

# Heart-Disease-Analysis

September 16, 2023

```
[24]: import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
```

```
[25]: data=pd.read_csv('heart/heart.csv')
```

```
[26]: # print first 5 rows
data.head(5)
```

```
[26]:
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	\
0	52	1	0	125	212	0	1	168	0	1.0	2	
1	53	1	0	140	203	1	0	155	1	3.1	0	
2	70	1	0	145	174	0	1	125	1	2.6	0	
3	61	1	0	148	203	0	1	161	0	0.0	2	
4	62	0	0	138	294	1	1	106	0	1.9	1	

	ca	thal	target
0	2	3	0
1	0	3	0
2	0	3	0
3	1	3	0
4	3	2	0

```
[27]: # print the last 5 rows
data.tail(5)
```

```
[27]:
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	\
1020	59	1	1	140	221	0	1	164	1	0.0	
1021	60	1	0	125	258	0	0	141	1	2.8	
1022	47	1	0	110	275	0	0	118	1	1.0	
1023	50	0	0	110	254	0	0	159	0	0.0	
1024	54	1	0	120	188	0	1	113	0	1.4	

	slope	ca	thal	target
1020	2	0	2	1

1021	1	1	3	0
1022	1	1	2	0
1023	2	0	2	1
1024	1	1	3	0

```
[28]: data.shape
```

```
[28]: (1025, 14)
```

```
[29]: print("number of rows: ",data.shape[0])
      print("number of coulums: ", data.shape[1])
```

```
number of rows: 1025
number of coulums: 14
```

```
[30]: # checking information related to data
      data.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
```

```
[31]: # checking for the null values
      data.isnull()
```

```
[31]:      age    sex    cp  trestbps    chol    fbs  restecg  thalach  exang  \
0     False  False  False      False  False  False    False    False  False
1     False  False  False      False  False  False    False    False  False
2     False  False  False      False  False  False    False    False  False
```

3	False	False	False	False	False	False	False	False	False
4	False	False	False	False	False	False	False	False	False
...	...	...	...	...	...	...	...	...	...
1020	False	False	False	False	False	False	False	False	False
1021	False	False	False	False	False	False	False	False	False
1022	False	False	False	False	False	False	False	False	False
1023	False	False	False	False	False	False	False	False	False
1024	False	False	False	False	False	False	False	False	False

	oldpeak	slope	ca	thal	target
0	False	False	False	False	False
1	False	False	False	False	False
2	False	False	False	False	False
3	False	False	False	False	False
4	False	False	False	False	False
...	...	...	...	...	...
1020	False	False	False	False	False
1021	False	False	False	False	False
1022	False	False	False	False	False
1023	False	False	False	False	False
1024	False	False	False	False	False

[1025 rows x 14 columns]

```
[32]: #converting null values to 0
data.isnull().sum()
```

```
[32]: 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
```

```
[33]: # checking duplicate data and dropping it
data_dup=data.duplicated().any()
print(data_dup)
```

True

```
[34]: # dropping
data=data.drop_duplicates()
```

```
[35]: # now just checking how much data reduced
data.shape
```

```
[35]: (302, 14)
```

```
[36]: # getting overall statistics for datasets
data.describe()
```

```
[36]:
```

	age	sex	cp	trestbps	chol	fbs	\
count	302.00000	302.000000	302.000000	302.000000	302.000000	302.000000	
mean	54.42053	0.682119	0.963576	131.602649	246.500000	0.149007	
std	9.04797	0.466426	1.032044	17.563394	51.753489	0.356686	
min	29.00000	0.000000	0.000000	94.000000	126.000000	0.000000	
25%	48.00000	0.000000	0.000000	120.000000	211.000000	0.000000	
50%	55.50000	1.000000	1.000000	130.000000	240.500000	0.000000	
75%	61.00000	1.000000	2.000000	140.000000	274.750000	0.000000	
max	77.00000	1.000000	3.000000	200.000000	564.000000	1.000000	

