HEART DISEASE ANALYSIS

February 20, 2025

IMPORTING LIBRARIES

[1]:
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

Step 2: Loading and Exploring the Dataset

[2]:
```
df = pd.read_csv(r"C:\Users\Josphat\Desktop\DATA ANALYSIS VIDEOS AND␣
↪DATASETS-DATA THINKERS\HEART DISEASE.csv")
```

[3]:
```
df.head()
```

[3]:

	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

[4]:
```
#finding the shape of the dataset
df.shape
print("Number of Rows:", df.shape[0])
print("Number of Columns:", df.shape[1])
```

```
Number of Rows: 1025
Number of Columns: 14
```

[5]:
```
#checking for null values
df.isnull().sum()
```

[5]:
```
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
```

[6]:
```
#finding information about the dataset
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
```

[7]:
```
#finding columns of the dataset
df.columns
```

[7]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[8]:
```
#finding the descriptive statistics of the dataset
df.describe()
```

[8]:

	age	sex	cp	trestbps	chol
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.00000
mean	54.434146	0.695610	0.942439	131.611707	246.00000
std	9.072290	0.460373	1.029641	17.516718	51.59251
min	29.000000	0.000000	0.000000	94.000000	126.00000
25%	48.000000	0.000000	0.000000	120.000000	211.00000
50%	56.000000	1.000000	1.000000	130.000000	240.00000
75%	61.000000	1.000000	2.000000	140.000000	275.00000
max	77.000000	1.000000	3.000000	200.000000	564.00000

	fbs	restecg	thalach	exang	oldpeak
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	0.149268	0.529756	149.114146	0.336585	1.071512
std	0.356527	0.527878	23.005724	0.472772	1.175053
min	0.000000	0.000000	71.000000	0.000000	0.000000
25%	0.000000	0.000000	132.000000	0.000000	0.000000
50%	0.000000	1.000000	152.000000	0.000000	0.800000
75%	0.000000	1.000000	166.000000	1.000000	1.800000
max	1.000000	2.000000	202.000000	1.000000	6.200000

	slope	ca	thal	target
count	1025.000000	1025.000000	1025.000000	1025.000000
mean	1.385366	0.754146	2.323902	0.513171
std	0.617755	1.030798	0.620660	0.500070
min	0.000000	0.000000	0.000000	0.000000
25%	1.000000	0.000000	2.000000	0.000000
50%	1.000000	0.000000	2.000000	1.000000
75%	2.000000	1.000000	3.000000	1.000000
max	2.000000	4.000000	3.000000	1.000000

[9]:
```
#checking for duplicated values and Dropping them
data_dup=df.duplicated().any()
print(data_dup)
```

```
True
```

[10]:
```
df.duplicated().sum()
```

[10]:
```
723
```

[11]:
```
df.shape
```

[11]:
```
(1025, 14)
```

[12]:
```
#dropping the duplicated values
df.drop_duplicates(inplace = True)
```

[13]:
```
df.shape
```

[13]:
```
(302, 14)
```

[14]:
```
df.describe()
```

[14]:

	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

[19]:
```
#Drawing correlation Matrix
plt.figure(figsize=(17,6))
sns.heatmap(df.corr(),annot = True)
```

[19]:
```
<Axes: >
```

Question 1: How many people have Heart disease and How many don't have heart disease in this Dataset

