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Title Of The Project : Heart Disease Diagnostic Analysis

Technologies : Data Science

Domain : Healthcare

Importing the Dependencie

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

Data Collection and Processing

```
# loading the csv data to a Pandas DataFrame
heart_data = pd.read_csv('/content/Heart Disease data.csv')
```

```
# Print the first 5 rows of the dataset
heart_data.head()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target
0	52	1	0	125	212	0	1	168	0	1.0	2	2	3	0
1	53	1	0	140	203	1	0	155	1	3.1	0	0	3	0
2	70	1	0	145	174	0	1	125	1	2.6	0	0	3	0
3	61	1	0	148	203	0	1	161	0	0.0	2	1	3	0
4	62	0	0	138	294	1	1	106	0	1.9	1	3	2	0

```
# number of rows and columns in the dataset
heart_data.shape
```

```
(1025, 14)
```

```
# Getting the info about the data
heart_data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1025 entries, 0 to 1024
Data columns (total 14 columns):
#   Column      Non-Null Count  Dtype
---  -
0    age         1025 non-null   int64
1    sex         1025 non-null   int64
2    cp          1025 non-null   int64
3    trestbps    1025 non-null   int64
4    chol        1025 non-null   int64
5    fbs         1025 non-null   int64
6    restecg     1025 non-null   int64
7    thalach     1025 non-null   int64
8    exang       1025 non-null   int64
9    oldpeak     1025 non-null   float64
10   slope       1025 non-null   int64
11   ca          1025 non-null   int64
12   thal        1025 non-null   int64
13   target      1025 non-null   int64
dtypes: float64(1), int64(13)
memory usage: 112.2 KB
```

```
# checking the missing values
heart_data.isnull().sum()
```

```
age      0
sex      0
cp       0
trestbps 0
chol     0
fbs      0
restecg  0
thalach  0
exang    0
oldpeak  0
```

```
slope      0
ca         0
thal       0
target     0
dtype: int64
```

```
# Statistical measures about the data
heart_data.describe()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	c
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	54.434146	0.695610	0.942439	131.611707	246.000000	0.149268	0.529756	149.114146	0.336585	1.000000
std	9.072290	0.460373	1.029641	17.516718	51.59251	0.356527	0.527878	23.005724	0.472772	1.000000
min	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	0.000000	71.000000	0.000000	0.000000
25%	48.000000	0.000000	0.000000	120.000000	211.000000	0.000000	0.000000	132.000000	0.000000	0.000000
50%	56.000000	1.000000	1.000000	130.000000	240.000000	0.000000	1.000000	152.000000	0.000000	0.000000
75%	61.000000	1.000000	2.000000	140.000000	275.000000	0.000000	1.000000	166.000000	1.000000	1.000000
max	77.000000	1.000000	3.000000	200.000000	564.000000	1.000000	2.000000	202.000000	1.000000	6.000000

```
#checking the distribution of Target Variable
heart_data['target'].value_counts()
```

```
1    526
0    499
Name: target, dtype: int64
```

1 for Defective Heart

0 for Healthy Heart

Splitting the Features and Target

```
X = heart_data.drop(columns='target', axis=1)
Y = heart_data['target']
```

```
print(X)
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak \
0	52	1	0	125	212	0	1	168	0	1.0
1	53	1	0	140	203	1	0	155	1	3.1
2	70	1	0	145	174	0	1	125	1	2.6
3	61	1	0	148	203	0	1	161	0	0.0
4	62	0	0	138	294	1	1	106	0	1.9
...
1020	59	1	1	140	221	0	1	164	1	0.0
1021	60	1	0	125	258	0	0	141	1	2.8
1022	47	1	0	110	275	0	0	118	1	1.0
1023	50	0	0	110	254	0	0	159	0	0.0
1024	54	1	0	120	188	0	1	113	0	1.4

	slope	ca	thal
0	2	2	3
1	0	0	3
2	0	0	3
3	2	1	3
4	1	3	2
...
1020	2	0	2
1021	1	1	3
1022	1	1	2
1023	2	0	2
1024	1	1	3

```
[1025 rows x 13 columns]
```

```
print(Y)
```

```
0    0
1    0
2    0
3    0
4    0
..
1020 1
```

```

1021    0
1022    0
1023    1
1024    0
Name: target, Length: 1025, dtype: int64

```

Splitting the Data into Training data & Test Data

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

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

```
(1025, 13) (820, 13) (205, 13)
```

Model Training

Logistic Regression

```
model = LogisticRegression()
```

```
# training the LogisticRegression model with Training data
model.fit(X_train, Y_train)
```

```

/usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge
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(
```

```
    LogisticRegression
```

```
    LogisticRegression())
```

Model Evaluation

Accuracy Score

```

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

```

```
print('Accuracy on Training data: ', training_data_accuracy)
```

```
Accuracy on Training data: 0.8524390243902439
```

```

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

```

```
print('Accuracy on Test data: ', test_data_accuracy)
```

```
Accuracy on Test data: 0.8048780487804879
```

Building a Predictive System

```
input_data = (43,0,0,132,341,1,0,136,1,3,1,0,3)
```

```

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

```

```

#reshape the numpy array as we are predicting for only on instance
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 person is healthy')
else:
    print('The person has Heart Disease')

```

```
[0]
```

```
The person is healthy
```

```
/usr/local/lib/python3.9/dist-packages/sklearn/base.py:439: UserWarning: X does not have valid feature names, but LogisticRegression
warnings.warn(
```



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