

## INTRODUCTION

Here our project aims to develop a neural network model that can classify the tumors as malignant or benign by using the **Breast Cancer Wisconsin Dataset** from `sklearn`. The goal of this project is to produce an accurate and reliable model that assists in early detection by applying machine learning to support medical diagnostics.

## IMPORTING THE LIBRARIES AND NECESSARY DEPENDENCIES

In [42]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import sklearn.datasets
from sklearn.model_selection import train_test_split
```

Below are the imports used:

**NumPy** for creating arrays and performing mathematical computations, **Pandas** for handling data in the form of **DataFrames**, and **Matplotlib** for visualizing results after training. Additionally, **sklearn.datasets** provides datasets, and **train\_test\_split** is used for splitting the data into training and testing sets.

## DATA COLLECTION AND PRE PROCESSING

### LOADING THE DATASET

In [43]:

```
dataset = sklearn.datasets.load_breast_cancer()
```

In [44]:

```
print(dataset)
```

```
{'data': array([[1.799e+01, 1.038e+01, 1.228e+02, ..., 2.654e-01, 4.601e-01,
 1.189e-01],
 [2.057e+01, 1.777e+01, 1.329e+02, ..., 1.860e-01, 2.750e-01,
 8.902e-02],
 [1.969e+01, 2.125e+01, 1.300e+02, ..., 2.430e-01, 3.613e-01,
 8.758e-02],
 ...,
 [1.660e+01, 2.808e+01, 1.083e+02, ..., 1.418e-01, 2.218e-01,
 7.820e-02],
 [2.060e+01, 2.933e+01, 1.401e+02, ..., 2.650e-01, 4.087e-01,
 1.240e-01],
 [7.760e+00, 2.454e+01, 4.792e+01, ..., 0.000e+00, 2.871e-01,
 7.039e-02]]), 'target': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 1, 1, 1,
 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0,
0, 0, 1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0, 0,
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0, 1, 1, 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 0,
0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 0, 1, 0, 0,
1 1 1 1 1 0 1 1 1 1 1 0 1 1 1 0 1 1 0 0 1 1
```

```

1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 1, 0,
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1, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 0, 0,
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1, 1, 1, 0, 1, 1, 0, 1, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 0, 1, 0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1]), 'frame': None, 'target_r
ames': array(['malignant', 'benign'], dtype='<U9'), 'DESCR': '.. _breast_cancer_dataset:\n
n\nBreast cancer wisconsin (diagnostic) dataset\n-----\n
----\n\n**Data Set Characteristics:**\n\n: Number of Instances: 569\n\n: Number of Attribu
tes: 30 numeric, predictive attributes and the class\n\n: Attribute Information:\n    - ra
dius (mean of distances from center to points on the perimeter)\n    - texture (standard
deviation of gray-scale values)\n    - perimeter\n    - area\n    - smoothness (local var
iation in radius lengths)\n    - compactness (perimeter^2 / area - 1.0)\n    - concavity
(severity of concave portions of the contour)\n    - concave points (number of concave po
rtions of the contour)\n    - symmetry\n    - fractal dimension ("coastline approximation
" - 1)\n\n    The mean, standard error, and "worst" or largest (mean of the three\n    wo
rst/largest values) of these features were computed for each image,\n    resulting in 30
features. For instance, field 0 is Mean Radius, field\n    10 is Radius SE, field 20 is
Worst Radius.\n\n    - class:\n        - WDBC-Malignant\n        - WDBC-Benign\n\n
\n: Summary Statistics:\n\n===== \n
Min    Max\n===== \nradius (mean):
6.981  28.11\ntexture (mean):          9.71  39.28\nperimeter (mean):
43.79  188.5\narea (mean):            143.5  2501.0\nsmoothness (mean):
0.053  0.163\ncompactness (mean):      0.019  0.345\nconcavity (mean):
0.0     0.427\nconcave points (mean):   0.0    0.201\nsymmetry (mean):
0.106  0.304\nfractal dimension (mean): 0.05   0.097\nradius (standard error)
:          0.112  2.873\ntexture (standard error):          0.36  4.885\nperimet
er (standard error):          0.757  21.98\narea (standard error):          6.802
542.2\nsmoothness (standard error):      0.002  0.031\ncompactness (standard error):
0.002  0.135\nconcavity (standard error): 0.0    0.396\nconcave points (standar
d error): 0.0    0.053\nsymmetry (standard error):          0.008  0.079\nfractal
dimension (standard error): 0.001  0.03\nradius (worst):          7.93
36.04\ntexture (worst):          12.02  49.54\nperimeter (worst):
50.41  251.2\narea (worst):            185.2  4254.0\nsmoothness (worst):
0.071  0.223\ncompactness (worst):      0.027  1.058\nconcavity (worst):
0.0     1.252\nconcave points (worst):   0.0    0.291\nsymmetry (worst):
0.156  0.664\nfractal dimension (worst): 0.055  0.208\n=====
===== \n\n: Missing Attribute Values: None\n\n: Class Distribution: 2
12 - Malignant, 357 - Benign\n\n: Creator: Dr. William H. Wolberg, W. Nick Street, Olvi L
. Mangasarian\n\n: Donor: Nick Street\n\n: Date: November, 1995\n\nThis is a copy of UCI ML
Breast Cancer Wisconsin (Diagnostic) datasets.\nhttps://goo.gl/U2Uwz2\n\nFeatures are com
puted from a digitized image of a fine needle\naspirate (FNA) of a breast mass. They des
cribe\ncharacteristics of the cell nuclei present in the image.\n\nSeparating plane descr
ibed above was obtained using\nMultisurface Method-Tree (MSM-T) [K. P. Bennett, "Decision
Tree\nConstruction Via Linear Programming." Proceedings of the 4th\nMidwest Artificial In
telligence and Cognitive Science Society,\npp. 97-101, 1992], a classification method whi
ch uses linear\nprogramming to construct a decision tree. Relevant features\nwere select
ed using an exhaustive search in the space of 1-4\nfeatures and 1-3 separating planes.\n\n
The actual linear program used to obtain the separating plane\nin the 3-dimensional spac
e is that described in:\n[K. P. Bennett and O. L. Mangasarian: "Robust Linear\nProgrammin
g Discrimination of Two Linearly Inseparable Sets",\nOptimization Methods and Software 1,
1992, 23-34].\n\nThis database is also available through the UW CS ftp server:\n\nftp ftp
.cs.wisc.edu\ncd math-prog/cpo-dataset/machine-learn/WDBC/\n\n.. dropdown:: References\n
n    - W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction\n    for
breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on\n    Electronic Imaging
: Science and Technology, volume 1905, pages 861-870,\n    San Jose, CA, 1993.\n    - O.L.
Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis and\n    prognosis via
linear programming. Operations Research, 43(4), pages 570-577,\n    July-August 1995.\n    - W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine learning techniques\n    to di
agnose breast cancer from fine-needle aspirates. Cancer Letters 77 (1994)\n    163-171.\n
', 'feature_names': array(['mean radius', 'mean texture', 'mean perimeter', 'mean area',
'mean smoothness', 'mean compactness', 'mean concavity',
'mean concave points', 'mean symmetry', 'mean fractal dimension',
'radius error', 'texture error', 'perimeter error', 'area error',
'smoothness error', 'compactness error', 'concavity error',
'concave points error', 'symmetry error',
'fractal dimension error', 'worst radius', 'worst texture',
'worst perimeter', 'worst area', 'worst smoothness',
'worst compactness', 'worst concavity', 'worst concave points',
'worst symmetry', 'worst fractal dimension'], dtype='<U23'), 'filename': 'breast_c

```

