299797e-04, 9.89704430e-01],
 [9.33316231e-01, 1.08463056e-02],
 [8.33260000e-01, 8.09682533e-02],
 [1.38626585e-03, 9.58235323e-01],
 [5.59933134e-04, 9.76477087e-01],
 [9.55433547e-01, 1.30898431e-02],
 [9.30512309e-01, 5.04098088e-03],
 [9.76345599e-01, 2.23975838e-03],
 [1.08914362e-08, 9.99970734e-01],
 [3.70614871e-05, 9.96860445e-01],
 [9.55898643e-01, 9.72562749e-03],
 [9.46212113e-01, 5.24495146e-04],
 [7.72765160e-01, 3.02293189e-02],
 [9.82399225e-01, 9.01362626e-04],
 [9.68568623e-01, 3.78753850e-03],
 [3.20463944e-09, 9.99991238e-01],
 [4.50073220e-02, 8.21912766e-01],
 [7.52093434e-01, 2.11587059e-03],
 [9.91524577e-01, 9.89714987e-04],
 [7.18579628e-03, 9.57393646e-01],
 [7.09191144e-01, 1.22571678e-03],
 [9.95527565e-01, 1.83623939e-04],
 [8.86224985e-01, 7.61702948e-04],
 [6.74261514e-07, 9.99415815e-01],
 [1.28702656e-03, 9.76919532e-01],
 [9.74253297e-01, 1.78660871e-03],
 [3.64623405e-02, 7.34966755e-01],
 [4.88777280e-01, 2.46961400e-01],
 [9.05061066e-01, 3.91130336e-03],
 [9.79775548e-01, 4.70038998e-04],
 [4.95718181e-01, 2.34145314e-01],
 [7.24772930e-01, 6.43579066e-02],
 [9.47778225e-01, 3.16638208e-04],
 [2.30452002e-04, 9.63313103e-01],
 [9.84664559e-01, 2.54050177e-03],
```

```
[8.15996885e-01, 7.85517767e-02],  
[2.14043539e-03, 9.31277215e-01],  
[9.86494899e-01, 1.08903891e-03],  
[8.36508843e-05, 9.95497286e-01],  
[4.16360563e-03, 8.83585155e-01],  
[9.31065381e-01, 2.64307903e-03],  
[1.48323625e-06, 9.99362111e-01],  
[2.74783233e-03, 9.63605404e-01],  
[4.38670486e-01, 1.14846162e-01],  
[2.60057777e-01, 6.23337217e-02],  
[6.14540651e-02, 6.10680342e-01],  
[2.29546975e-04, 9.92862225e-01],  
[9.46358562e-01, 2.77371169e-03],  
[1.52478134e-02, 9.22355473e-01],  
[9.86157715e-01, 3.17322672e-04],  
[1.06969764e-02, 9.36964154e-01],  
[9.35326517e-01, 2.55872938e-03],  
[9.90373254e-01, 9.76463896e-04],  
[6.53627098e-01, 8.44691247e-02],  
[2.68205479e-02, 9.03193891e-01],  
[1.52896289e-04, 9.90437508e-01],  
[7.53849233e-03, 9.32610810e-01],  
[4.61395830e-05, 9.97916758e-01],  
[9.51771140e-01, 9.77563206e-03],  
[9.29120004e-01, 5.08035999e-03],  
[2.57168531e-01, 2.49635205e-01],  
[9.96572077e-01, 2.61942710e-04],  
[9.87862170e-01, 7.43782613e-04],  
[9.73691404e-01, 8.20897426e-03],  
[3.91975254e-06, 9.98498261e-01],  
[9.80087698e-01, 2.14376533e-03],  
[9.09832656e-01, 2.88372356e-02],  
[9.97190773e-01, 5.36317064e-04],  
[4.92907013e-04, 9.52523053e-01],  
[1.38734635e-02, 9.19794023e-01],  
[9.77464557e-01, 2.48204288e-03],  
[1.70654857e-05, 9.98443663e-01],  
[1.42859906e-04, 9.85853493e-01],  
[7.73116529e-01, 2.43944768e-02],  
[9.49195504e-01, 6.48476183e-04],  
[9.80180800e-01, 3.17703205e-04],  
[4.84219491e-02, 8.77735734e-01],  
[6.30656132e-08, 9.99676943e-01],  
[2.06084081e-07, 9.99889791e-01],  
[8.47639740e-01, 8.97210930e-03],  
[9.94968712e-01, 1.84790217e-04],  
[9.98517811e-01, 9.22357067e-05],  
[9.90152776e-01, 4.10761219e-04],  
[9.17411208e-01, 1.63754885e-05],  
[5.05371332e-01, 5.50724491e-02],  
[8.09481753e-06, 9.96149719e-01],  
[5.73819443e-06, 9.99302149e-01],  
[3.40470187e-02, 4.52843338e-01],  
[6.62702776e-04, 9.85725522e-01]], dtype=float32)
```

In []:

Prediction Using random Data

```
In [74]: from sklearn.preprocessing import StandardScaler
```

```
In [75]: SCALER = StandardScaler()
```

```
In [82]: Random_data = (0,21.6,74.72,427.9,0.08637,0.04966,0.01657,0.01115,0.1495,0.0588)

Random_data_N = np.asarray(Random_data)

Random_data_N = Random_data_N.reshape(1,-1)

Random_data_NS = SCALER.fit_transform(Random_data_N)
```

```
In [83]: MODEL.predict(Random_data_NS)
```

```
1/1 [=====] - 0s 23ms/step
Out[83]: array([[0.5062147 , 0.42554605]], dtype=float32)
```