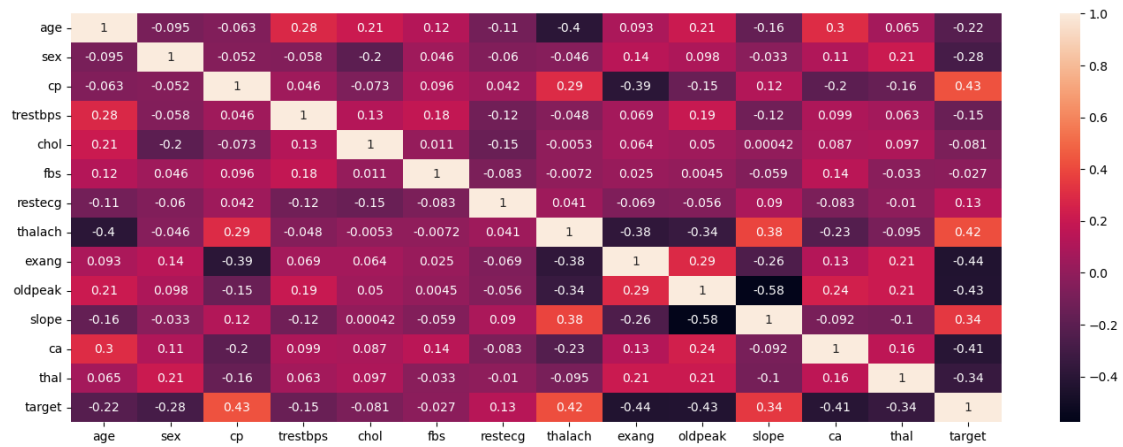
	restecg	thalach	exang	oldpeak	slope	ca	\
count	302.000000	302.000000	302.000000	302.000000	302.000000	302.000000	
mean	0.526490	149.569536	0.327815	1.043046	1.397351	0.718543	
std	0.526027	22.903527	0.470196	1.161452	0.616274	1.006748	
min	0.000000	71.000000	0.000000	0.000000	0.000000	0.000000	
25%	0.000000	133.250000	0.000000	0.000000	1.000000	0.000000	
50%	1.000000	152.500000	0.000000	0.800000	1.000000	0.000000	
75%	1.000000	166.000000	1.000000	1.600000	2.000000	1.000000	
max	2.000000	202.000000	1.000000	6.200000	2.000000	4.000000	

	thal	target
count	302.000000	302.000000
mean	2.314570	0.543046
std	0.613026	0.498970
min	0.000000	0.000000
25%	2.000000	0.000000
50%	2.000000	1.000000
75%	3.000000	1.000000
max	3.000000	1.000000

```
[37]: # Now to check the relationship between different column forming the
      ↪Correlation Matrix
      # data.corr()
      plt.figure(figsize=(17,6))
```

```
sns.heatmap(data.corr(),annot=True)
plt.show()
```

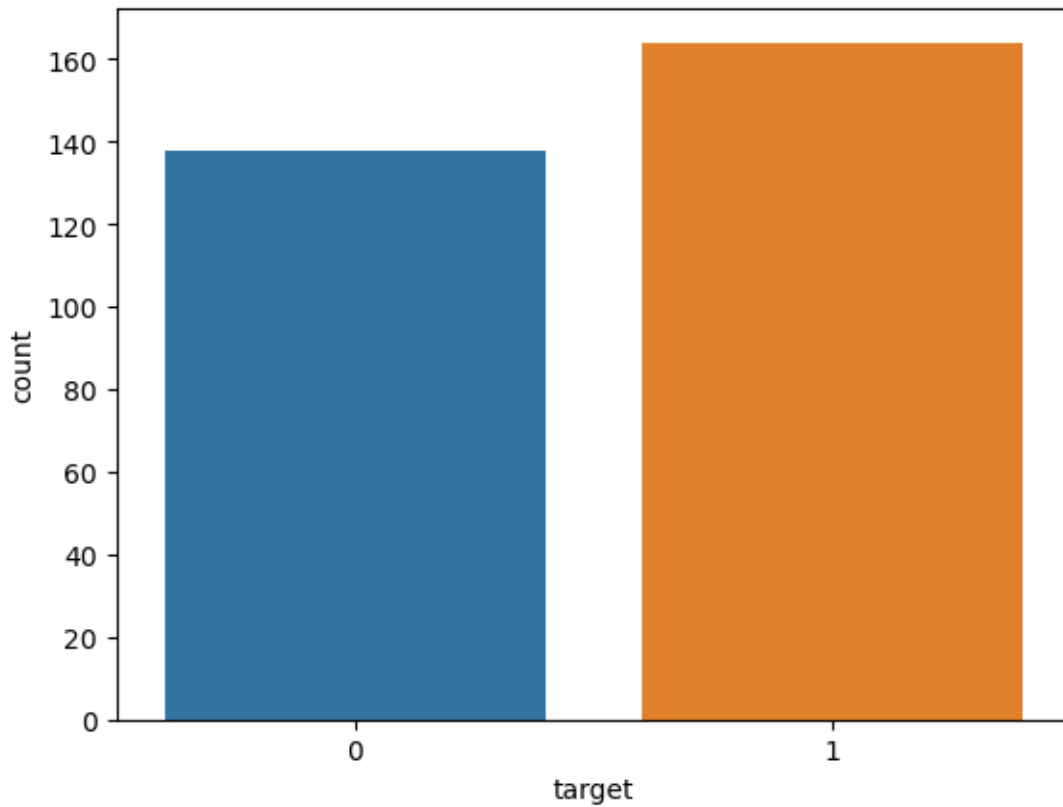


```
[38]: # what correlation matrix do
```

```
[39]: # calculating how many people have heart diseases and how many don't have this
data['target'].value_counts()
```

```
[39]: target
1    164
0    138
Name: count, dtype: int64
```