```
worst_symmetry, worst_fractal_dimension ], dtype= <object> , filename : breast_cancer.csv', 'data_module': 'sklearn.datasets.data'}
```

LOADING THE DATA IN DATAFRAME

In [45]:

```
finaldataset= pd.DataFrame(dataset.data , columns = dataset.feature_names )
```

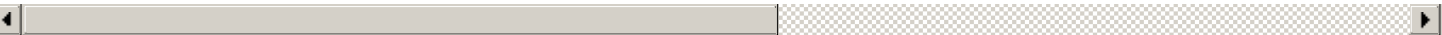
In [46]:

```
finaldataset.head()
```

Out[46]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	...	worst radius	worst texture
0	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001	0.14710	0.2419	0.07871	...	25.38	17.33
1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869	0.07017	0.1812	0.05667	...	24.99	23.44
2	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1974	0.12790	0.2069	0.05999	...	23.57	25.54
3	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2414	0.10520	0.2597	0.09744	...	14.91	26.50
4	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980	0.10430	0.1809	0.05883	...	22.54	16.66

5 rows x 30 columns



In [47]:

```
finaldataset.tail()
```

Out[47]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	...	worst radius	worst texture
564	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	0.13890	0.1726	0.05623	...	25.450	23.44
565	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	0.09791	0.1752	0.05533	...	23.690	34.56
566	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	0.05302	0.1590	0.05648	...	18.980	34.56
567	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	0.15200	0.2397	0.07016	...	25.740	34.56
568	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	0.00000	0.1587	0.05884	...	9.456	34.56

5 rows x 30 columns



ADDING THE TARGET COLUMN TO DATAFRAME

In [48]:

```
finaldataset['label']= dataset.target
```

In [49]:

```
finaldataset.head()
```

Out[49]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	...	worst radius	worst texture
0	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001	0.14710	0.2419	0.07871	...	17.33	17.33

1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869	0.07017	0.1812	0.05667	...	23.41	15.67
2	mean	mean	mean	mean	mean	mean	mean	mean	mean	mean	...	mean	mean
3	radius	texture	perimeter	area	smoothness	compactness	concavity	convex points	symmetry	fractal dimension	...	texture	perimeter
3	11.42	20.36	77.56	366.1	0.14250	0.28390	0.2414	0.10520	0.2597	0.09744	...	26.50	15.67
4	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980	0.10430	0.1809	0.05883	...	16.67	15.67

**5 rows x 31 columns**

◀ ▶

In [50]:

```
finaldataset.tail()
```

Out[50]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	...	worst texture	per
564	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	0.13890	0.1726	0.05623	...	26.40	
565	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	0.09791	0.1752	0.05533	...	38.25	
566	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	0.05302	0.1590	0.05648	...	34.12	
567	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	0.15200	0.2397	0.07016	...	39.42	
568	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	0.00000	0.1587	0.05884	...	30.37	

**5 rows x 31 columns**

◀ | | ▶

In [51]:

```
finaldataset.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

RangeIndex: 569 entries, 0 to 568

Data columns (total 31 columns):

#	Column	Non-Null Count	Dtype
0	mean radius	569 non-null	float64
1	mean texture	569 non-null	float64
2	mean perimeter	569 non-null	float64
3	mean area	569 non-null	float64
4	mean smoothness	569 non-null	float64
5	mean compactness	569 non-null	float64
6	mean concavity	569 non-null	float64
7	mean concave points	569 non-null	float64
8	mean symmetry	569 non-null	float64
9	mean fractal dimension	569 non-null	float64
10	radius error	569 non-null	float64
11	texture error	569 non-null	float64
12	perimeter error	569 non-null	float64
13	area error	569 non-null	float64
14	smoothness error	569 non-null	float64
15	compactness error	569 non-null	float64
16	concavity error	569 non-null	float64
17	concave points error	569 non-null	float64
18	symmetry error	569 non-null	float64
19	fractal dimension error	569 non-null	float64
20	worst radius	569 non-null	float64
21	worst texture	569 non-null	float64
22	worst perimeter	569 non-null	float64
23	worst area	569 non-null	float64
24	worst smoothness	569 non-null	float64
25	worst compactness	569 non-null	float64
26	worst concavity	569 non-null	float64
27	worst concave points	569 non-null	float64
28	worst symmetry	569 non-null	float64
29	worst fractal dimension	569 non-null	float64
30	label	569 non-null	int64

dtypes: float64(30), int64(1)  
memory usage: 137.9 KB

In [52]:

```
finaldataset.shape
```

Out[52]:

(569, 31)

In [53]:

```
finaldataset.describe()
```

Out[53]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	dim
count	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.
mean	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	0.088799	0.048919	0.181162	0.
std	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	0.079720	0.038803	0.027414	0.
min	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	0.000000	0.000000	0.106000	0.
25%	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	0.029560	0.020310	0.161900	0.
50%	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	0.061540	0.033500	0.179200	0.
75%	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	0.130700	0.074000	0.195700	0.
max	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	0.426800	0.201200	0.304000	0.

8 rows x 31 columns



In [54]:

```
finaldataset.isnull().sum( )
```

Out[54]:

	0
mean radius	0
mean texture	0
mean perimeter	0
mean area	0
mean smoothness	0
mean compactness	0
mean concavity	0
mean concave points	0
mean symmetry	0
mean fractal dimension	0
radius error	0
texture error	0
perimeter error	0
area error	0
smoothness error	0
compactness error	0
concavity error	0

concavity error	0
concave points error	0
symmetry error	0
fractal dimension error	0
worst radius	0
worst texture	0
worst perimeter	0
worst area	0
worst smoothness	0
worst compactness	0
worst concavity	0
worst concave points	0
worst symmetry	0
worst fractal dimension	0
label	0

dtype: int64

```
In [55]:
finaldataset['label'].value_counts()
```

Out[55]:

	count
label	
1	357
0	212

dtype: int64

in this dataset as we can see thers no imbalance on large extent so we can ignore it that we will not have to use methods like upsampeling or downsampeling.

here

**1 = Benign(non-cancerous)**

A benign tumor or condition is non-cancerous. It generally does not spread to other parts of the body and is not considered life-threatening. While it can cause symptoms due to its size or location, it usually does not pose a significant risk to health.

**0 =Malignant(cancerous)**

A malignant tumor or condition is cancerous. It has the potential to grow uncontrollably and spread to other parts of the body (a process known as metastasis). Malignant tumors can be life-threatening and often require more aggressive treatment.

```
In [56]:
finaldataset.groupby('label').mean()
```

Out[56]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	..
--	-------------	--------------	----------------	-----------	-----------------	------------------	----------------	---------------------	---------------	------------------------	----

label	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension
0	17.462830	21.604906	115.365377	978.376415	0.102898	0.145188	0.160775	0.070969	0.192909	0.062866
label	12.146524	17.914762	78.075406	462.790196	0.092478	0.080085	0.046058	0.025717	0.174186	0.062867

2 rows x 30 columns

## SPLITTING THE FEATURES AND TAEGET VARIABLES

In [57]:

```
X = finaldataset.drop(columns='label',axis=1)
Y = finaldataset['label']
```

