KNN_HEARTATTACK_PREDICTION

December 8, 2024

HEART ATTCK PREDICTION

[504]:

Heart Attack Prediction using machine learning involves using various factors (features) related to a person's health and medical history to predict whether they are at risk of a heart attack. The key goal is to create a model that can make predictions based on available data, which can help in preventive care and early diagnosis.

```
!unzip /content/archive (88).zip
      /bin/bash: -c: line 1: syntax error near unexpected token `('
      /bin/bash: -c: line 1: `unzip /content/archive (88).zip'
[505]: import pandas as pd
       import numpy as np
       import matplotlib.pyplot as plt
       import seaborn as sns
[506]: df=pd.read_csv('/content/archive (88).zip')
      UNDERSTANDING THE DATA
[507]: df
[507]:
            Age Sex ChestPainType
                                     RestingBP
                                                 Cholesterol
                                                               FastingBS RestingECG
                                                                              Normal
       0
             40
                   Μ
                                ATA
                                            140
                                                          289
       1
             49
                   F
                                NAP
                                            160
                                                          180
                                                                       0
                                                                              Normal
                                                                                  ST
       2
             37
                   Μ
                                ATA
                                            130
                                                          283
                                                                       0
       3
                                ASY
                                                          214
                                                                       0
                                                                              Normal
             48
                   F
                                            138
       4
                                NAP
                                            150
                                                          195
                                                                       0
                                                                              Normal
             54
                   Μ
                                                                              Normal
                                 TA
                                                          264
       913
             45
                   Μ
                                            110
                                                                       0
       914
             68
                                ASY
                                            144
                                                          193
                                                                       1
                                                                              Normal
                   М
       915
             57
                                ASY
                                           130
                                                          131
                                                                       0
                                                                              Normal
                   Μ
       916
                                ATA
                                            130
                                                          236
                                                                       0
                                                                                 LVH
             57
                   F
       917
             38
                   Μ
                                NAP
                                            138
                                                          175
                                                                       0
                                                                              Normal
            MaxHR ExerciseAngina
                                    Oldpeak ST_Slope
                                                       HeartDisease
               172
                                        0.0
       0
                                 Ν
                                                   Uр
               156
       1
                                        1.0
                                 N
                                                 Flat
                                                                   1
```

2	98 108		N Y	0.0 1.5	Up Flat		0 1
4	122		N	0.0	Up		0
	•••	•••	•••	•••		•••	
913	132		N	1.2	Flat		1
914	141		N	3.4	Flat		1
915	115		Y	1.2	Flat		1
916	174		N	0.0	Flat		1
917	173		N	0.0	Up		0

[918 rows x 12 columns]

[508]: df.shape

[508]: (918, 12)

[509]: df.dtypes

[509]: Age int64 Sex object ${\tt ChestPainType}$ object RestingBP int64 Cholesterol int64 FastingBS int64 RestingECG object MaxHR int64 ExerciseAngina object Oldpeak float64 ST_Slope object HeartDisease int64

dtype: object

[510]: df.describe()

[510]: FastingBS Age RestingBP Cholesterol MaxHR \ 918.000000 count 918.000000 918.000000 918.000000 918.000000 mean 53.510893 132.396514 0.233115 136.809368 198.799564 std 9.432617 18.514154 109.384145 0.423046 25.460334 min 28.000000 0.000000 0.000000 0.000000 60.000000 25% 47.000000 120.000000 173.250000 0.000000 120.000000 50% 54.000000 130.000000 223.000000 0.000000 138.000000 75% 60.000000 140.000000 267.000000 0.000000 156.000000 max 77.000000 200.000000 603.000000 1.000000 202.000000

Oldpeak HeartDisease count 918.000000 918.000000 mean 0.887364 0.553377

```
0.497414
std
         1.066570
min
        -2.600000
                        0.000000
25%
                        0.000000
         0.000000
50%
         0.600000
                        1.000000
75%
         1.500000
                        1.000000
max
         6.200000
                        1.000000
```

[511]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 918 entries, 0 to 917
Data columns (total 12 columns):

