


Importing the dependencies (i.e, libraries and functions)

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

data collection preprocessing

```
heart_data=pd.read_csv("/heart_disease_data.csv") #reading csv file
###heart_data=pd.read_csv("/heart.csv",skiprows=5) #reading csv file and skipping five ro
```

```
heart_data.head()
```



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

```
heart_data.tail()
```

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

```
heart_data.shape
```

```
(303, 14)
```

```
heart_data.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

```
#to check wheteher our dataset contains any missing value or not
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
```

```
#to check statistical data in dataset
heart_data.describe()
```

```

age      sex      cp      trestbps      chol      fbs      restecg
#checking the distribution of target variable
heart_data['target'].value_counts()

1      165
0      138
Name: target, dtype: int64

```

1---->heart disease exists

0---->heart disease does not exists [link text](#)

splitting the feature and target columns

```

X=heart_data.drop(columns='target',axis =1) #when we remove column then we write axis=1 and
Y=heart_data['target']

```

```
print(X)
```

```

      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]

```

```
print(Y)
```

```

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 training and testing data

```

X_train,X_test,Y_train,Y_test=train_test_split(X,Y,test_size=0.2,stratify=Y,random_state=2
#by the use of stratify our target data is uniformly divided otherwise it might happen tha

```

```
print(X.shape)
```

```
(303, 13)
```

```
print(X_train.shape)
```

```
(242, 13)
```

```
print(X_test.shape)
```

```
(61, 13)
```

```
print(Y.shape,Y_test.shape,Y_train.shape)
```

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

model training

logistic regression

```
model=LogisticRegression()
```

```
#training our logistic regression model with training data
```

```
model.fit(X_train,Y_train)
```

```

/usr/local/lib/python3.7/dist-packages/sklearn/linear_model/_logistic.py:818: 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

```
extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG,
LogisticRegression()
```

model evaluation

```
#model 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 is ",training_data_accuracy)
```

```
accuracy score on training data is  0.8512396694214877
```

```
#model accuracy on testing data
X_test_prediction=model.predict(X_test)
accuracy_x_test=accuracy_score(X_test_prediction,Y_test)
```

```
print("accuracy score of test data is:",accuracy_x_test)
```

```
accuracy score of test data is: 0.819672131147541
```

build a predictive system

```
input_data=(52,1,2,172,199,1,1,162,0,0.5,2,0,3)
#change the input array to numpy array
input_data_to_numpy=np.asarray(input_data)
#reshape the numpy array as we are predicting for only 1 instance
reshaped_data=input_data_to_numpy.reshape(1,-1)
prediction=model.predict(reshaped_data)
print(prediction)
if(prediction[0]==0):
    print("the person is having a heart disease")
else:
    print("the person dont have aheart disease")
```

```
[1]
the person dont have aheart disease
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:451: UserWarning: X does not have valid feature names, but"
```

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