

Heart-Disease-Prediction

Given clinical parameters about a patient, can we predict whether or not they have heart disease?

MAJOR PROJECT 1

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Predicting heart disease using machine learning

Problem Definition

In a statement,

Given clinical parameters about a patient, can we predict whether or not they have heart disease?

Features

This is where you'll get different information about each of the features in your data. You can do this via doing your own research (such as looking at the links above) or by talking to a subject matter expert (someone who knows about the dataset).

Dataset Details:

1. age - age in years
2. sex - (1 = male; 0 = female)
3. cp - chest pain type
 - 0: Typical angina: chest pain related decrease blood supply to the heart
 - 1: Atypical angina: chest pain not related to heart
 - 2: Non-anginal pain: typically esophageal spasms (non heart related)
 - 3: Asymptomatic: chest pain not showing signs of disease
4. trestbps - resting blood pressure (in mm Hg on admission to the hospital) anything above 130-140 is typically cause for concern
5. chol - serum cholestoral in mg/dl
 - serum = LDL + HDL + .2 * triglycerides
 - above 200 is cause for concern
6. fbs - (fasting blood sugar > 120 mg/dl) (1 = true; 0 = false)
 - '>126' mg/dL signals diabetes
7. restecg - resting electrocardiographic results
 - 0: Nothing to note
 - 1: ST-T Wave abnormality
 - can range from mild symptoms to severe problems
 - signals non-normal heart beat
 - 2: Possible or definite left ventricular hypertrophy
 - Enlarged heart's main pumping chamber
8. thalach - maximum heart rate achieved
9. exang - exercise induced angina (1 = yes; 0 = no)
10. oldpeak - ST depression induced by exercise relative to rest looks at stress of heart during exercise unhealthy heart will stress more
11. slope - the slope of the peak exercise ST segment
 - 0: Upsloping: better heart rate with exercise (uncommon)
 - 1: Flatsloping: minimal change (typical healthy heart)
 - 2: Downsloping: signs of unhealthy heart

12. ca - number of major vessels (0-3) colored by flourosopy
- colored vessel means the doctor can see the blood passing through
 - the more blood movement the better (no clots)
13. thal - thalium stress result
- 1,3: normal
 - 6: fixed defect: used to be defect but ok now
 - 7: reversable defect: no proper blood movement when excercising
14. target - have disease or not (1=yes, 0=no) (= the predicted attribute)



Aniverthy/Heart-Disease-Prediction



<https://heartdiseasepredictbyaniverthy.herokuapp.com/>

ML CODE

```
#MAJOR PROJECT - 1
#Choose any dataset of your choice and apply a suitable CLASSIFIER/REGRESSOR and if
possible deploy it on heroku.
#NOTE: Heroku deployment is not compulsory
#Dataset = "https://github.com/Aniverthy/Heart-Disease-
Prediction/blob/main/heart.csv"
```

```
#step1:dataframe creation
import pandas as pd
import matplotlib.pyplot as plt
df=pd.read_csv("https://raw.githubusercontent.com/Aniverthy/Heart-Disease-
Prediction/main/heart.csv")
df
```

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
...
1020	59	1	1	140	221	0	1	164	1	0.0	2	0	2	1
1021	60	1	0	125	258	0	0	141	1	2.8	1	1	3	0
1022	47	1	0	110	275	0	0	118	1	1.0	1	1	2	0
1023	50	0	0	110	254	0	0	159	0	0.0	2	0	2	1
1024	54	1	0	120	188	0	1	113	0	1.4	1	1	3	0

```
#Step2: Exploratory Data Analysis
df.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

df.shape
```

```
(1025, 14)
```

```
df.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
```

```
df=df.drop(['slope'], axis=1)
```

```
df.describe()
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	ca
count	1025.000000	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				
std	9.072290	0.460373	1.029641	17.516718	51.59251	0.356527	0.527878				
min	29.000000	0.000000	0.000000	94.000000	126.00000	0.000000	0.000000				
25%	48.000000	0.000000	0.000000	120.000000	211.00000	0.000000	0.000000				
50%	56.000000	1.000000	1.000000	130.000000	240.00000	0.000000	1.000000				
75%	61.000000	1.000000	2.000000	140.000000	275.00000	0.000000	1.000000				
max	77.000000	1.000000	3.000000	200.000000	564.00000	1.000000	2.000000				