#	Column	Non-Null Count	Dtype
0	Age	918 non-null	int64
1	Sex	918 non-null	object
2	${\tt ChestPainType}$	918 non-null	object
3	RestingBP	918 non-null	int64
4	Cholesterol	918 non-null	int64
5	FastingBS	918 non-null	int64
6	RestingECG	918 non-null	object
7	MaxHR	918 non-null	int64
8	ExerciseAngina	918 non-null	object
9	Oldpeak	918 non-null	float64
10	ST_Slope	918 non-null	object
11	HeartDisease	918 non-null	int64
dtype	es: float64(1),	int64(6), object	(5)

dtypes. 110ato4(1), 111to4(0), object

memory usage: 86.2+ KB

[512]: df.corr(numeric_only=True)

[512]:		Age	${\tt RestingBP}$	Cholesterol	${\tt FastingBS}$	${\tt MaxHR}$	Oldpeak	\
	Age	1.000000	0.254399	-0.095282	0.198039	-0.382045	0.258612	
	RestingBP	0.254399	1.000000	0.100893	0.070193	-0.112135	0.164803	
	Cholesterol	-0.095282	0.100893	1.000000	-0.260974	0.235792	0.050148	
	FastingBS	0.198039	0.070193	-0.260974	1.000000	-0.131438	0.052698	
	MaxHR	-0.382045	-0.112135	0.235792	-0.131438	1.000000	-0.160691	
	Oldpeak	0.258612	0.164803	0.050148	0.052698	-0.160691	1.000000	
	HeartDisease	0.282039	0.107589	-0.232741	0.267291	-0.400421	0.403951	

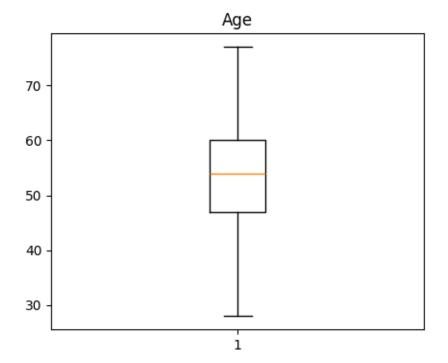
HeartDisease Age 0.282039 RestingBP 0.107589 Cholesterol -0.232741 FastingBS 0.267291 MaxHR -0.400421 Oldpeak 0.403951

HeartDisease 1.000000

DATA CLEANING

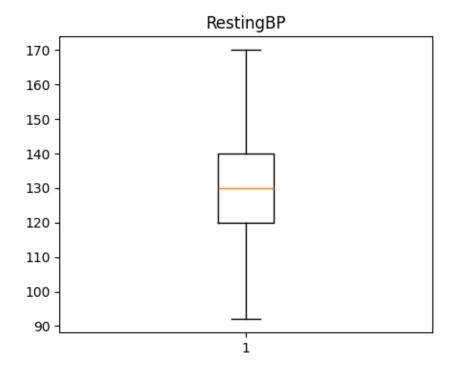
```
[513]: df.isna().sum()
                          0
[513]: Age
                          0
       Sex
       ChestPainType
                          0
       RestingBP
                          0
       Cholesterol
                          0
       FastingBS
                          0
       RestingECG
                          0
       MaxHR
                          0
       ExerciseAngina
                          0
       Oldpeak
                          0
       ST_Slope
                          0
       HeartDisease
                          0
       dtype: int64
[514]: df.duplicated().sum()
[514]: 0
[515]: df['Sex'].value_counts()
[515]: Sex
       М
            725
       F
            193
       Name: count, dtype: int64
[516]: df['ChestPainType'].value_counts()
[516]: ChestPainType
       ASY
              496
       NAP
              203
       ATA
              173
       TA
               46
       Name: count, dtype: int64
[517]: df['RestingECG'].value_counts()
[517]: RestingECG
       Normal
                 552
       LVH
                 188
       ST
                 178
       Name: count, dtype: int64
```

```
[518]: df['ExerciseAngina'].value_counts()
[518]: ExerciseAngina
      N
           547
       Y
            371
      Name: count, dtype: int64
[519]: df['ST_Slope'].value_counts()
[519]: ST_Slope
      Flat
               460
      Uр
               395
                63
      Down
      Name: count, dtype: int64
      OUTLIER DETECTION
[520]: import matplotlib.pyplot as plt
       plt.figure(figsize=(5,4))
      plt.boxplot(df['Age'])
      plt.title('Age')
      plt.show()
```