In [58]:

```
print(X)
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	\
0	17.99	10.38	122.80	1001.0	0.11840	
1	20.57	17.77	132.90	1326.0	0.08474	
2	19.69	21.25	130.00	1203.0	0.10960	
3	11.42	20.38	77.58	386.1	0.14250	
4	20.29	14.34	135.10	1297.0	0.10030	
..	...	...	...	...	...	
564	21.56	22.39	142.00	1479.0	0.11100	
565	20.13	28.25	131.20	1261.0	0.09780	
566	16.60	28.08	108.30	858.1	0.08455	
567	20.60	29.33	140.10	1265.0	0.11780	
568	7.76	24.54	47.92	181.0	0.05263	

	mean compactness	mean concavity	mean concave points	mean symmetry	\
0	0.27760	0.30010	0.14710	0.2419	
1	0.07864	0.08690	0.07017	0.1812	
2	0.15990	0.19740	0.12790	0.2069	
3	0.28390	0.24140	0.10520	0.2597	
4	0.13280	0.19800	0.10430	0.1809	
..	...	...	...	...	
564	0.11590	0.24390	0.13890	0.1726	
565	0.10340	0.14400	0.09791	0.1752	
566	0.10230	0.09251	0.05302	0.1590	
567	0.27700	0.35140	0.15200	0.2397	
568	0.04362	0.00000	0.00000	0.1587	

	mean fractal dimension	...	worst radius	worst texture	\
0	0.07871	...	25.380	17.33	
1	0.05667	...	24.990	23.41	
2	0.05999	...	23.570	25.53	
3	0.09744	...	14.910	26.50	
4	0.05883	...	22.540	16.67	
..	...	...	...	...	
564	0.05623	...	25.450	26.40	
565	0.05533	...	23.690	38.25	
566	0.05648	...	18.980	34.12	
567	0.07016	...	25.740	39.42	
568	0.05884	...	9.456	30.37	

	worst perimeter	worst area	worst smoothness	worst compactness	\
0	184.60	2019.0	0.16220	0.66560	
1	158.80	1956.0	0.12380	0.18660	
2	152.50	1709.0	0.14440	0.42450	
3	98.87	567.7	0.20980	0.86630	
4	152.20	1575.0	0.13740	0.20500	
..	...	...	...	...	
564	166.10	2027.0	0.14100	0.21130	
565	155.00	1731.0	0.11660	0.19220	
566	126.70	1124.0	0.11390	0.30940	
567	184.60	1821.0	0.16500	0.86810	
568	59.16	268.6	0.08996	0.06444	

	worst concavity	worst concave points	worst symmetry \
0	0.7119	0.2654	0.4601
1	0.2416	0.1860	0.2750
2	0.4504	0.2430	0.3613
3	0.6869	0.2575	0.6638
4	0.4000	0.1625	0.2364
..	...	...	...
564	0.4107	0.2216	0.2060
565	0.3215	0.1628	0.2572
566	0.3403	0.1418	0.2218
567	0.9387	0.2650	0.4087
568	0.0000	0.0000	0.2871

	worst fractal dimension
0	0.11890
1	0.08902
2	0.08758
3	0.17300
4	0.07678
..	...
564	0.07115
565	0.06637
566	0.07820
567	0.12400
568	0.07039

[569 rows x 30 columns]

In [59]:

```
print(Y)
```

```
0      0
1      0
2      0
3      0
4      0
..
564    0
565    0
566    0
567    0
568    1
```

Name: label, Length: 569, dtype: int64

## TRAIN\_TEST SPLIT

In [60]:

```
X_train , X_test , Y_train , Y_test = train_test_split(X,Y,test_size=.2,random_state=42)
```

In [61]:

```
X.shape,X_train.shape,X_test.shape
```

Out[61]:

```
((569, 30), (455, 30), (114, 30))
```

## STANDAEDIZE THE DATA

In [62]:

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

In [63]:

```
scaler = StandardScaler()
X_train_std = scaler.fit_transform(X_train)
```



```
X_test_std = scaler.fit_transform(X_test)
```

# BUILDING A NEURAL NETWORK

In [64]:

```
import tensorflow as tf
tf.random.set_seed(3)
from tensorflow import keras
```

TensorFlow is a deep learning library developed by Google, and it's used quite widely to build neural networks because of its extensive functionality. Keras is a wrapper for TensorFlow, often referred to as such, which makes building neural networks much easier with its very simple API. Until the appearance of TensorFlow and Keras, neural networks were considerably harder to develop.

As mentioned earlier, during the training process in neural networks, the weights and parameters get initialized randomly; hence, for each run of the model, there may be minute changes in the values of accuracy. We used `random.set_seed(3)`, which fixes this random initialization and thus always presents the same accuracy score over different runs for reproducibility.

## SETTING KERAS NETWORK (CREATING LAYERS OF NN IE.INPUT LAYER,HIDDEN LAYER AND OUTPUT LAYER)

In [65]:

```
model = keras.Sequential([
    keras.layers.Flatten(input_shape=(30,)),
    keras.layers.Dense(40,activation = 'relu'),
    keras.layers.Dense(2,activation = 'sigmoid')
])
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/reshaping/flatten.py:37: UserWarning: Do not pass an `input_shape`/`input_dim` argument to a layer. When using Sequential models, prefer using an `Input(shape)` object as the first layer in the model instead.
  super().__init__(**kwargs)
```

**Flatten** is used to transform the data of `X_train` and `X_test` into a uni-dimensional array. The number `30` indicates the number of input features, that is, the 30 columns in the dataset.

In the **hidden layer**, the number `40` represents the number of neurons in that layer. For the **output layer**, 2 neurons are used because the number of neurons is equal to the number of classes in the target variable. This process is called the **firing of neurons**, where if one neuron outputs `1` (for example, for class 0), the other neuron will not fire (output `0`) and vice versa. This setup ensures accurate classification.

## COMPLING THE NEURAL NETWORK

In [66]:

```
model.compile(optimizer='adam',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])
```

### NOTE:









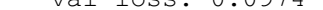











while using categorical variables in dataet like 0,1 we use 'sparse\_categorical\_crossentropy' and when we have one hot encoded labels we have 'categorical\_crossentropy'

## TRAINING THE NEURAL NETWORK

In [67]:

```
history = model.fit(X_train_std,Y_train , validation_split =.1 ,epochs=20)
```

