

In [9]:

```
"""
Importing the Dependencies
"""

import numpy as np
import pandas as pd
import sklearn.datasets
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score

"""Data Collection & Processing"""

# loading the data from sklearn
breast_cancer_dataset = sklearn.datasets.load_breast_cancer()

print(breast_cancer_dataset)

# loading the data to a data frame
data_frame = pd.DataFrame(breast_cancer_dataset.data, columns = breast_cancer_dataset.fe

# print the first 5 rows of the dataframe
data_frame.head()

# adding the 'target' column to the data frame
data_frame['label'] = breast_cancer_dataset.target

# print last 5 rows of the dataframe
data_frame.tail()

# number of rows and columns in the dataset
data_frame.shape

# getting some information about the data
data_frame.info()

# checking for missing values
data_frame.isnull().sum()

# statistical measures about the data
data_frame.describe()

# checking the distribution of Target Varibale
data_frame['label'].value_counts()

"""1 --> Benign

0 --> Malignant
"""

data_frame.groupby('label').mean()

"""Separating the features and target"""

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

print(X)

print(Y)

"""Splitting the data into training data & Testing data"""

X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2, random_state=2)
```

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print(X.shape, X_train.shape, X_test.shape)

"""Model Training

Logistic Regression
"""

model = LogisticRegression()

# training the Logistic Regression model using Training data

model.fit(X_train, Y_train)

"""Model Evaluation

Accuracy Score
"""

# accuracy on training data
X_train_prediction = model.predict(X_train)
training_data_accuracy = accuracy_score(Y_train, X_train_prediction)

print('Accuracy on training data = ', training_data_accuracy)

# accuracy on test data
X_test_prediction = model.predict(X_test)
test_data_accuracy = accuracy_score(Y_test, X_test_prediction)

print('Accuracy on test data = ', test_data_accuracy)

"""Building a Predictive System"""

input_data = (13.54,14.36,87.46,566.3,0.09779,0.08129,0.06664,0.04781,0.1885,0.05766,0.2

# change the input data to a numpy array
input_data_as_numpy_array = np.asarray(input_data)

# reshape the numpy array as we are predicting for one datapoint
input_data_reshaped = input_data_as_numpy_array.reshape(1,-1)

prediction = model.predict(input_data_reshaped)
print(prediction)

if (prediction[0] == 0):
    print('The Breast cancer is Malignant')

else:
    print('The Breast Cancer is Benign')