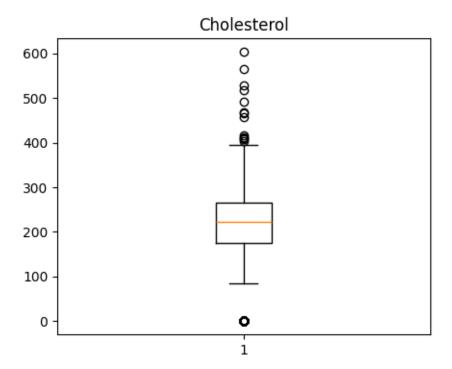


```
[521]: q1=df['RestingBP'].quantile(0.25) q1
```

```
[521]: 120.0
[522]: q3=df['RestingBP'].quantile(0.75)
[522]: 140.0
[523]: iqr=q3-q1
       min_range=q1-1.5*iqr
       print(min_range)
       max_range=q3+1.5*iqr
       print(max_range)
      90.0
      170.0
[524]: df=df[(df['RestingBP'] <= max_range) & (df['RestingBP'] >= min_range)]
[525]: import matplotlib.pyplot as plt
       plt.figure(figsize=(5,4))
       plt.boxplot(df['RestingBP'])
       plt.title('RestingBP')
       plt.show()
```

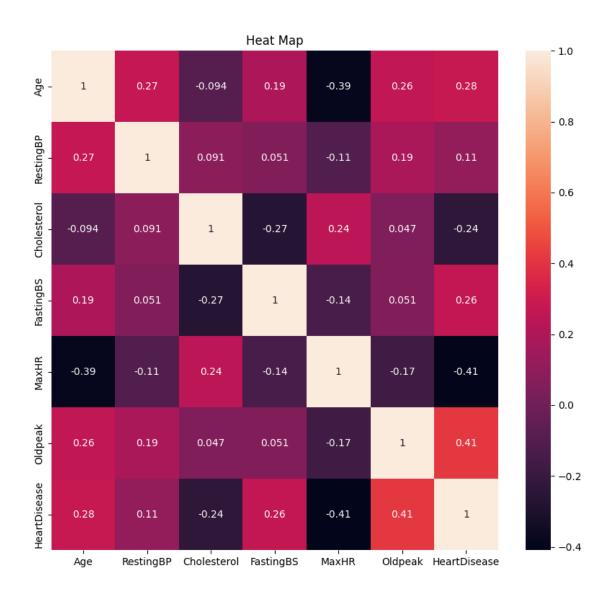


```
[526]: import matplotlib.pyplot as plt
plt.figure(figsize=(5,4))
plt.boxplot(df['Cholesterol'])
plt.title('Cholesterol')
plt.show()
```



HEAT MAP

```
[527]: plt.figure(figsize=(10,9))
  var1=df.corr(numeric_only=True)
  sns.heatmap(var1,annot=True)
  plt.title('Heat Map')
  plt.show()
```



CATEGORICAL VALUE ENCODING

```
[528]: obj1=[]
    for i in df:
        if df[i].dtype=='object':
            obj1.append(i)
[529]: obj1
```

[529]: ['Sex', 'ChestPainType', 'RestingECG', 'ExerciseAngina', 'ST_Slope']