```

Epoch 1/20
13/13  1s 23ms/step - accuracy: 0.7222 - loss: 0.5641 - val_accuracy: 0.8261 - val_loss: 0.3642
Epoch 2/20
13/13  0s 5ms/step - accuracy: 0.8859 - loss: 0.2999 - val_accuracy: 0.8913 - val_loss: 0.2458
Epoch 3/20
13/13  0s 6ms/step - accuracy: 0.9237 - loss: 0.2148 - val_accuracy: 0.9348 - val_loss: 0.1891
Epoch 4/20
13/13  0s 5ms/step - accuracy: 0.9467 - loss: 0.1732 - val_accuracy: 0.9565 - val_loss: 0.1572
Epoch 5/20
13/13  0s 5ms/step - accuracy: 0.9503 - loss: 0.1478 - val_accuracy: 0.9565 - val_loss: 0.1369
Epoch 6/20
13/13  0s 4ms/step - accuracy: 0.9564 - loss: 0.1301 - val_accuracy: 0.9783 - val_loss: 0.1229
Epoch 7/20
13/13  0s 6ms/step - accuracy: 0.9564 - loss: 0.1166 - val_accuracy: 1.0000 - val_loss: 0.1123
Epoch 8/20
13/13  0s 7ms/step - accuracy: 0.9646 - loss: 0.1059 - val_accuracy: 1.0000 - val_loss: 0.1041
Epoch 9/20
13/13  0s 4ms/step - accuracy: 0.9696 - loss: 0.0972 - val_accuracy: 1.0000 - val_loss: 0.0974
Epoch 10/20
13/13  0s 4ms/step - accuracy: 0.9747 - loss: 0.0898 - val_accuracy: 1.0000 - val_loss: 0.0918
Epoch 11/20
13/13  0s 4ms/step - accuracy: 0.9795 - loss: 0.0834 - val_accuracy: 1.0000 - val_loss: 0.0871
Epoch 12/20
13/13  0s 4ms/step - accuracy: 0.9795 - loss: 0.0780 - val_accuracy: 1.0000 - val_loss: 0.0832
Epoch 13/20
13/13  0s 5ms/step - accuracy: 0.9873 - loss: 0.0733 - val_accuracy: 1.0000 - val_loss: 0.0798
Epoch 14/20
13/13  0s 4ms/step - accuracy: 0.9873 - loss: 0.0690 - val_accuracy: 1.0000 - val_loss: 0.0768
Epoch 15/20
13/13  0s 4ms/step - accuracy: 0.9873 - loss: 0.0653 - val_accuracy: 1.0000 - val_loss: 0.0743
Epoch 16/20
13/13  0s 5ms/step - accuracy: 0.9873 - loss: 0.0619 - val_accuracy: 1.0000 - val_loss: 0.0721
Epoch 17/20
13/13  0s 5ms/step - accuracy: 0.9873 - loss: 0.0588 - val_accuracy: 1.0000 - val_loss: 0.0701
Epoch 18/20
13/13  0s 5ms/step - accuracy: 0.9895 - loss: 0.0561 - val_accuracy: 1.0000 - val_loss: 0.0683
Epoch 19/20
13/13  0s 5ms/step - accuracy: 0.9869 - loss: 0.0536 - val_accuracy: 1.0000 - val_loss: 0.0667
Epoch 20/20
13/13  0s 4ms/step - accuracy: 0.9869 - loss: 0.0512 - val_accuracy: 1.0000 - val_loss: 0.0653

```

We can observe in the training process that when **loss** decreases, the **accuracy** score increases. It shows that as the model is learning, it is getting better in classifying the data, which is reflected by the decrease in loss and an increase in accuracy. Moreover, the **validation accuracy** is quite high, which shows that the model performs well on unseen data.

However, in the above example, it is necessary to **standardize** the data before training by using a technique such as **StandardScaler**. Standardizing the data ensures that all features have a similar scale, which helps the model converge faster and improves overall performance. This is especially important for neural networks, where feature scaling can significantly impact training efficiency and model accuracy.

## visualizing accuracy and loss

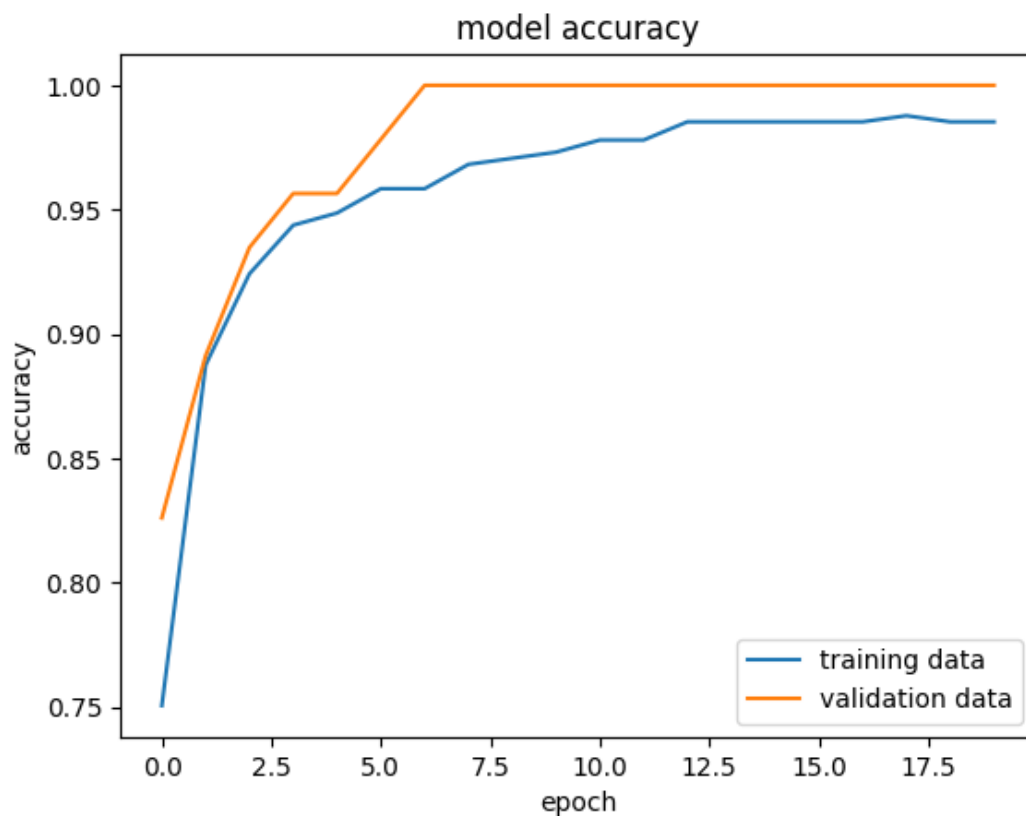
below are the plots for loss and accuracy vs epoch

In [68]:

```
plt.plot(history.history['accuracy'])
plt.plot(history.history['val_accuracy'])
plt.title('model accuracy')
plt.ylabel('accuracy')
plt.xlabel('epoch')
plt.legend(['training data', 'validation data'], loc = 'lower right')
```

Out[68]:

<matplotlib.legend.Legend at 0x7c7d39fc24a0>



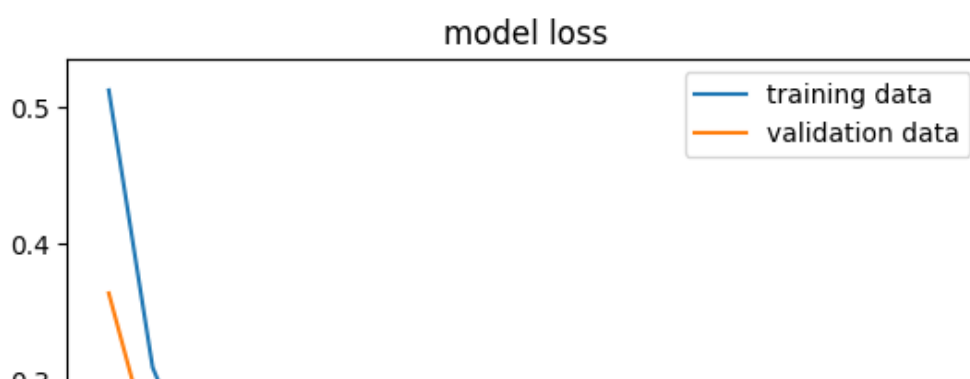
In [69]:

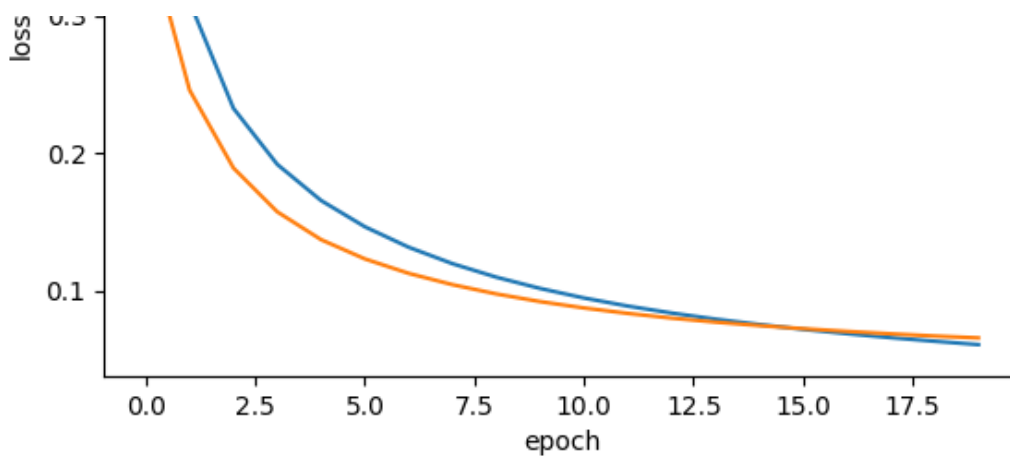
```
plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])

plt.title('model loss')
plt.ylabel('loss')
plt.xlabel('epoch')
plt.legend(['training data', 'validation data'], loc = 'upper right')
```

Out[69]:

<matplotlib.legend.Legend at 0x7c7d49d41b70>





## ACCURACY ON TEST DATA

In [70]:

```
loss, accuracy = model.evaluate(X_test_std, Y_test)
```

4/4 ————— 0s 4ms/step - accuracy: 0.9794 - loss: 0.0762

In [71]:

```
print(accuracy)
```

0.9824561476707458

In [72]:

```
print(loss)
```

0.0646393746137619

In [73]:

```
print(X_test_std.shape)
```

(114, 30)

In [74]:

```
print(X_test_std[0])
```

```
[-0.4877952 -0.25088379 -0.46378664 -0.51543986  0.05784012 -0.0262922
 -0.10351188 -0.31929301  0.40877389  0.11968017 -0.0858903 -0.39144008
 -0.22090758 -0.2944085  -0.10670318 -0.38822597 -0.05805714 -0.18386347
 -0.30097441 -0.05389032 -0.29186744 -0.2492245  -0.34337233 -0.37693497
  0.32080655 -0.13570975  0.0219982  -0.21989091  0.23083947  0.17173277]
```

In [75]:

```
Y_pred = model.predict(X_test_std)
```

4/4 ————— 0s 12ms/step

Here 'model.predict' gives Prediction Probability of each class for that data point

In [76]:

```
print(Y_pred.shape)
```

(114, 2)

In [77]:

```
print(Y_pred[0])
```

[0.36871758 0.90940666]

In [78]:

```
print(X_test_std)
```

```
[[-0.4877952 -0.25088379 -0.46378664 ... -0.21989091  0.23083947
  0.17173277]
 [ 1.37325734  0.36318719  1.29254295 ...  0.93019499 -0.58958803
 -0.972629   ]
 [ 0.37225845 -0.05148067  0.38772954 ...  0.52157274 -0.08280128
 -0.21536971]
 ...
 [-0.76105639 -1.08248546 -0.76208585 ... -0.30042664 -0.3893541
 -0.32657562]
 [ 0.01558068  1.84511495  0.00969106 ... -0.50681802 -1.76086788
 -0.33345979]
 [ 1.88813896  2.58607883  1.79246268 ...  0.73702811 -0.02786985
 -0.10998887]]
```

In [79]:

```
print(Y_pred)
```

```
[[3.68717581e-01 9.09406662e-01]
 [9.49734807e-01 7.82909710e-03]
 [7.29148269e-01 9.83393714e-02]
 [2.69956619e-01 9.90913510e-01]
 [1.73727691e-01 9.97892797e-01]
 [9.99541104e-01 5.48638927e-05]
 [9.93384242e-01 4.11466142e-04]
 [6.52125776e-01 2.39761755e-01]
 [6.35126650e-01 8.01352561e-01]
 [1.03908285e-01 9.93952036e-01]
 [3.97986621e-01 9.39810693e-01]
 [7.58947074e-01 1.29158661e-01]
 [2.69596845e-01 9.51986790e-01]
 [6.08443141e-01 2.40193516e-01]
 [2.14384705e-01 9.93139327e-01]
 [9.01194811e-01 2.61751581e-02]
 [1.35579452e-01 9.70830321e-01]
 [2.72782177e-01 9.99250591e-01]
 [3.12361091e-01 9.99953985e-01]
 [9.57143962e-01 4.34301095e-03]
 [3.58143598e-01 8.58646989e-01]
 [3.98802489e-01 9.81752574e-01]
 [9.96306062e-01 4.00131918e-04]
 [1.42152593e-01 9.99471426e-01]
 [2.04580054e-01 9.97151375e-01]
 [1.39170587e-01 9.84659612e-01]
 [1.44612283e-01 9.90306020e-01]
 [1.75287321e-01 9.94423091e-01]
 [2.51135588e-01 9.85127568e-01]
 [9.91486132e-01 4.28293226e-03]
 [2.44109750e-01 9.97621357e-01]
 [1.20472841e-01 9.97034967e-01]
 [2.45707691e-01 9.95579302e-01]
 [2.31070369e-01 9.63147998e-01]
 [2.97606260e-01 9.98362780e-01]
 [3.10764879e-01 9.87725317e-01]
 [6.89205289e-01 3.55033189e-01]
 [1.19845495e-01 9.84520197e-01]
 [9.30966496e-01 1.28668100e-02]
 [3.33247572e-01 8.59483004e-01]
 [2.32073113e-01 9.98748541e-01]
 [7.99586177e-01 6.81976974e-02]
 [2.84895897e-01 9.83313203e-01]
 [1.89191088e-01 9.94748175e-01]
 [2.87213981e-01 8.36745203e-01]
 [3.88636976e-01 9.70685661e-01]
 [4.43138897e-01 9.97547388e-01]
 [2.17191473e-01 9.98739541e-01]
 [5.35536528e-01 9.72234130e-01]]
```