{'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,
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1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1]), 'frame': None, 'target
_names': array(['malignant', 'benign'], dtype='<U9'), 'DESCR': '.. _breast_cancer_datase
t:\n\nBreast cancer wisconsin (diagnostic) dataset\n-----
-----\n\n**Data Set Characteristics:**\n\n      :Number of Instances: 569\n\n      :Numbe
r of Attributes: 30 numeric, predictive attributes and the class\n\n      :Attribute Infor
mation:\n      - radius (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 variation in radius lengths)\n      - compactness
(perimeter^2 / area - 1.0)\n      - concavity (severity of concave portions of the con
tour)\n      - concave points (number of concave portions of the contour)\n      - s
ymmetry\n      - fractal dimension ("coastline approximation" - 1)\n\n      The mea
n, standard error, and "worst" or largest (mean of the three\n      worst/largest valu
es) 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-Be
nign\n\n      :Summary Statistics:\n\n      =====
=====
=====\n\n      Min      Max\n      =====
=====
=====\n      radius (mean):      6.981  28.11\n
texture (mean):      9.71  39.28\n      perimeter (mean):
      43.79  188.5\n      area (mean):      143.5  2501.0\n      smoothn
ess (mean):      0.053  0.163\n      compactness (mean):
0.019  0.345\n      concavity (mean):      0.0  0.427\n      concave points
(mean):      0.0  0.201\n      symmetry (mean):      0.106  0.
304\n      fractal dimension (mean):      0.05  0.097\n      radius (standard erro
r):      0.112  2.873\n      texture (standard error):      0.36  4.885\n
      perimeter (standard error):      0.757  21.98\n      area (standard error):
      6.802  542.2\n      smoothness (standard error):      0.002  0.031\n      compa
ctness (standard error):      0.002  0.135\n      concavity (standard error):
0.0  0.396\n      concave points (standard error):      0.0  0.053\n      symmetry (stan
dard error):      0.008  0.079\n      fractal dimension (standard error):      0.001
0.03\n      radius (worst):      7.93  36.04\n      texture (worst):
      12.02  49.54\n      perimeter (worst):      50.41  251.2\n
      area (worst):      185.2  4254.0\n      smoothness (worst):
      0.071  0.223\n      compactness (worst):      0.027  1.058\n      conca
vity (worst):      0.0  1.252\n      concave points (worst):
0.0  0.291\n      symmetry (worst):      0.156  0.664\n      fractal dimens
ion (worst):      0.055  0.208\n      =====
=====
=====\n\n      :Missing Attribute Values: None\n\n      :Class Distribution: 212 - Malignan
t, 357 - Benign\n\n      :Creator: Dr. William H. Wolberg, W. Nick Street, Olvi L. Mangas
arian\n\n      :Donor: Nick Street\n\n      :Date: November, 1995\n\nThis is a copy of UCI M
L Breast Cancer Wisconsin (Diagnostic) datasets.\nhttps://goo.gl/U2Uwz2\n\nFeatures are
computed from a digitized image of a fine needle\naspirate (FNA) of a breast mass. They
describe\ncharacteristics of the cell nuclei present in the image.\n\nSeparating plane d
escribed above was obtained using\nMultisurface Method-Tree (MSM-T) [K. P. Bennett, "Dec

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ision Tree\nConstruction Via Linear Programming." Proceedings of the 4th\nMidwest Artificial Intelligence and Cognitive Science Society, \npp. 97-101, 1992], a classification method which uses linear\nprogramming to construct a decision tree. Relevant features\nwere selected using an exhaustive search in the space of 1-4\nfeatures and 1-3 separating planes.\n\nThe actual linear program used to obtain the separating plane\nin the 3-dimensional space is that described in:\n[K. P. Bennett and O. L. Mangasarian: "Robust Linear\nProgramming 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.. topic:: References\n\n - W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction\nfor breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on\nElectronic Imaging: Science and Technology, volume 1905, pages 861-870,\nSan Jose, CA, 1993.\n - O.L. Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis and\nprognosis via linear programming. Operations Research, 43(4), pages 570-577,\nJuly-August 1995.\n - W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine learning techniques\nto diagnose breast cancer from fine-needle aspirates. Cancer Letters 77 (1994)\n163-171.', '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_cancer.csv', 'data_module': 'sklearn.datasets.data'}
<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	int32

dtypes: float64(30), int32(1)

memory usage: 135.7 KB

	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]
```

```

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: int32
```

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

```
Accuracy on training data = 0.9384615384615385
```

```
Accuracy on test data = 0.9298245614035088
```

```
[1]
```

```
The Breast Cancer is Benign
```

```

C:\Users\vakap\anaconda3\lib\site-packages\sklearn\linear_model\_logistic.py:814: Conver
genceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

```

```
Increase the number of iterations (max_iter) or scale the data as shown in:
```

```
https://scikit-learn.org/stable/modules/preprocessing.html
```

```
Please also refer to the documentation for alternative solver options:
```

```
https://scikit-learn.org/stable/modules/linear\_model.html#logistic-regression
```

```
n_iter_i = _check_optimize_result(
```

```

C:\Users\vakap\anaconda3\lib\site-packages\sklearn\base.py:450: UserWarning: X does not
have valid feature names, but LogisticRegression was fitted with feature names
warnings.warn(

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In [ ]:
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