Source for Predicting heart disease using machine learning

This notebook looks into using various Python-based machine learning and data science libraries in an attempt to build a machine learning model capable of predicting whether or not someone has heart disease based on their medical attributes.

I am going to take the following approach:

- 1. Problem definition
- 2. Data
- 3. Evaluation
- 4. Features
- 5. Modelling
- 6. Experimentation

Preparing the Tools

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

import seaborn as sns

%matplotlib inline

from sklearn.linear model import LogisticRegression

from sklearn.neighbors import KNeighborsClassifier

from sklearn.ensemble import RandomForestClassifier

from sklearn.model selection import train test split, cross val score

from sklearn.model selection import RandomizedSearchCV, GridSearchCV

from sklearn.metrics import confusion matrix, classification report

from sklearn.metrics import precision score, recall score, fl score, RocCurveDisplay

Load Data

df = pd.read_csv("Downloads/healthcare_dataset.csv.zip") df.shape

df.shape

OUTPUT:

(55500, 15)

Data Exploration (exploratory data analysis or EDA)

PROGRAM:

df.head()

OUTPUT:

	Name	Age	Gender	Blood Type	Medical Condition	Date of Admission	Doctor	Hospital	Insurance Provider	Billing Amount	Room Number	Admission Type	Discharge Date	Medication	Test Results
0	Bobby JacksOn	30	Male	B-	Cancer	2024-01- 31	Matthew Smith	Sons and Miller	Blue Cross	18856.281306	328	Urgent	2024-02- 02	Paracetamol	Normal
1	LesLie TErRy	62	Male	A+	Obesity	2019-08- 20	Samantha Davies	Kim Inc	Medicare	33643.327287	265	Emergency	2019-08- 26	Ibuprofen	Inconclusive
2	DaNnY sMitH	76	Female	Α-	Obesity	2022-09- 22	Tiffany Mitchell	Cook PLC	Aetna	27955.096079	205	Emergency	2022-10- 07	Aspirin	Normal
3	andrEw waTtS	28	Female	0+	Diabetes	2020-11- 18	Kevin Wells	Hernandez Rogers and Vang,	Medicare	37909.782410	450	Elective	2020-12- 18	Ibuprofen	Abnormal
4	adrIENNE bEll	43	Female	AB+	Cancer	2022-09-	Kathleen Hanna	White- White	Aetna	14238.317814	458	Urgent	2022-10-	Penicillin	Abnormal

PROGRAM:

df.tail()

OUTPUT:

	Name	Age	Gender	Blood Type	Medical Condition	Date of Admission	Doctor	Hospital	Insurance Provider	Billing Amount	Room Number	Admission Type	Discharge Date	Medication	R
55495	eLIZABeTH jaCkSOn	42	Female	0+	Asthma	2020-08- 16	Joshua Jarvis	Jones- Thompson	Blue Cross	2650.714952	417	Elective	2020-09- 15	Penicillin	Abn
55496	KYle pEREz	61	Female	AB-	Obesity	2020-01- 23	Taylor Sullivan	Tucker- Moyer	Cigna	31457.797307	316	Elective	2020-02- 01	Aspirin	N
55497	HEATher WaNG	38	Female	B+	Hypertension	2020-07- 13	Joe Jacobs DVM	and Mahoney Johnson Vasquez,	UnitedHealthcare	27620.764717	347	Urgent	2020-08- 10	Ibuprofen	Abn
55498	JENniFER JOneS	43	Male	0-	Arthritis	2019-05- 25	Kimberly Curry	Jackson Todd and Castro,	Medicare	32451.092358	321	Elective	2019-05- 31	Ibuprofen	Abn
55499	jAMES GARCIA	53	Female	0+	Arthritis	2024-04- 02	Dennis Warren	Henry Sons and	Aetna	4010.134172	448	Urgent	2024-04- 29	Ibuprofen	Abn

PROGRAM:

df["Gender"].value_counts()

```
Gender
```

Male 27774 Female 27726

Name: count, dtype: int64

PROGRAM:

df["Blood Type"].value counts()

OUTPUT:

Blood Type

A- 6969

A+ 6956

AB+ 6947

AB- 6945

B+ 6945

B- 6944

O+ 6917

O- 6877

Name: count, dtype: int64

PROGRAM:

df["Test Results"].value_counts()

OUTPUT:

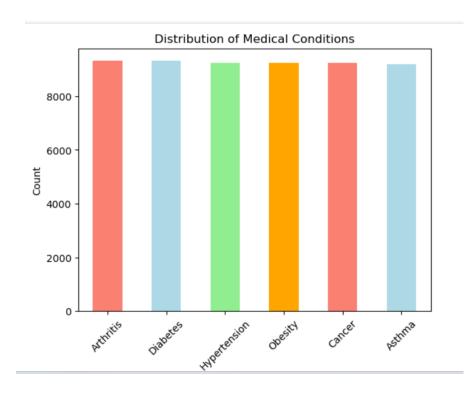
Test Results

Abnormal 18627 Normal 18517 Inconclusive 18356 Name: count, dtype: int64

```
df["Medical Condition"].value_counts().plot(kind='bar', color=["salmon", "lightblue", "lightgreen", "orange"])
plt.title("Distribution of Medical Conditions")
plt.xlabel("Condition")
plt.ylabel("Count")
plt.xticks(rotation=45)
```

plt.show()

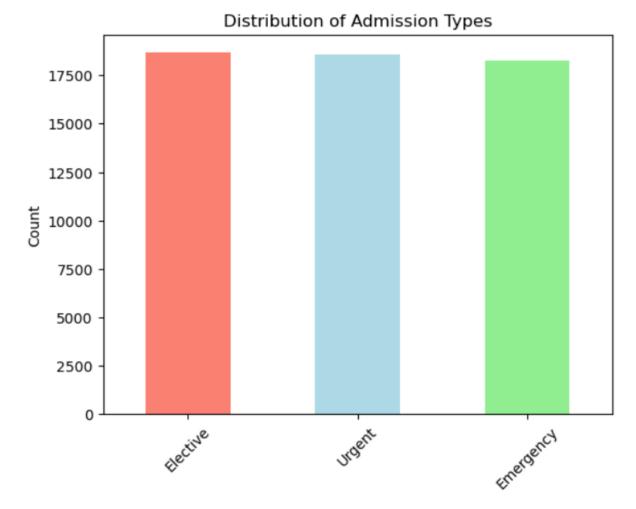
OUTPUT:



PROGRAM:

df.info()

```
<class 'pandas.core.frame.DataFrame'>
 RangeIndex: 55500 entries, 0 to 55499
Data columns (total 15 columns):
      Column
                           Non-Null Count Dtype
      ____
                           -----
  0
      Name
                           55500 non-null object
  1
                           55500 non-null int64
      Age
  2
      Gender
                           55500 non-null object
      Blood Type
                           55500 non-null object
  3
      Medical Condition
  4
                           55500 non-null object
      Date of Admission
  5
                           55500 non-null object
                           55500 non-null object
  6
      Doctor
      Hospital
                           55500 non-null object
  7
                           55500 non-null object
  8
      Insurance Provider
      Billing Amount
                           55500 non-null float64
  10 Room Number
                           55500 non-null int64
  11 Admission Type
                           55500 non-null object
  12 Discharge Date
                           55500 non-null object
  13 Medication
                           55500 non-null object
  14 Test Results
                           55500 non-null object
 dtypes: float64(1), int64(2), object(12)
memory usage: 6.4+ MB
PROGRAM:
df["Admission Type"].value counts().plot(kind='bar', color=["salmon", "lightblue",
"lightgreen"])
plt.title("Distribution of Admission Types")
plt.xlabel("Type")
plt.ylabel("Count")
plt.xticks(rotation=45)
plt.show()
```



df.isna().sum()

OUTPUT:

Name	0	
Age	0	
Gender	0	
Blood Type	0	
Medical Condition	0	
Date of Admission	0	
Doctor	0	
Hospital	0	
Insurance Provider	0	
Billing Amount	0	
Room Number	0	
Admission Type	0	
Discharge Date	0	
Medication	0	
Test Results	0	
dtype: int64		

df.describe()

OUTPUT:

	Age	Billing Amount	Room Number
count	55500.000000	55500.000000	55500.000000
mean	51.539459	25539.316097	301.134829
std	19.602454	14211.454431	115.243069
min	13.000000	-2008.492140	101.000000
25%	35.000000	13241.224652	202.000000
50%	52.000000	25538.069376	302.000000
75 %	68.000000	37820.508436	401.000000
max	89.000000	52764.276736	500.000000

PROGRAM:

df.Gender.value_counts()

OUTPUT:

Gender

Male 27774 Female 27726

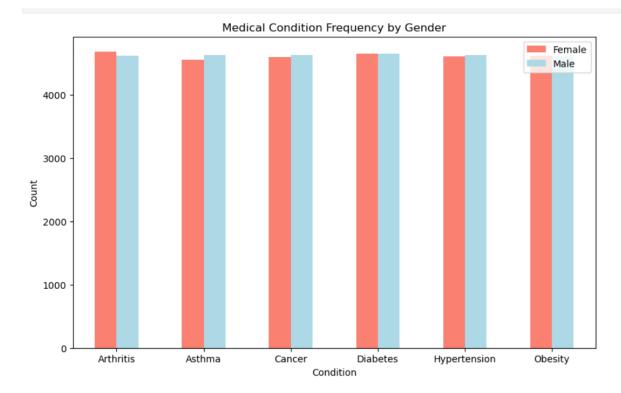
Name: count, dtype: int64

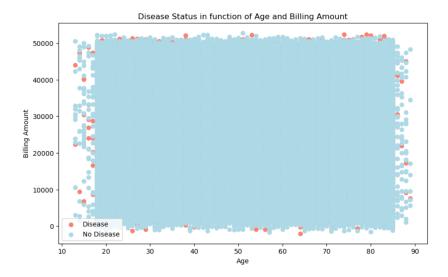
PROGRAM:

pd.crosstab(df["Medical Condition"], df["Gender"])

Gender	Female	Male
Medical Condition		
Arthritis	4686	4622
Asthma	4553	4632
Cancer	4602	4625
Diabetes	4651	4653
Hypertension	4612	4633
Obesity	4622	4609

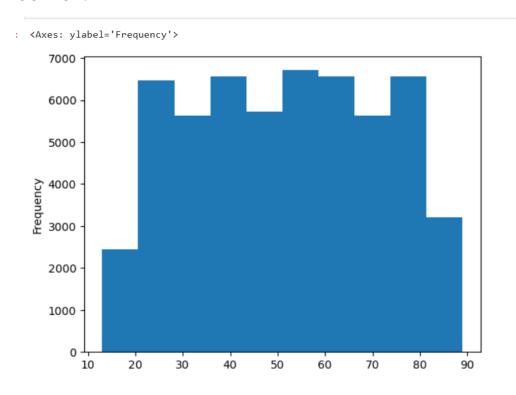
```
pd.crosstab(df["Medical Condition"], df["Gender"]).plot(kind="bar", figsize=(10, 6), color=["salmon", "lightblue"])
plt.title("Medical Condition Frequency by Gender")
plt.xlabel("Condition")
plt.ylabel("Count")
plt.legend(["Female", "Male"])
plt.xticks(rotation=0)
plt.show()
```





df.Age.plot.hist()

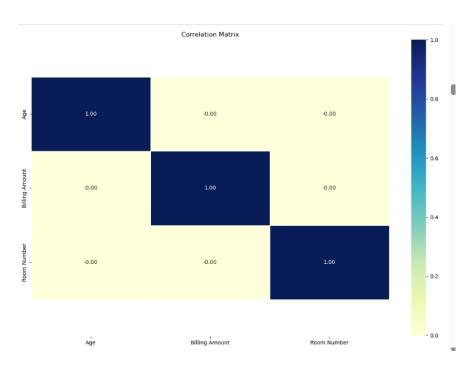
OUTPUT:



correlation matrix

```
numeric_df = df.select_dtypes(include=[np.number])
corr_matrix = numeric_df.corr()
fig, ax = plt.subplots(figsize=(15, 10))
sns.heatmap(corr_matrix, annot=True, linewidths=0.5, fmt=".2f", cmap="YlGnBu", ax=ax)
bottom, top = ax.get_ylim()
ax.set_ylim(bottom + 0.5, top - 0.5)
plt.title("Correlation Matrix")
plt.show()
```

OUTPUT:



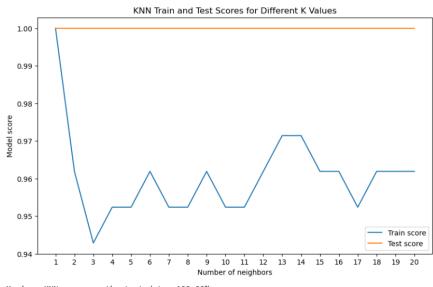
PROGRAM:

import numpy as np

import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split

```
from sklearn.neighbors import KNeighborsClassifier
from sklearn.datasets import load iris
from sklearn.metrics import accuracy score
data = load iris()
X = data.data
y = data.target
X train, X test, y train, y test = train test split(X, y, test size=0.3, random state=42)
train scores = [] test scores = [] neighbors = range(1, 21)
for k in neighbors:
knn = KNeighborsClassifier(n neighbors=k)
knn.fit(X train, y train)
train_accuracy = accuracy_score(y_train, knn.predict(X_train))
test accuracy = accuracy score(y test, knn.predict(X test))
train scores.append(train accuracy)
test scores.append(test accuracy)
plt.figure(figsize=(10,6))
plt.plot(neighbors, train scores, label="Train score")
plt.plot(neighbors, test scores, label="Test score")
plt.xticks(np.arange(1, 21, 1))
plt.xlabel("Number of neighbors")
plt.ylabel("Model score")
plt.title("KNN Train and Test Scores for Different K Values")
plt.legend()
plt.show()
print(f"Maximum KNN score on the test data: {max(test scores)*100:.2f}%")
OUTPUT:
```



Maximum KNN score on the test data: 100 00%

PROGRAM:

import seaborn as sns

import matplotlib.pyplot as plt

```
# Visualize distribution of Age

plt.figure(figsize=(10,6))

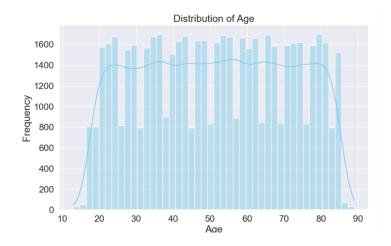
sns.histplot(df['Age'], kde=True, color='skyblue')

plt.title('Distribution of Age')

plt.xlabel('Age')

plt.ylabel('Frequency')

plt.show()
```



plt.figure(figsize=(8,6))

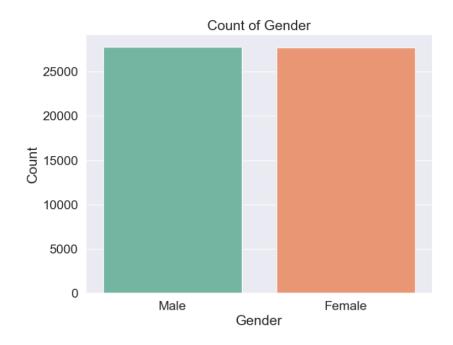
sns.countplot(x='Gender', data=df, palette='Set2')

plt.title('Count of Gender')

plt.xlabel('Gender')

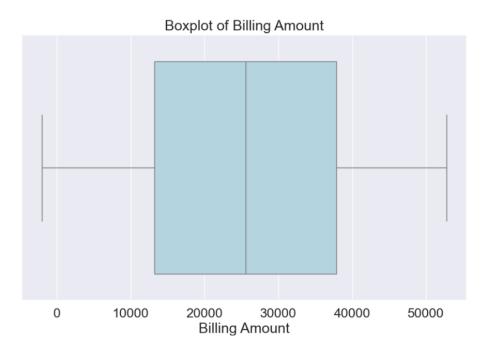
plt.ylabel('Count')

plt.show()



```
plt.figure(figsize=(10,6))
sns.boxplot(x='Billing Amount', data=df, color='lightblue')
plt.title('Boxplot of Billing Amount')
plt.xlabel('Billing Amount')
plt.show()
```

OUTPUT:



Outliers using IQR (Interquartile Range)

```
Q1 = df[['Age', 'Billing Amount']].quantile(0.25) Q3 = df[['Age', 'Billing Amount']].quantile(0.75) IQR = Q3 - Q1 outliers = ((df[['Age', 'Billing Amount']] < (Q1 - 1.5 * IQR)) | (df[['Age', 'Billing Amount']] > (Q3 + 1.5 * IQR)))
```

```
outlier_count = outliers.sum()
print("\nOutliers per column:")
print(outlier_count)
df_cleaned_no_outliers = df[~outliers.any(axis=1)]
OUTPUT:
```

Outliers per column:

Age 0

Billing Amount 0

dtype: int64

duplicate rows

PROGRAM:

```
duplicates = df.duplicated().sum()
print(f"Number of duplicate rows: {duplicates}")
df_cleaned_no_duplicates = df.drop_duplicates()
```

OUTPUT:

Number of duplicate rows: 534