UNIFIED MENTOR

programe: Data Science Advance Course

Task: Exploratory Data Analysis on Heart Disease Data

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As a Health Data Analyst my main obejective for this analysis includes:

- 1. Cleaning and Transformation of Data
- 2. Number of patients diagnosed with heart disease.
- 3. Distribution of diseased patients based on gender.
- 4. Analysis concerning age:
 - 1. Gender-specific impact of age on cholesterol levels.
 - 2. Influence of age on blood pressure.
 - 3. Relationship between heart rate and age among diseased patients.
- 5. Examination of diseased patients with respect to categorical features:
 - 1. 'Resting Electrocardiogram (ECG)'.
 - 2. Presence of angina.
 - 3. Slope of ST segment.
 - 4. Number of major vessels colored by fluoroscopy('ca').
 - 5. Type of thalassemia.
 - 6. Chest pain type.
 - 7. Fasting blood sugar levels.

```
#Import libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import plotly as px
import missingno as msno
```

#Load The DataSet

df=pd.read_csv('/content/drive/MyDrive/UM_python_assignment/Heart Disease data.csv')
df.head(5)

	age	sex	ср	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	ılı
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	

df.shape

(1025, 14)

This Dataset Consist of 1025 rows and 14 columns

df.info()

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1025 entries, 0 to 1024
Data columns (total 14 columns):
# Column Non-Null Count Dtype
```

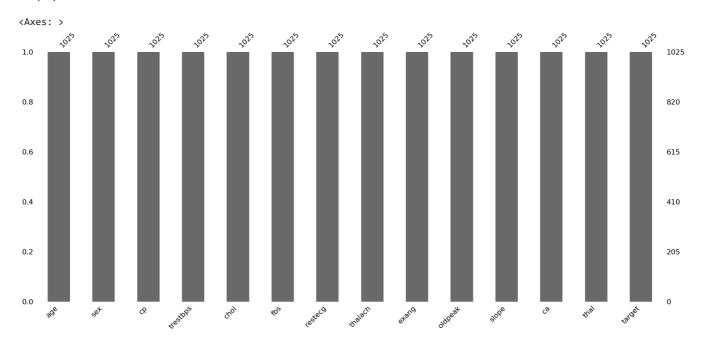
```
0
     age
               1025 non-null
                               int64
1
               1025 non-null
     sex
2
               1025 non-null
                               int64
     ср
3
     trestbps
               1025 non-null
                               int64
               1025 non-null
     chol
                               int64
5
               1025 non-null
                               int64
6
     restecg
               1025 non-null
                               int64
     thalach
               1025 non-null
                               int64
     exang
               1025 non-null
9
     oldpeak
               1025 non-null
                               float64
10
     slope
               1025 non-null
                               int64
11 ca
               1025 non-null
12 thal
               1025 non-null
                               int64
13 target
               1025 non-null
dtypes: float64(1), int64(13)
memory usage: 112.2 KB
```

Cleaning and Transformation of Dataset

```
#check for null values if any
df.isnull().any()
```

age	False
sex	False
ср	False
trestbps	False
chol	False
fbs	False
restecg	False
thalach	False
exang	False
oldpeak	False
slope	False
ca	False
thal	False
target	False
<pre>dtype: bool</pre>	

msno.bar(df)



#check for duplicate rows if any
df.duplicated().sum()

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#droping duplicates
df.drop_duplicates(inplace=True)