ORDINAL ENCODING

```
[530]: from sklearn.preprocessing import OrdinalEncoder
       obj=OrdinalEncoder(categories=[['Normal','ST','LVH']])
       df['RestingECG']=obj.fit_transform(df[['RestingECG']])
      <ipython-input-530-e43b1fad3ba5>:3: SettingWithCopyWarning:
      A value is trying to be set on a copy of a slice from a DataFrame.
      Try using .loc[row_indexer,col_indexer] = value instead
      See the caveats in the documentation: https://pandas.pydata.org/pandas-
      docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
        df['RestingECG']=obj.fit transform(df[['RestingECG']])
[531]: from sklearn.preprocessing import OrdinalEncoder
       obj=OrdinalEncoder(categories=[['Up','Flat','Down']])
       df['ST_Slope']=obj.fit_transform(df[['ST_Slope']])
      <ipython-input-531-7b6285265aa4>:3: SettingWithCopyWarning:
      A value is trying to be set on a copy of a slice from a DataFrame.
      Try using .loc[row_indexer,col_indexer] = value instead
      See the caveats in the documentation: https://pandas.pydata.org/pandas-
      docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
        df['ST_Slope']=obj.fit_transform(df[['ST_Slope']])
      LABEL ENCODING
[532]: label_Enc=[]
       Onehot_Enc=[]
       for i in df:
         if df[i].dtype=='object' and df[i].nunique()>2:
           Onehot_Enc.append(i)
         elif df[i].dtype=='object' and df[i].nunique()<=2:</pre>
           label_Enc.append(i)
[533]: label_Enc
[533]: ['Sex', 'ExerciseAngina']
[534]: df['Sex']
[534]: 0
              Μ
              F
       1
       2
              Μ
       3
              F
              Μ
       913
             М
       914
              M
```

```
915
     Μ
     F
  916
  917
     Μ
  Name: Sex, Length: 890, dtype: object
[535]: from sklearn.preprocessing import LabelEncoder
  le=LabelEncoder()
  df['Sex'] = le.fit_transform(df['Sex'])
  <ipython-input-535-5473b90f9496>:3: SettingWithCopyWarning:
  A value is trying to be set on a copy of a slice from a DataFrame.
  Try using .loc[row_indexer,col_indexer] = value instead
  See the caveats in the documentation: https://pandas.pydata.org/pandas-
  docs/stable/user guide/indexing.html#returning-a-view-versus-a-copy
   df['Sex']=le.fit_transform(df['Sex'])
[536]: le.inverse_transform(df['Sex'])
'M', 'M', 'M', 'M', 'M', 'M',
       'F'. 'F'.
                       'M', 'M',
     'M', 'F', 'M', 'M', 'M', 'F', 'F', 'M',
       'M', 'F',
           'M',
     'M', 'M',
     'M', 'M', 'F',
           'M', 'F', 'M', 'M',
                   'M', 'M',
     'M', 'M', 'M', 'M',
             'M', 'M', 'M', 'M', 'F', 'M', 'M', 'M',
     'M', 'M',
     'M', 'M', 'M', 'M', 'M', 'M', 'M', 'F', 'M', 'F',
     'F', 'M',
           'M',
             'M', 'M', 'M',
                   'M', 'M', 'M', 'M',
                           'Μ',
```

```
'F', 'M', 'F',
'F'.
'M', 'M',
'F', 'M', 'M', 'M', 'F', 'M'], dtype=object)
```

```
[537]: from sklearn.preprocessing import LabelEncoder le_ex=LabelEncoder() df['ExerciseAngina']=le_ex.fit_transform(df['ExerciseAngina'])
```

<ipython-input-537-d4e7f18dc591>:3: SettingWithCopyWarning:

A value is trying to be set on a copy of a slice from a DataFrame. Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy df['ExerciseAngina']=le_ex.fit_transform(df['ExerciseAngina'])

[538]: le_ex.inverse_transform(df['Sex']) 'N'. 'N'. 'Y', 'N', 'Y', 'Y', 'Y', 'Y', 'Y', 'Y', 'N', 'Y', 'Y', 'N', 'Y', 'Y', 'N', 'Y', 'N', 'N', 'Y', 'Y', 'Y', 'Y',

```
'Y'.
'Y', 'N', 'Y',
'Y', 'Y',
'N', 'Y', 'Y', 'Y', 'N', 'Y'], dtype=object)
```

ONE HOT ENCODING

```
[542]:
            ChestPainType_ATA ChestPainType_NAP
                                                   ChestPainType_TA
                           1.0
                                              0.0
                                                                 0.0
       0
                                                                 0.0
       1
                           0.0
                                               1.0
       2
                           1.0
                                              0.0
                                                                 0.0
       3
                           0.0
                                              0.0
                                                                 0.0
       4
                           0.0
                                               1.0
                                                                 0.0
                                                                 1.0
       885
                           0.0
                                              0.0
       886
                           0.0
                                              0.0
                                                                 0.0
                                              0.0
                                                                 0.0
       887
                          0.0
       888
                           1.0
                                              0.0
                                                                 0.0
       889
                           0.0
                                              1.0
                                                                 0.0
       [890 rows x 3 columns]
[543]: df=df.drop(columns=Onehot_Enc)
[544]: df.reset_index(drop=True, inplace=True)
[545]: df=df.join(result)
[546]: x=df.drop(columns=['HeartDisease'])
       y=df['HeartDisease']
      SAMPLING
[547]: from imblearn.over_sampling import SMOTE
       sampler=SMOTE()
       x_resample,y_resample=sampler.fit_resample(x,y)
       y_resample.value_counts()
[547]: HeartDisease
            490
       1
            490
       Name: count, dtype: int64
      SCALING
[548]: from sklearn.preprocessing import MinMaxScaler
       minmax=MinMaxScaler()
       x_scaled=minmax.fit_transform(x_resample)
      DATA SPLITTING
[549]: from sklearn.model_selection import train_test_split
       x_train,x_test,y_train,y_test=train_test_split(x_scaled,y_resample,test_size=0.
        \hookrightarrow 2, random state=50)
```

```
[550]: x_train.shape
[550]: (784, 13)
[551]: x_test.shape
[551]: (196, 13)
[552]: y_train.shape
[552]: (784,)
[553]: (y_test.shape
[553]: (196,)
```