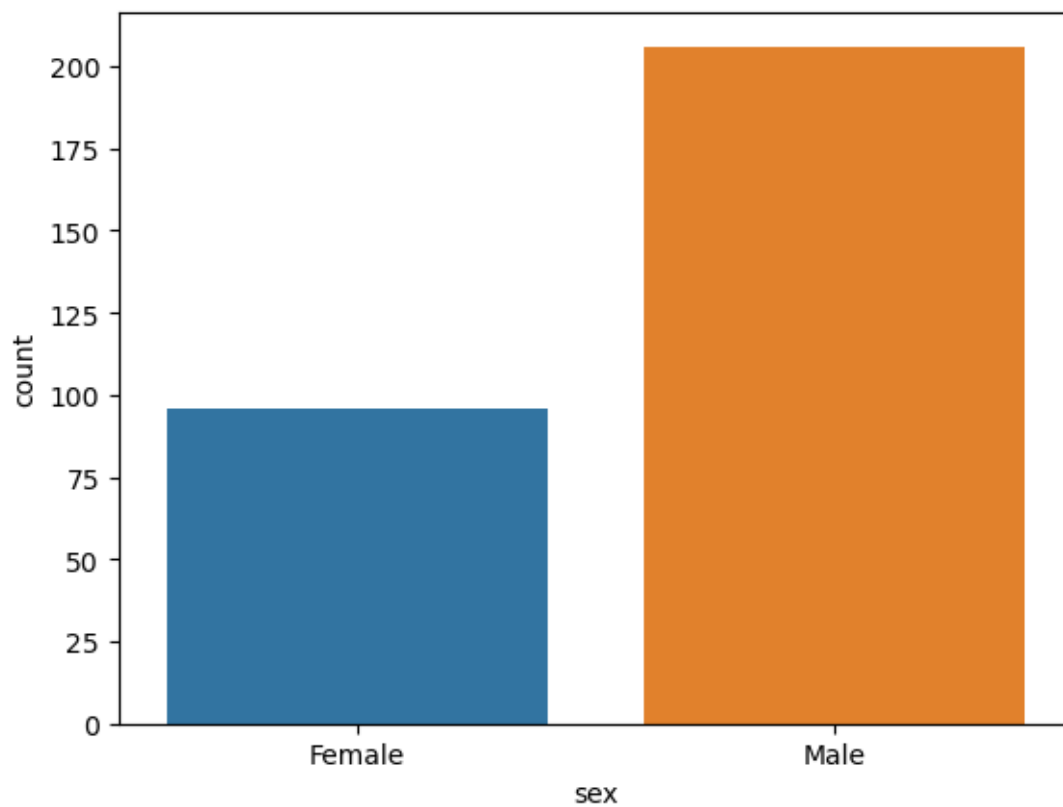
```
[40]: # checking the
sns.countplot(x=data['target'])
plt.show()
# sns.countplot(x="target",data=data)
```