```
df.shape (302, 14)
```

After dropping duplicates now the length of dataset is 302 rows and 14 columns

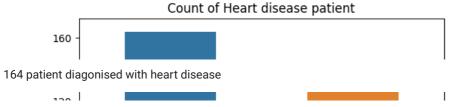
```
df.dtypes
                                             int64
            age
                                             int64
            sex
                                             int64
            trestbps
                                            int64
            chol
                                             int64
            fbs
                                             int64
            restecg
                                             int64
            thalach
                                             int64
                                             int64
            exang
                                       float64
            oldpeak
                                            int64
            slope
                                             int64
            ca
            thal
                                             int64
                                             int64
            target
            dtype: object
#Changing the datatype of columns according to entries
convert_dict={ 'sex': object,
                                  'cp': object,
                                 'fbs': object,
'restecg': object,
'exang' : object,
'slope' : object,
'ca' : object,
'thal' : object,
'target' : object
}
df = df.astype(convert_dict)
print(df.dtypes)
            age
                                             int64
                                           object
            sex
            ср
                                           object
            trestbps
                                            int64
            chol
                                            int64
            fbs
                                           object
            restecg
                                         object
            thalach
                                            int64
                                           object
            exang
            oldpeak
                                       float64
            slope
                                           object
                                           object
            ca
            thal
                                           object
            target
                                           object
            dtype: object
#Renaming the features
df=df.rename({'age':'Age','sex':'Sex','cp':'Chest_pain','trestbps':'Resting_BP','chol':'Cholestrol','fbs':'Fasting_bloodSuga
                          'restecg':'Resting_ECG','thalach':'Max_heartRate','exang':'angina','oldpeak':'ST_Depression','slope':'Slope_ST_Se
                          'ca':'ca','thal':'Thalassemia','target':'Target'},axis=1)
df.columns
           dtype='object')
#Replace the values of columns for better understanding
df['Sex']=df['Sex'].replace([0,1],['Female','Male'])
\label{lem:df-condition} $$ df['Resting_ECG'].replace([0,1,2],['Norm','abnormal','hyper']) $$ $$ df['Resting_ECG'].$$ $$ df[
\tt df['Fasting\_bloodSugar'] = df['Fasting\_bloodSugar'].replace([1,0],['Yes','No'])
```

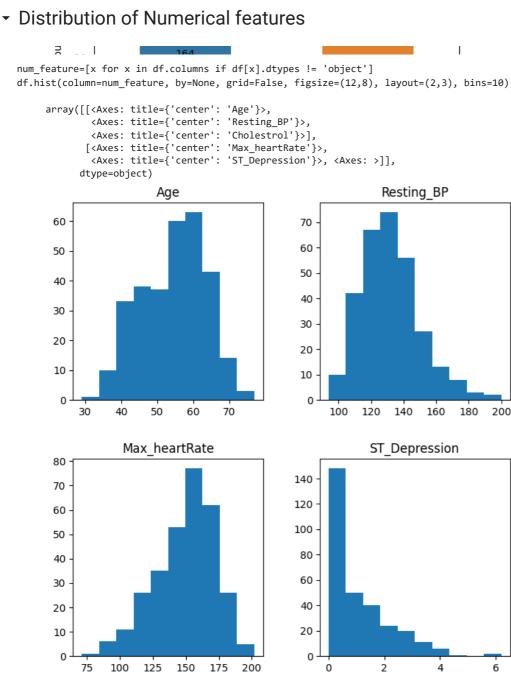
```
df['angina'] = df['angina'].replace([1,0],['Yes','No'])
df['Slope_ST_Segment']=df['Slope_ST_Segment'].replace([0,1,2],['Up', 'flat', 'down'])
df['Target'] = df['Target'].replace([1,0],['Yes','No'])
#unique entries in dataset
df.nunique()
                          41
    Age
    Sex
                          2
     Chest_pain
                          4
                          49
    Resting_BP
                         152
    Cholestrol
    Fasting_bloodSugar
                       2
    Resting_ECG
                          3
    Max_heartRate
                        91
    angina
    ST_Depression
                          40
                         3
    Slope_ST_Segment
    Thalassemia
    Target
    dtype: int64
```

df.describe()

	Age	Resting_BP	Cholestrol	Max_heartRate	ST_Depression	
count	302.00000	302.000000	302.000000	302.000000	302.000000	ılı
mean	54.42053	131.602649	246.500000	149.569536	1.043046	
std	9.04797	17.563394	51.753489	22.903527	1.161452	
min	29.00000	94.000000	126.000000	71.000000	0.000000	
25%	48.00000	120.000000	211.000000	133.250000	0.000000	
50%	55.50000	130.000000	240.500000	152.500000	0.800000	
75%	61.00000	140.000000	274.750000	166.000000	1.600000	
max	77.00000	200.000000	564.000000	202.000000	6.200000	

Number of patients diagnosed with heart disease.

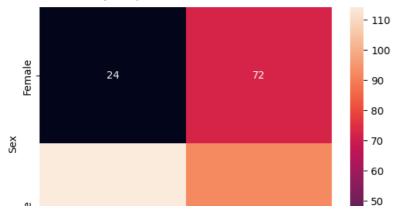




Distribution of diseased patients based on gender.