```
In [86]: Label = [np.argmax(MODEL.predict(Random_data_NS))]
Label
```

```
1/1 [=====] - 0s 22ms/step
Out[86]: [0]
```

```
In [87]: if(Label[0] == 0):
        print('The tumor is Malignant')

        else:
        print('The tumor is Benign')
```

The tumor is Malignant

Primary Tumor Dataset

```
In [113... Col_namesT = ["Class", "Age", "Sex", "Histological_type", "Degree_of_diff", "Bo
                "Peritoneum", "Liver", "Brain", "Skin", "Neck", "Supraclavicular
                ]
```

```
In [114... TumorData = pd.read_csv("primary-tumorD.csv", names= Col_namesT)
```

```
In [ ]:
```

```
In [116... TumorData["Age"].unique()
```

```
Out[116]: array([1, 2, 3])
```

```
In [120... TumorData.head()
```

Out[120]:

	Class	Age	Sex	Histological_type	Degree_of_diff	Bone	Bone-marow	lung	Pluera	Peritoneiu
0	1	1	1	?		3	2	2	1	2
1	1	1	1	?		3	2	2	2	2
2	1	1	2	2		3	1	2	2	2
3	1	1	2	?		3	1	2	1	1
4	1	1	2	?		3	1	2	1	1

In [121... TumorData.tail()

Out[121]:

	Class	Age	Sex	Histological_type	Degree_of_diff	Bone	Bone-marow	lung	Pluera	Peritoneiu
334	22	2	2	2		?	2	2	2	2
335	22	2	2	2		?	2	2	2	2
336	22	2	2	?		?	1	2	2	2
337	22	3	2	2		2	2	2	2	2
338	22	3	2	2		2	2	2	2	2

In [122... TumorData.shape

Out[122]: (339, 18)

In [124... TumorData.info

```

Out[124]: <bound method DataFrame.info of          Class  Age Sex Histological_type Degree_
of_diff  Bone  Bone-marow lung  \
0         1     1     1          ?           3     2           2     1
1         1     1     1          ?           3     2           2     2
2         1     1     2          2           3     1           2     2
3         1     1     2          ?           3     1           2     1
4         1     1     2          ?           3     1           2     1
..      ...   ...   ..          ...         ...   ...         ...   ...
334       22     2     2          2           ?     2           2     2
335       22     2     2          2           ?     2           2     2
336       22     2     2          ?           ?     1           2     2
337       22     3     2          2           2     2           2     2
338       22     3     2          2           2     2           2     2

      Pluera  Peritoneium  Liver  Brain  Skin  Neck  Supraclavicular  axillar  \
0           2           2     2     2     2     2           2           2
1           2           2     1     2     2     2           1           2
2           2           2     2     2     2     2           2           2
3           1           2     2     2     2     2           2           2
4           1           2     2     2     2     2           2           2
..      ...         ...   ...   ...   ...   ...         ...         ...
334         2           2     2     2     2     2           2           1
335         2           2     2     2     2     2           2           1
336         2           2     2     2     2     2           1           1
337         2           2     2     2     2     1           1           1
338         2           2     2     2     2     2           1           1

      Mediastinum  Abdominal
0                2          2
1                1          2
2                1          2
3                1          2
4                1          2
..              ...         ...
334              2          2
335              2          2
336              2          2
337              2          2
338              2          2

[339 rows x 18 columns]>

```

```
In [126... TumorData.describe()
```

```

Out[126]:

```

	Class	Age	Bone	Bone-marow	lung	Pluera	Peritoneiu
count	339.000000	339.000000	339.000000	339.000000	339.000000	339.000000	339.000000
mean	8.678466	2.247788	1.722714	1.979351	1.778761	1.778761	1.719761
std	7.052624	0.568362	0.448321	0.142416	0.415695	0.415695	0.449761
min	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000
25%	2.000000	2.000000	1.000000	2.000000	2.000000	2.000000	1.000000
50%	7.000000	2.000000	2.000000	2.000000	2.000000	2.000000	2.000000
75%	14.000000	3.000000	2.000000	2.000000	2.000000	2.000000	2.000000
max	22.000000	3.000000	2.000000	2.000000	2.000000	2.000000	2.000000

Checking for Null values, according to dataset description here null value indicated by "?"

```
In [137... TumorData.isnull().sum()
```

```
Out[137]: Class          0
Age            0
Sex            0
Histological_type  0
Degree_of_diff  0
Bone           0
Bone-marow     0
lung           0
Pluera         0
Peritoneium    0
Liver          0
Brain          0
Skin           0
Neck           0
Supraclavicular 0
axillar        0
Mediastinum    0
Abdominal      0
dtype: int64
```

```
In [140... TumorDataR = TumorData.replace("?", "0")
TumorDataR.head()
```

```
Out[140]:
```

	Class	Age	Sex	Histological_type	Degree_of_diff	Bone	Bone-marow	lung	Pluera	Peritoneiu
0	1	1	1	0	3	2	2	1	2	
1	1	1	1	0	3	2	2	2	2	
2	1	1	2	2	3	1	2	2	2	
3	1	1	2	0	3	1	2	1	1	
4	1	1	2	0	3	1	2	1	1	

```
In [141... TumorDataR["Class"].value_counts()
```

```
Out[141]:
```

1	84
5	39
18	29
11	28
14	24
22	24
2	20
12	16
7	14
4	14
17	10
3	9
13	7
8	6
19	6
10	2
15	2
20	2
6	1
16	1
21	1

```
Name: Class, dtype: int64
```

```
In [ ]:
```