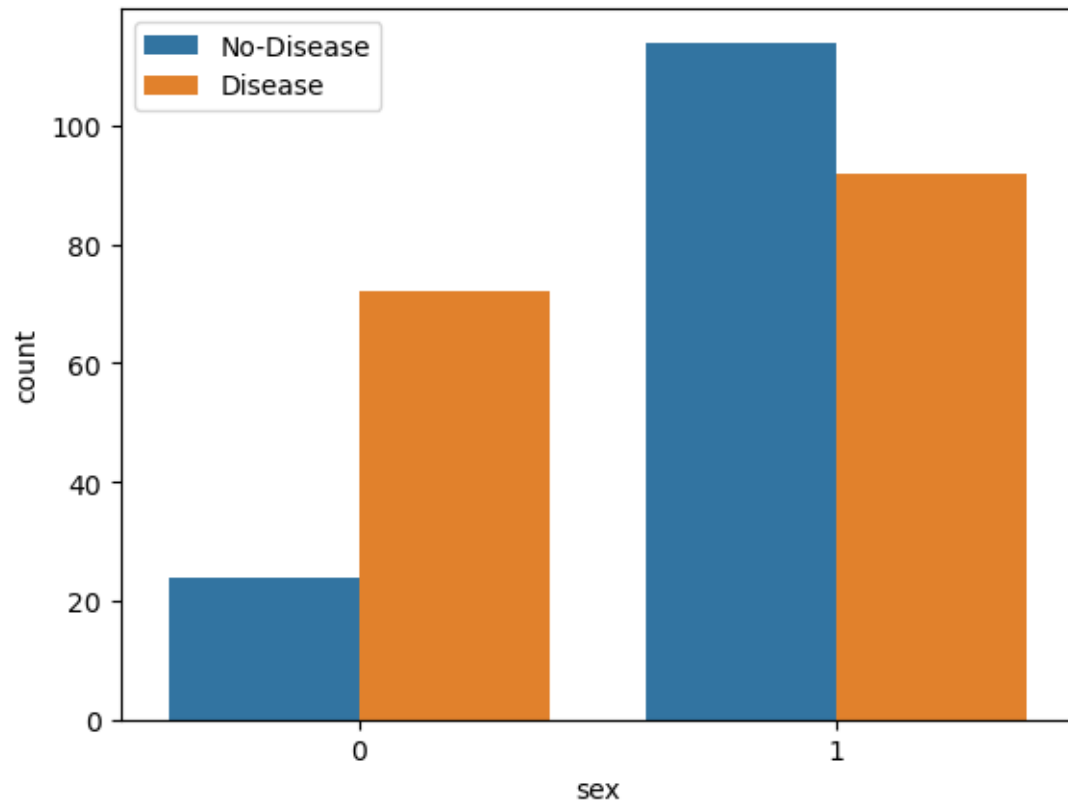
```
[41]: # counting number of male and females  
data['sex'].value_counts()
```

```
[41]: sex  
1    206  
0     96  
Name: count, dtype: int64
```

```
[42]: # showing graphically the males and female  
# sns.countplot(data['sex'])  
sns.countplot(x='sex',data=data)  
plt.xticks([0,1],['Female','Male'])  
plt.show()
```

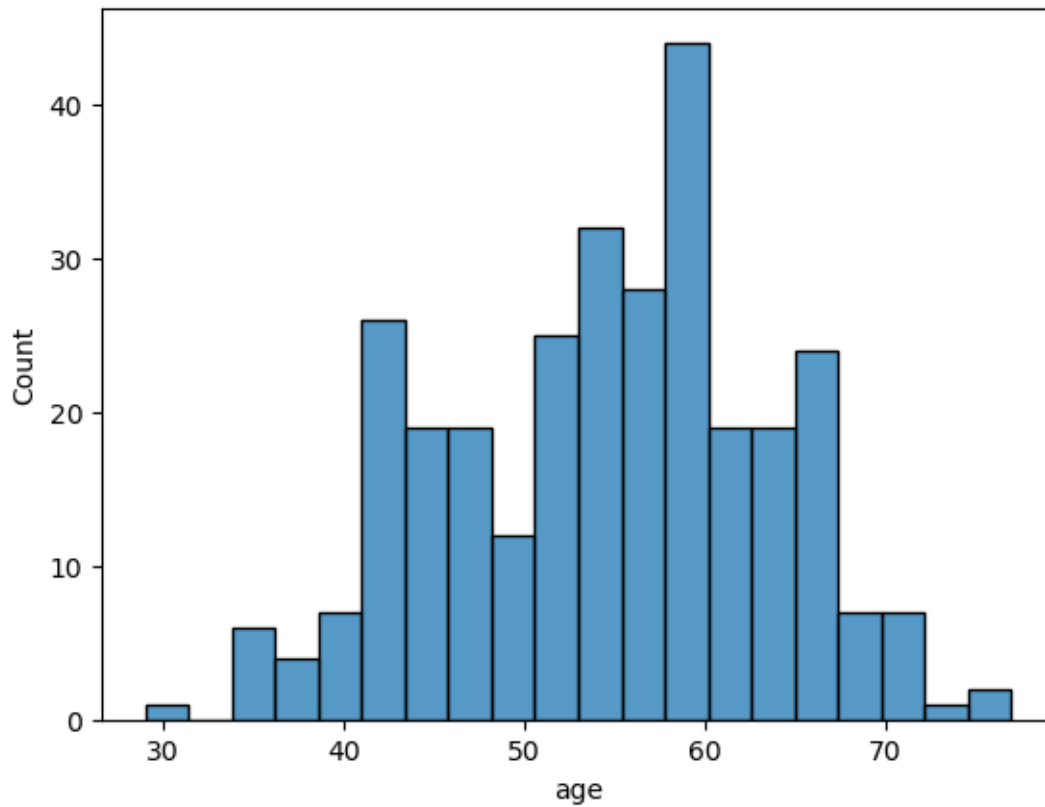


```
[43]: sns.countplot(x='sex',hue="target",data=data)
      # plt.xticks([0,1],['Male','Female'])
      plt.legend(labels=['No-Disease','Disease'])
      plt.show()
```