```
sns.heatmap(pd.crosstab(df['Sex'],df['Target']),annot=True)
```

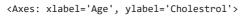


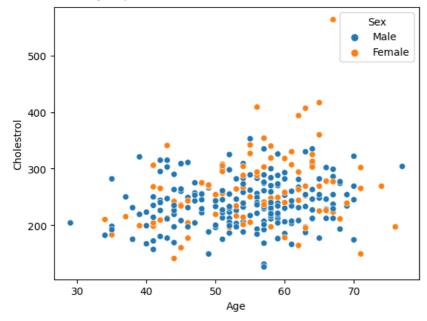


→ Analysis concerning age:

#Gender-specific impact of age on cholesterol levels

sns.scatterplot(data=df,x='Age',y='Cholestrol',hue='Sex')

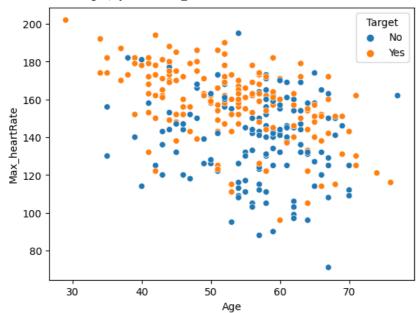




#Influence of age on blood pressure
sns.scatterplot(data=df,x='Age',y='Resting_BP')

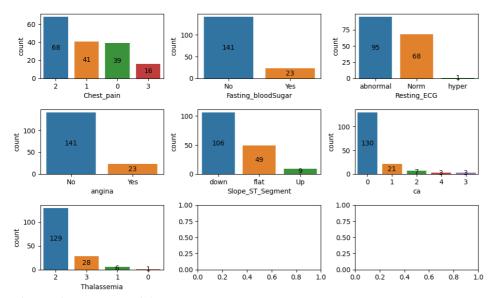
#Relationship between heart rate and age among diseased patients
sns.scatterplot(data=df,x='Age',y='Max_heartRate',hue='Target')

<Axes: xlabel='Age', ylabel='Max_heartRate'>



- Examination of diseased patients with respect to categorical features:

```
#heart-disesed patient dataset
df1=df[df['Target']=='Yes']
plt.rcParams["figure.figsize"] = [10.00, 6.00]
plt.rcParams["figure.autolayout"] = True
fig, ax =plt.subplots(nrows=3,ncols=3)
plt.figure(figsize=(12,8))
g=sns.countplot(data=df1,x='Chest_pain',order=df1['Chest_pain'].value_counts(ascending=False).index,ax=ax[0,0])
g.bar_label(container=g.containers[0],labels=df1['Chest_pain'].value_counts(ascending=False).values,label_type='center')
g=sns.countplot(data=df1,x='Fasting_bloodSugar',order=df1['Fasting_bloodSugar'].value_counts(ascending=False).index,ax=ax[0,
g.bar_label(container=g.containers[0],labels=df1['Fasting_bloodSugar'].value_counts(ascending=False).values,label_type='cent
g=sns.countplot(data=df1,x='Resting_ECG',order=df1['Resting_ECG'].value_counts(ascending=False).index,ax=ax[0,2])
g.bar_label(container=g.containers[0],labels=df1['Resting_ECG'].value_counts(ascending=False).values,label_type='center')
{\tt g=sns.countplot(data=df1,x='angina',order=df1['angina'].value\_counts(ascending=False).index,ax=ax[1,0])}
{\tt g.bar\_label(container=g.containers[0],labels=df1['angina'].value\_counts(ascending=False).values,label\_type='center')}
g=sns.countplot(data=df1,x='Slope\_ST\_Segment',order=df1['Slope\_ST\_Segment'].value\_counts(ascending=False).index,ax=ax[1,1])
{\tt g.bar\_label(container=g.containers[0],labels=df1['Slope\_ST\_Segment'].value\_counts(ascending=False).values,label\_type='center']}.
g=sns.countplot(data=df1,x='ca',order=df1['ca'].value_counts(ascending=False).index,ax=ax[1,2])
g. bar\_label (container=g. containers [0], labels=df1['ca']. value\_counts (ascending=False). values, label\_type='center')
g=sns.countplot(data=df1,x='Thalassemia',order=df1['Thalassemia'].value_counts(ascending=False).index,ax=ax[2,0])
g.bar_label(container=g.containers[0],labels=df1['Thalassemia'].value_counts(ascending=False).values,label_type='center')
fig.show()
```



<Figure size 1200x800 with 0 Axes>