

## Importing libraries

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

## Data Collection

```
#loading a diabetes dataset to the pandas dataframe
heart_disease_df=pd.read_csv("/content/heart_disease_data.csv")
```

```
#print first 5 rows of te dataset
heart_disease_df.head()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target
0	63	1	3	145	233	1	0	150	0	2.3	0	0	1	1
1	37	1	2	130	250	0	1	187	0	3.5	0	0	2	1
2	41	0	1	130	204	0	0	172	0	1.4	2	0	2	1
3	56	1	1	120	236	0	1	178	0	0.8	2	0	2	1
4	57	0	0	120	354	0	1	163	1	0.6	2	0	2	1

```
heart_disease_df.tail()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target
298	57	0	0	140	241	0	1	123	1	0.2	1	0	3	0
299	45	1	3	110	264	0	1	132	0	1.2	1	0	3	0
300	68	1	0	144	193	1	1	141	0	3.4	1	2	3	0
301	57	1	0	130	131	0	1	115	1	1.2	1	1	3	0
302	57	0	1	130	236	0	0	174	0	0.0	1	1	2	0

```
# checking no.of rows and columns in the dataset
heart_disease_df.shape
```

```
(303, 14)
```

```
#getting info about the dataset
heart_disease_df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):
 #   Column      Non-Null Count  Dtype
---  -
 0   age         303 non-null    int64
 1   sex         303 non-null    int64
 2   cp          303 non-null    int64
 3   trestbps    303 non-null    int64
 4   chol        303 non-null    int64
 5   fbs         303 non-null    int64
 6   restecg     303 non-null    int64
 7   thalach     303 non-null    int64
 8   exang       303 non-null    int64
 9   oldpeak     303 non-null    float64
10   slope       303 non-null    int64
11   ca          303 non-null    int64
12   thal        303 non-null    int64
13   target      303 non-null    int64
dtypes: float64(1), int64(13)
memory usage: 33.3 KB
```

```
# checking for missing value
heart_disease_df.isnull().sum()
```

```

      0
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 measure about the dataset
heart_disease_df.describe()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	s
count	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000
mean	54.366337	0.683168	0.966997	131.623762	246.264026	0.148515	0.528053	149.646865	0.326733	1.039604	1.395000
std	9.082101	0.466011	1.032052	17.538143	51.830751	0.356198	0.525860	22.905161	0.469794	1.161075	0.610000
min	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	0.000000	71.000000	0.000000	0.000000	0.000000
25%	47.500000	0.000000	0.000000	120.000000	211.000000	0.000000	0.000000	133.500000	0.000000	0.000000	1.000000
50%	55.000000	1.000000	1.000000	130.000000	240.000000	0.000000	1.000000	153.000000	0.000000	0.800000	1.000000
75%	61.000000	1.000000	2.000000	140.000000	274.500000	0.000000	1.000000	166.000000	1.000000	1.600000	2.000000
max	77.000000	1.000000	3.000000	200.000000	564.000000	1.000000	2.000000	202.000000	1.000000	6.200000	2.000000

```
heart_disease_df['target'].value_counts()
```

```

      count
target
1         165
0         138

```

dtype: int64

1 -> heart disease 0 -> no heart disease

Splitting the features and target

```
x = heart_disease_df.drop(columns='target',axis=1)
y = heart_disease_df['target']
```

x,y

```

(   age  sex  cp  trestbps  chol  fbs  restecg  thalach  exang  oldpeak  \
0     63   1   3     145    233   1         0     150     0       2.3
1     37   1   2     130    250   0         1     187     0       3.5

```

```

2    41    0    1    130    204    0    0    172    0    1.4
3    56    1    1    120    236    0    1    178    0    0.8
4    57    0    0    120    354    0    1    163    1    0.6
...    ...    ..    ..    ...    ...    ...    ...    ...    ...
298  57    0    0    140    241    0    1    123    1    0.2
299  45    1    3    110    264    0    1    132    0    1.2
300  68    1    0    144    193    1    1    141    0    3.4
301  57    1    0    130    131    0    1    115    1    1.2
302  57    0    1    130    236    0    0    174    0    0.0

```

```

      slope  ca  thal
0         0  0    1
1         0  0    2
2         2  0    2
3         2  0    2
4         2  0    2
..      ...  ..   ...
298      1  0    3
299      1  0    3
300      1  2    3
301      1  1    3
302      1  1    2

```

```

[303 rows x 13 columns],
0      1
1      1
2      1
3      1
4      1
..
298    0
299    0
300    0
301    0
302    0
Name: target, Length: 303, dtype: int64)

```

```

# splitting the data into train and test
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)
```

```
(303, 13) (242, 13) (61, 13)
```

## Model Training

### Logistic regression

```
model = LogisticRegression()
```

```
model.fit(x_train,y_train)
```

```

/usr/local/lib/python3.12/dist-packages/sklearn/linear_model/_logistic.py:465: ConvergenceWarning: lbfgs failed to converge (sta
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](https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression)

```
n_iter_i = _check_optimize_result(
```

▼ **LogisticRegression** ⓘ ?

```
LogisticRegression()
```

## Model Evaluation

### Accuracy Score

```

#Accuracy score on training data
x_train_prediction = model.predict(x_train)
training_data_accuracy = accuracy_score(x_train_prediction,y_train)

```

```
print("Accuracy score on training data :",training_data_accuracy*100)
```

```
Accuracy score on training data : 85.12396694214877
```

```
#Accuracy score on testing data
x_test_prediction = model.predict(x_test)
testing_data_accuracy = accuracy_score(x_test_prediction,y_test)
```

```
print("Accuracy score on testing data :",testing_data_accuracy*100)
```

```
Accuracy score on testing data : 81.9672131147541
```

### Predictive system

```
input = (68,1,0,144,193,1,1,141,0,3.4,1,2,3)

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

#reshape the array as we are predicting for one 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 don't have any heart disease")
else:
    print("The person is detective of heart disease")
```

```
[0]
The person don't have any heart disease
/usr/local/lib/python3.12/dist-packages/sklearn/utils/validation.py:2739: UserWarning: X does not have valid feature names, but
warnings.warn(
```

### Saving the trained model

```
import pickle
```

```
filename = 'heart_disease_model.sav'
pickle.dump(model,open(filename,'wb'))
```

```
#loaded the saved model
loaded_model = pickle.load(open('heart_disease_model.sav','rb'))
```

```
for column in x.columns:
    print(column)
```

```
age
sex
cp
trestbps
chol
fbs
restecg
thalach
exang
oldpeak
slope
ca
thal
```