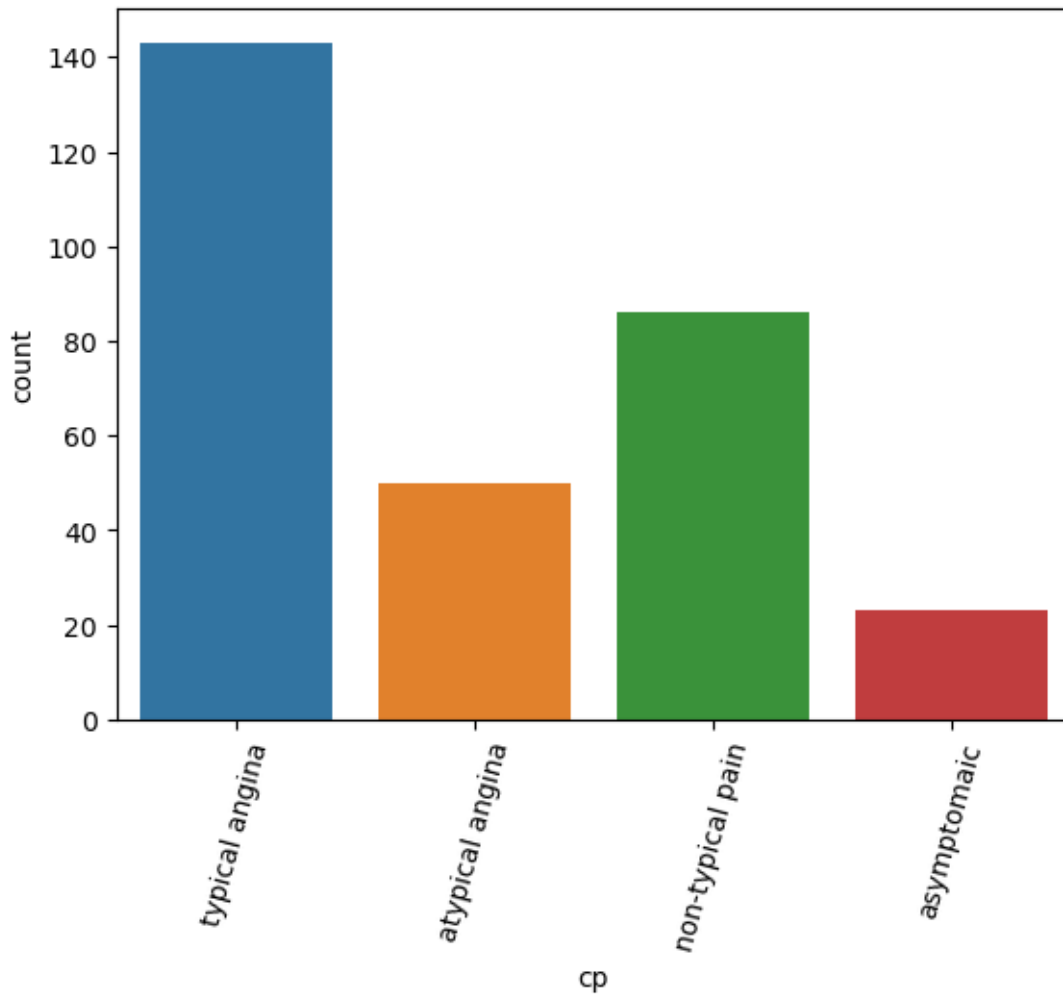


```
[44]: # checking for the people on age basis who were victim of heart disease
sns.histplot(data['age'],bins=20)
plt.show()
```





