

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	fb	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	fb	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	slope
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	1.393604
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()
```

target	count
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

```
n_iter_i = _check_optimize_result(
```

```
  ▾ LogisticRegression (i) (?)
```

```
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
```