```
df.target.value_counts()
```

```
#gives no of person have heart disease
```

```
#1 represents having heart disease and 0 represents donot having heart disease
```

```
1    526
0    499
```

```
Name: target, dtype: int64
```

```
#Step 3: Data visualization
```

```
plt.figure(figsize=(15, 15))
```

```
for i, column in enumerate(df):
```

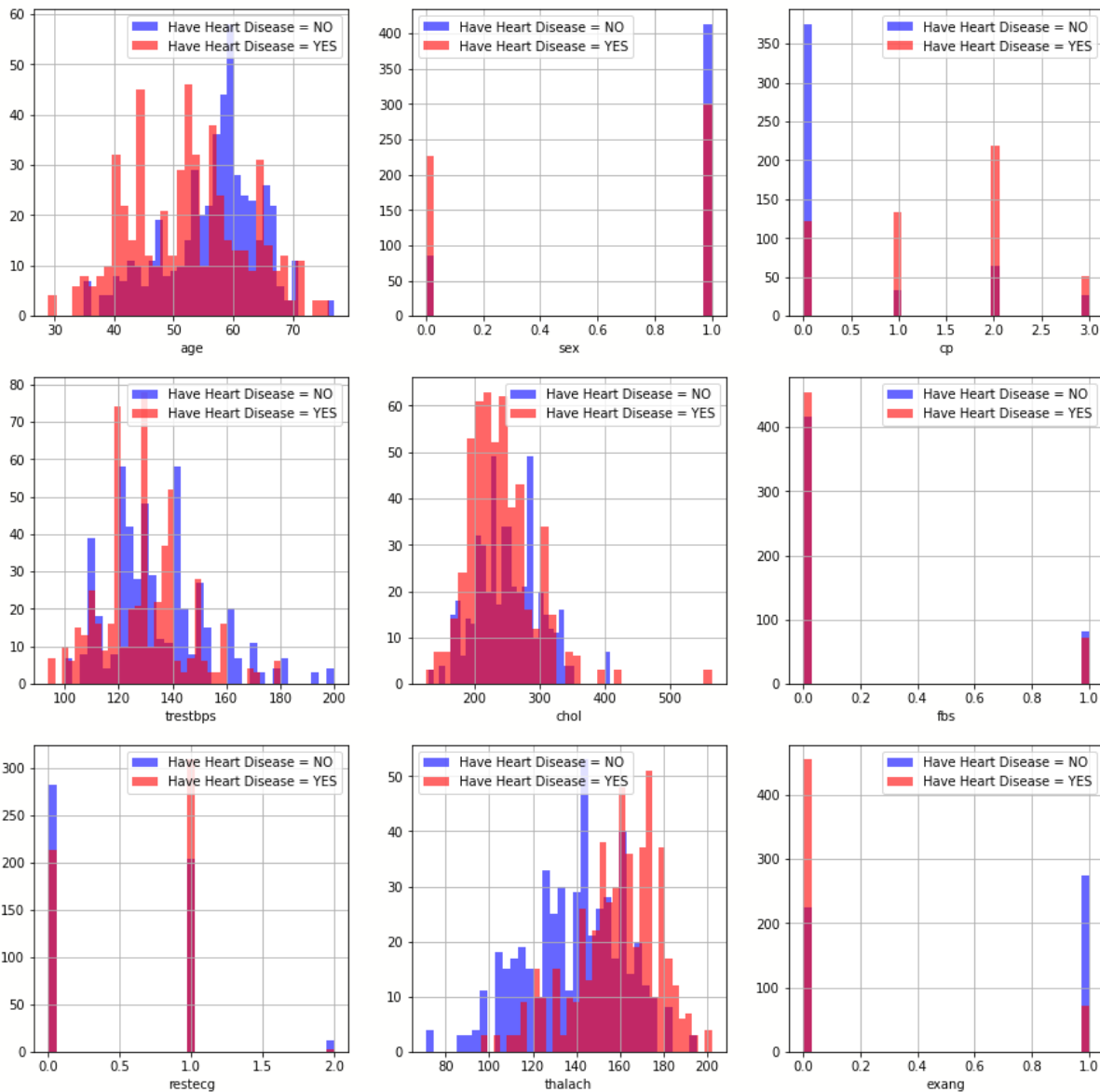
```
    try:
```