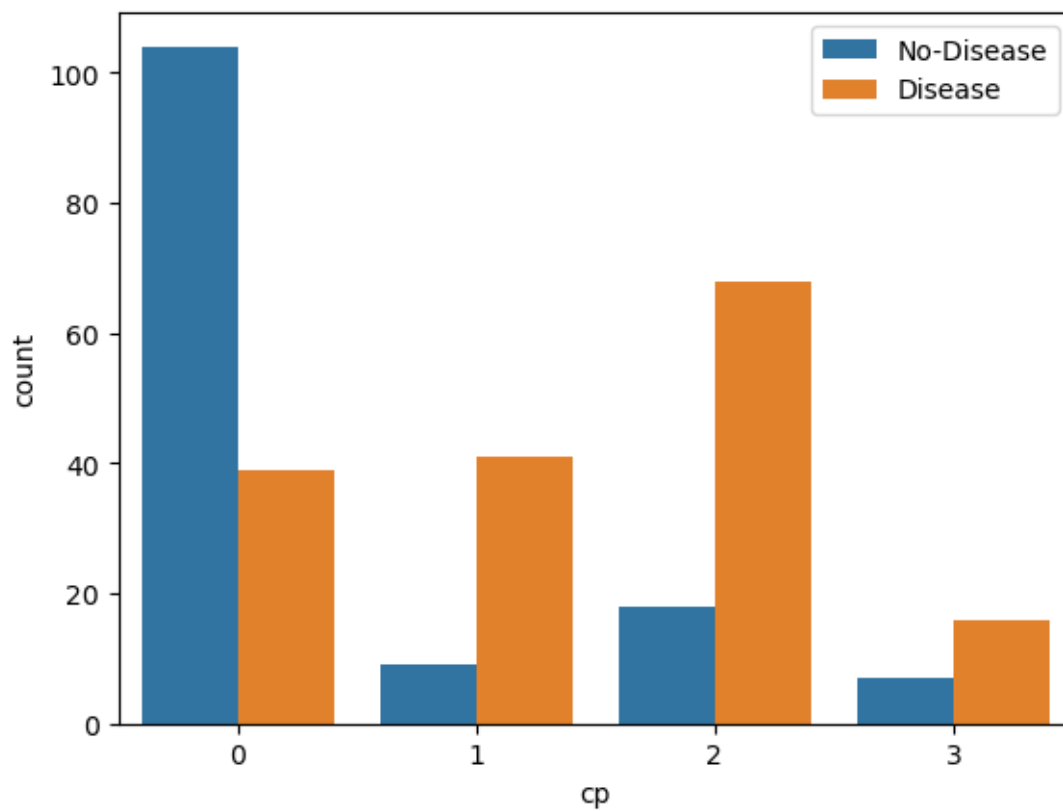
```
[45]: #checking chest pain type
sns.countplot(x=data['cp'])
plt.xticks([0,1,2,3],["typical angina","atypical angina","non-typical_
    ↳pain","asymptomaic"])
plt.xticks(rotation=75)
plt.show()
```



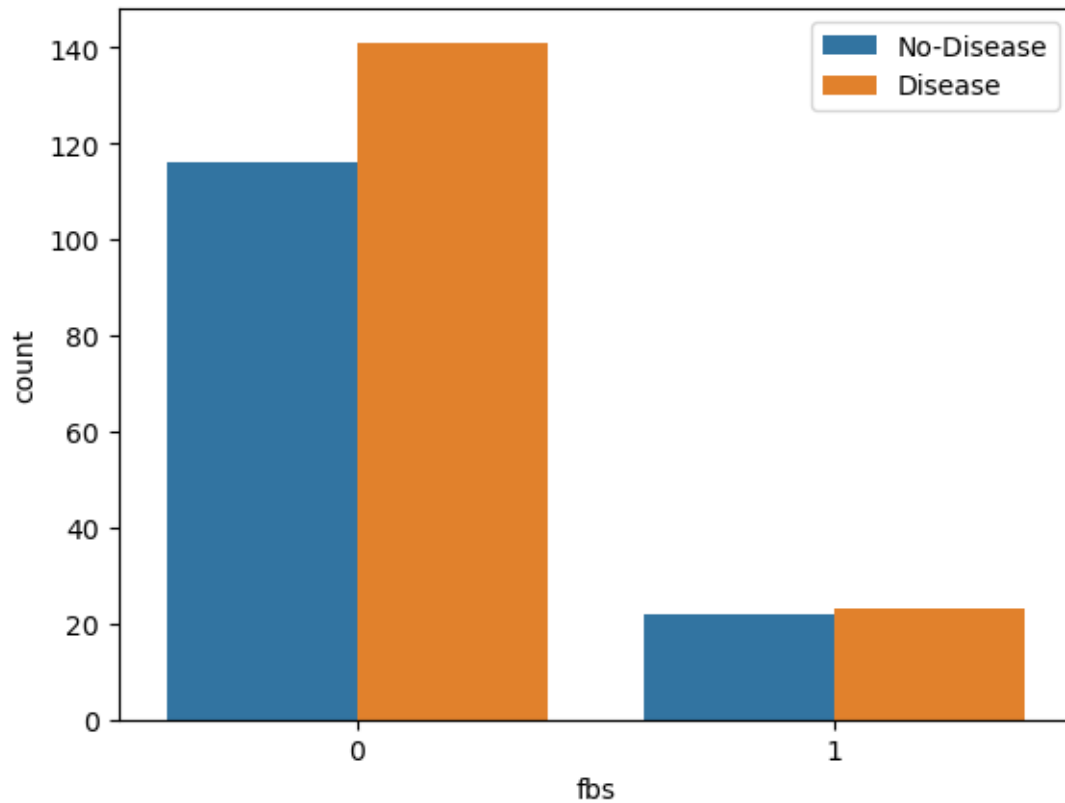
```
[46]: data.columns
```

```
[46]: Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',  
        'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],  
        dtype='object')
```

```
[47]: # identifying who suffer from the chest pain most  
sns.countplot(x="cp",hue="target",data=data)  
plt.legend(labels=["No-Disease","Disease"])  
plt.show()
```



```
[48]: # showing fasting blood sugar distribution according to target variable
sns.countplot(x="fbs",hue="target",data=data)
plt.legend(labels=["No-Disease","Disease"])
plt.show()
```