```
[9.26552191e-02 9.91803646e-01]
[9.15397942e-01 1.68135520e-02]
[9.89276230e-01 1.70424394e-03]
[7.37309039e-01 8.75177383e-01]
[1.50190189e-01 8.90652359e-01]
[1.58921897e-01 9.97007370e-01]
[1.90719977e-01 9.76380706e-01]
[2.18253106e-01 9.97604430e-01]
[9.99399364e-01 2.21786922e-05]
[5.73976874e-01 4.10342872e-01]
[1.29326105e-01 9.97449756e-01]
[3.00388932e-01 9.78869557e-01]
[9.72675323e-01 4.45455639e-03]
[9.77223098e-01 8.09118326e-04]
[2.67274857e-01 9.39317882e-01]
[1.95790946e-01 9.95521903e-01]
[2.50207543e-01 9.56876695e-01]
[9.48473930e-01 1.04553411e-02]
[9.97850418e-01 8.08905519e-04]
[3.81773025e-01 9.95706320e-01]
[3.78076702e-01 9.55845773e-01]
[8.34117353e-01 1.15504995e-01]
[9.32898521e-01 7.01000020e-02]
[1.97022721e-01 9.81200039e-01]
[8.39150906e-01 6.50138780e-02]
[1.82290182e-01 9.99854863e-01]
[3.76961142e-01 9.57012296e-01]
[2.73487031e-01 9.37870979e-01]
[5.60811222e-01 6.47986591e-01]
[1.01737000e-01 9.99294698e-01]
[4.64193881e-01 9.74773765e-01]
[8.67051184e-01 9.24002752e-02]
[1.46397665e-01 9.99150157e-01]
[5.58540404e-01 3.93063664e-01]
[9.76572633e-01 1.33499876e-03]
[8.52360785e-01 6.32489771e-02]
[7.72722006e-01 1.06965967e-01]
[9.91553724e-01 1.03936143e-01]
[8.43380690e-01 4.01753820e-02]
[4.21411358e-02 9.98379052e-01]
[2.66970366e-01 9.64466214e-01]
[3.62523645e-01 9.90390599e-01]
[6.35508895e-01 7.32306659e-01]
[2.54293740e-01 7.98824787e-01]
[1.93164989e-01 9.98160183e-01]
[5.32054484e-01 9.95406032e-01]
[2.59332120e-01 9.97613966e-01]
[9.69400644e-01 2.31236150e-03]
[9.85464156e-01 9.27838776e-03]
[1.61653757e-01 9.97992814e-01]
[9.35303867e-01 1.99486967e-02]
[9.08898532e-01 6.85537681e-02]
[1.22395575e-01 9.99907076e-01]
[9.53924596e-01 3.81063446e-02]
[8.63132238e-01 3.60085070e-02]
[2.75752455e-01 9.90567207e-01]
[4.25820380e-01 9.32571173e-01]
[2.46723190e-01 9.80000079e-01]
[9.97015119e-01 2.53661594e-04]
[3.17434072e-01 8.59048128e-01]
[3.43554467e-01 9.16883230e-01]
[7.76087940e-01 8.72754976e-02]
[1.99169278e-01 9.91953909e-01]
[5.47413409e-01 7.88669765e-01]
[9.99077260e-01 2.87298190e-05]]
```

**NOW USING 'argmax' FUNCTION**

In [80]:

```
my_list = [0.25, 0.56]
```

```
index_of_max_value = np.argmax(my_list)
print(my_list)
print(index_of_max_value)
```

```
[0.25, 0.56]
1
```

## NOW CONVERT PREDICTION PROBABILITY TO CLASS LABELS

In [81]:

```
Y_pred_labels = [np.argmax(i) for i in Y_pred]
print(Y_pred_labels)
```

```
[1, 0, 0, 1, 1, 0, 0, 0, 1, 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0,
1, 1, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 0, 0, 1, 1, 1, 1, 1, 0, 0, 1,
1, 0, 0, 1, 1, 1, 0, 0, 1, 1, 0, 0, 1, 0, 1, 1, 1, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1,
1, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 0, 1, 1, 0]
```

## BUILDING THE PREDICTIVE SYSTEM

In [82]:

```
input_data = (11.76,21.6,74.72,427.9,0.08637,0.04966,0.01657,0.01115,0.1495,0.05888,0.40
62,1.21,2.635,28.47,0.005857,0.009758,0.01168,0.007445,0.02406,0.001769,12.98,25.72,82.98
,516.5,0.1085,0.08615,0.05523,0.03715,0.2433,0.06563)
```

```
#CONVERT INPUT DATA INTO NUMPY ARRAY
```

```
input_data_as_numpy_array = np.asarray(input_data)
```

```
#RESHAPE THE NUMPY ARRAY AS WE ARE PREDICTING FOR ONE DATA POINT
```

```
input_data_resaped = input_data_as_numpy_array.reshape(1,-1)
```

```
#STANDARDIZING THE INPUT DATA
```

```
input_data_std = scaler.transform(input_data_resaped)
```

```
prediction = model.predict(input_data_std)
```

```
print(prediction)
```

```
prediction_label = [np.argmax(prediction)]
```

```
print(prediction_label)
```

```
if(prediction_label[0] == 0):
```

```
    print('The tumor is Malignant')
```

```
else:
```

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

```
1/1  0s 21ms/step
```

```
[[0.19964646 0.98835504]]
```

```
[1]
```

```
The tumor is Benign
```

```
/usr/local/lib/python3.10/dist-packages/sklearn/utils/validation.py:2739: UserWarning: X
does not have valid feature names, but StandardScaler was fitted with feature names
warnings.warn(
```

## Conclusion

The project successfully built a neural network to classify breast tumors as malignant or benign using the Breast Cancer Wisconsin Dataset. The preprocessing of data, standardizing, and using a neural network model helped us achieve a high accuracy in classification. The performance of the model clearly showed the inverse relationship between loss and accuracy, as expected, where the loss decreased and accuracy increased in each epoch.

The use of **StandardScaler** ensured the data was standardized, which helped improve the model's convergence and overall performance. The project shows the effectiveness of machine learning, especially neural networks, in assisting medical diagnosis, particularly for early detection of cancerous cases of the breast, and also points

out the importance of preprocessing the data to get reliable and precise results.