```
        plt.subplot(3, 3, i+1)
```

```

df[df["target"] == 0][column].hist(bins=35, color='blue', label='Have Heart Dis
ease = NO', alpha=0.6)
df[df["target"] == 1][column].hist(bins=35, color='red', label='Have Heart Dise
ase = YES', alpha=0.6)
plt.legend()
plt.xlabel(column)
except ValueError:
    break

```



```

# Create another figure
plt.figure(figsize=(10, 8))

# Scatter with postivie examples
plt.scatter(df.age[df.target==1],
            df.thalach[df.target==1],
            c="salmon")

# Scatter with negative examples
plt.scatter(df.age[df.target==0],
            df.thalach[df.target==0],
            c="lightblue")

```

```
# Add some helpful info
plt.title("Heart Disease in function of Age and Max Heart Rate")
plt.xlabel("Age")
plt.ylabel("Max Heart Rate")
plt.legend(["Disease", "No Disease"]);
```



```
#step 4 divide into input and output:
#step 5 TEST and TRAIN data
from sklearn.model_selection import train_test_split

x= df.drop('target', axis=1).values
y= df.target.values

x_train, x_test, y_train, y_test = train_test_split(x, y, random_state=42)

#step 6:SCALING or NORMALISATION -DONE ONLY FOR INPUTS
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
x_train = scaler.fit_transform(x_train)
x_test = scaler.fit_transform(x_test)
```



```
#step 7:model classification
from sklearn.linear_model import LogisticRegression
```

```
model = LogisticRegression()
```

```
#step 8: fitting the training values
```

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

```
LogisticRegression()
```

```
#step 9:predict the output
```

```
y_pred=model.predict(x_test)
```

```
y_pred
```

```
array([[1, 1, 0, 1, 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0,
        1, 0, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 1,
        0, 0, 0, 0, 1, 1, 1, 0, 1, 0, 1, 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0,
        1, 1, 0, 1, 1, 0, 0, 1, 1, 1, 0, 1, 1, 1, 0, 0, 0, 0, 1, 0, 1, 0,
        0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 0, 1, 0, 1, 1, 0, 1, 0, 1,
        1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1, 1, 0, 0, 0, 0, 1, 1, 1, 1, 1, 0,
        1, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1, 0, 1, 1,
        0, 0, 1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 0, 1, 1,
        1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0,
        1, 1, 1, 1, 1, 0, 0, 0, 1, 1, 0, 1, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0,
        1, 0, 0, 1, 1, 1, 1, 1, 0, 1, 1, 0, 1, 1, 0, 1, 1, 0, 0, 0, 1, 1,
        1, 0, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0, 1, 0])
```

```
y_test
```

```
array([[1, 1, 0, 1, 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 0, 0, 0, 1, 1, 0, 1, 0,
        0, 0, 1, 1, 1, 1, 0, 0, 0, 1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0,
        0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 1, 0, 1, 0, 0,
        1, 0, 0, 1, 0, 0, 0, 1, 1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 1, 0,
        0, 0, 0, 0, 1, 1, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1,
        0, 1, 1, 0, 1, 1, 0, 1, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 1, 1, 0,
        1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 1, 1,
        0, 0, 0, 1, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 1, 0, 1, 1, 0, 1, 1,
        1, 0, 0, 1, 1, 0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 0, 0, 0, 0, 1, 0, 0,
        1, 1, 1, 1, 1, 0, 0, 0, 1, 1, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 1, 0,
        1, 0, 0, 1, 1, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0,
        1, 0, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0])
```

```
#accuracy
test_score = accuracy_score(y_test, model.predict(x_test)) * 100
train_score = accuracy_score(y_train, model.predict(x_train)) * 100

results_df = pd.DataFrame(data=[["Logistic Regression", train_score, test_score]],
                           columns=['Model', 'Training Accuracy %', 'Testing Accuracy %'])
results_df
```

	Model Training	Accuracy %	Testing Accuracy %
0	Logistic Regression	86.328125	80.544747

```
#individual prediction
model.predict([x_train[10]])
#1 implies presence of heart disease
```

This is deployed in a Web APP using Heroku
Cloud Data



[Aniverthy/Heart-Disease-Prediction](https://github.com/Aniverthy/Heart-Disease-Prediction)



<https://heartdiseasepredictbyaniverthy.herokuapp.com/>