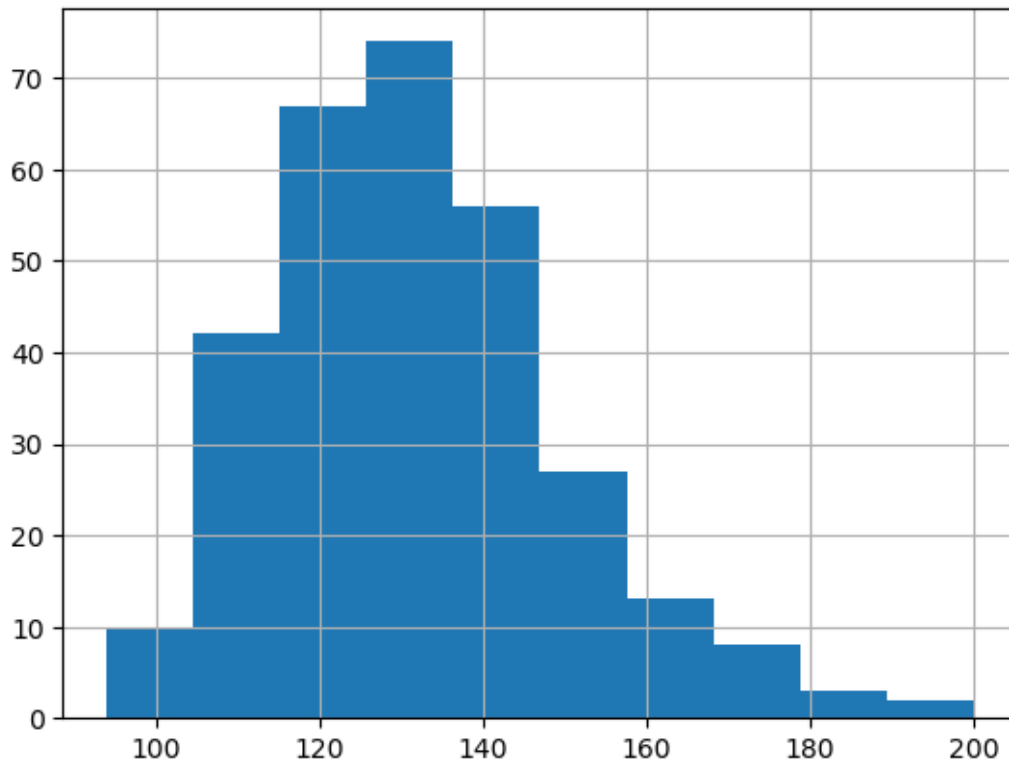


```
[49]: # checking resting blood pressure distribution  
data.columns
```

```
[49]: Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',  
        'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],  
       dtype='object')
```