MODEL TRAINING

K NEAREST NEIGHBOR

K-Nearest Neighbors (KNN) is a simple, intuitive, and versatile machine learning algorithm used for both classification and regression tasks. It is based on the idea that similar data points tend to exist near each other in the feature space.

```
[554]: from sklearn.neighbors import KNeighborsClassifier knn_model=KNeighborsClassifier() knn_model.fit(x_test,y_test)
```

[554]: KNeighborsClassifier()

```
[555]: y_pred=knn_model.predict(x_test)
```

MODEL EVALUATION

```
[556]: from sklearn.metrics import

accuracy_score,classification_report,confusion_matrix

print(accuracy_score(y_test,y_pred))
```

0.8979591836734694

```
[557]: models=[KNeighborsClassifier()]
for model in models:
    model.fit(x_train,y_train)
    y_pred=model.predict(x_test)
    print(f"Accuracy :{accuracy_score(y_test,y_pred)}")
    print("-----")
    print(f"Confusion_matrix:\n {confusion_matrix(y_test,y_pred)}")
    print()
    print(f"classification_report:\n{classification_report(y_test,y_pred)}")
```

print("\n\n\n")

Accuracy :0.8520408163265306

Confusion_matrix:

[[84 15] [14 83]]

classification_report:

	precision	recall	f1-score	support
0	0.86	0.85	0.85	99
1	0.85	0.86	0.85	97
accuracy			0.85	196
macro avg	0.85	0.85	0.85	196
weighted avg	0.85	0.85	0.85	196

```
[558]: [jupyter nbconvert --to pdf /content/KNN_HEARTATTACK_PREDICTION (2).ipynb
```

```
/bin/bash: -c: line 1: syntax error near unexpected token `('
/bin/bash: -c: line 1: `jupyter nbconvert --to pdf
/content/KNN HEARTATTACK_PREDICTION (2).ipynb'
```

The heart attack prediction model, implemented using the K-Nearest Neighbors (KNN) algorithm, has achieved an 86% accuracy in predicting the likelihood of heart attacks. This is a promising result, especially considering the model's ability to effectively distinguish between the two classes (heart attack or no heart attack). The high accuracy can be attributed to the balanced nature of the dataset, which was achieved through the use of the SMOTE (Synthetic Minority Over-sampling Technique) technique. By applying SMOTE, the dataset was resampled to ensure that both classes (positive and negative heart attack cases) were equally represented, thereby preventing the model from being biased towards predicting the majority class.

KNN, being a distance-based algorithm, calculates the proximity of new data points to the training examples. It works well for this problem due to its simplicity and the fact that heart attack predictions are often influenced by a combination of numerical and categorical features, such as age, cholesterol levels, blood pressure, and chest pain type. These features, when properly preprocessed, allow the KNN model to classify a new patient's data based on similarities to known examples.

The performance of the model can be further evaluated using a classification report, which provides important metrics such as precision, recall, and F1-score. These metrics give a more detailed insight into how well the model performs in detecting both heart attack and non-heart attack cases. A high precision indicates that when the model predicts a heart attack, it is correct most of the time. On the other hand, a high recall suggests that the model is good at identifying most of the true

heart attack cases, reducing the chances of false negatives.

Overall, this model provides a reliable approach for heart attack prediction, which can be useful in clinical decision-making, potentially aiding healthcare professionals in making faster, data-driven decisions. While 86% accuracy is impressive, future improvements could involve fine-tuning the model's hyperparameters, exploring other algorithms, or incorporating additional features such as family history or lifestyle factors to further enhance prediction accuracy.