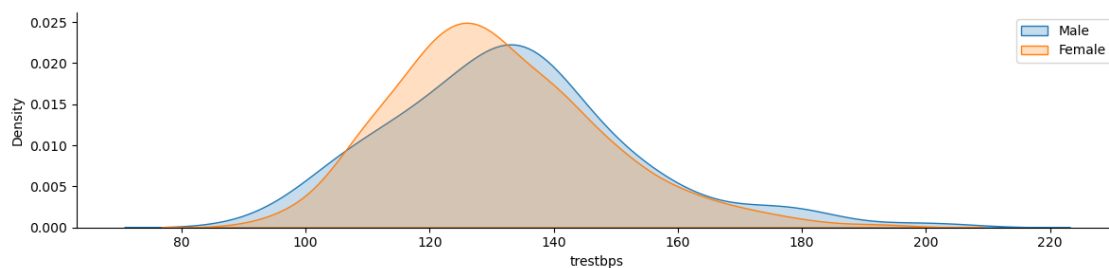
```
[50]: data['trestbps'].hist()
```

```
[50]: <AxesSubplot:>
```



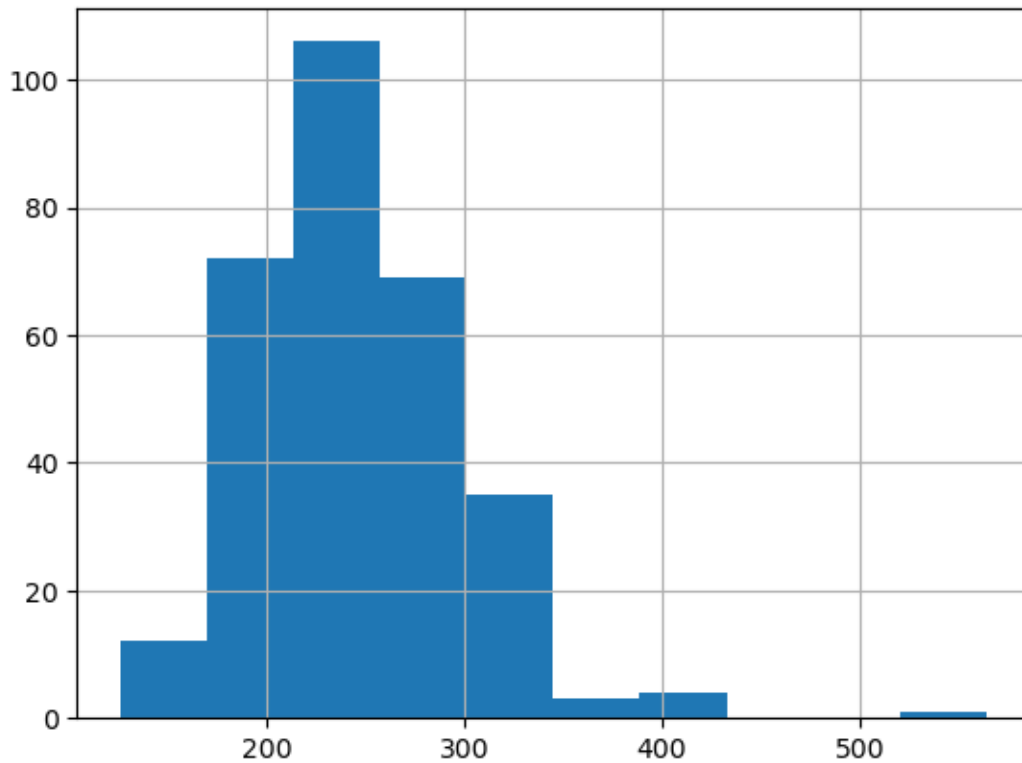
```
[51]: # comparing resting blood pressure as per sex column
g=sns.FacetGrid(data,hue="sex",aspect=4)
g.map(sns.kdeplot,'trestbps',shade=True)
plt.legend(labels=['Male','Female'])
```

```
[51]: <matplotlib.legend.Legend at 0x7f241aff9c00>
```



```
[52]: # showing distribution of serum cholesterol
data['chol'].hist()
```

```
[52]: <AxesSubplot:>
```



```
[53]: # plotting continuous variables
cate_val=[]
cont_val=[]
for column in data.columns:
    if data[column].nunique()<=10:
        cate_val.append(column)
    else:
        cont_val.append(column)
```

```
[54]: cate_val
```

```
[54]: ['sex', 'cp', 'fbs', 'restecg', 'exang', 'slope', 'ca', 'thal', 'target']
```

```
[55]: cont_val
```

```
[55]: ['age', 'trestbps', 'chol', 'thalach', 'oldpeak']
```

```
[56]: data.hist(cont_val,figsize=(15,6))
plt.tight_layout()
plt.show()
```