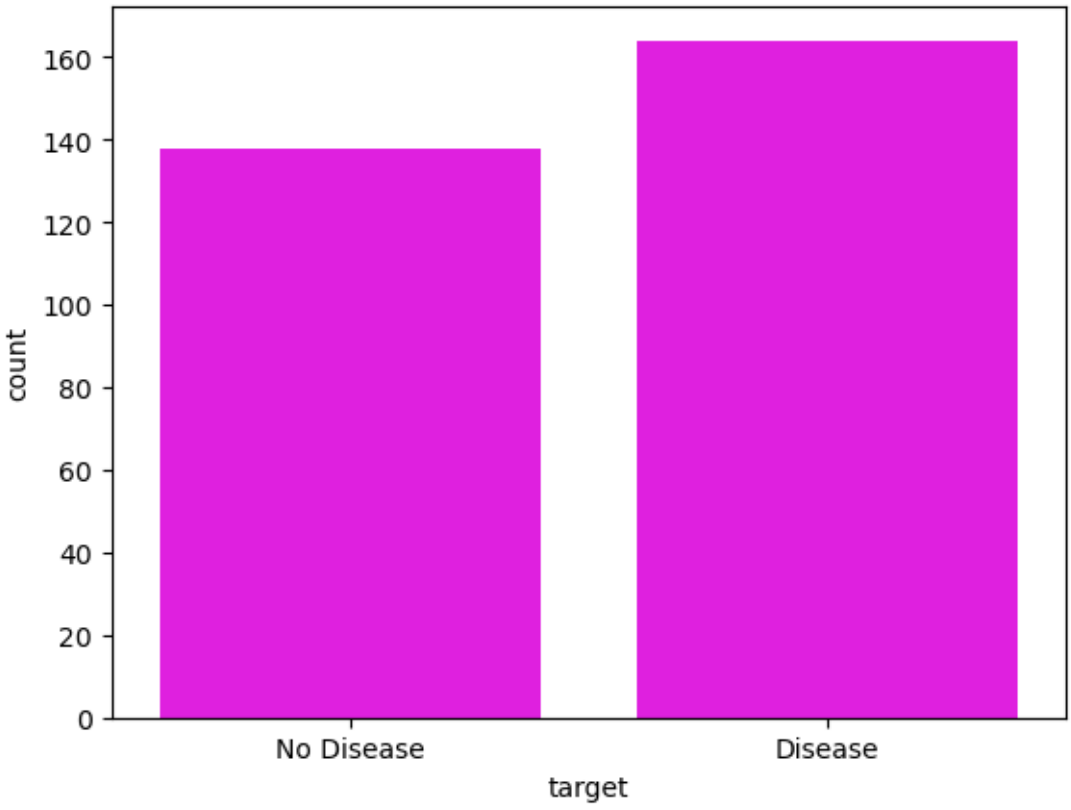
[21]:
```
df.columns
```

[21]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[23]:
```
df["target"].value_counts()
```

[23]:
```
target
1    164
0    138
Name: count, dtype: int64
```

[108]:
```
sns.countplot(x='target', data=df, color='magenta')
plt.xticks([1,0],['Disease','No Disease'])
plt.show()
```



164 people have heart disease and 138 people have no heart disease

Question 2: Find count of Male and Female in the Dataset

[27]:
```
df.columns
```

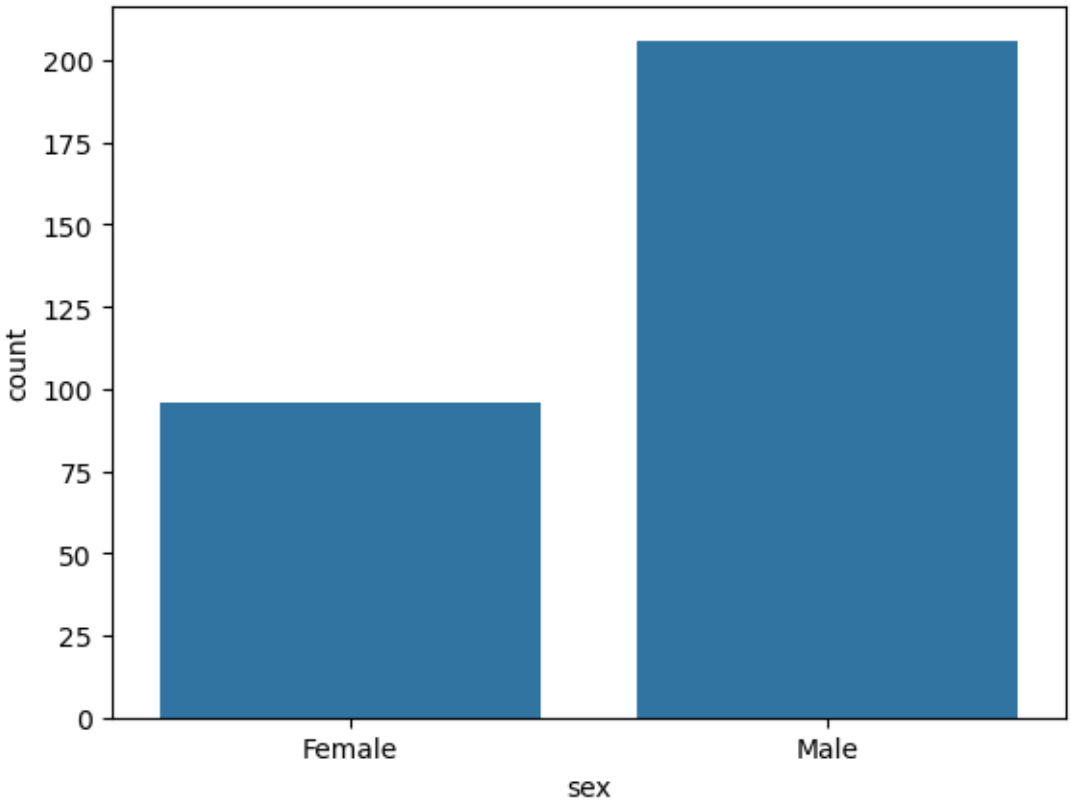
[27]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[29]:
```
df['sex'].value_counts()
```

[29]:
```
sex
1    206
0     96
Name: count, dtype: int64
```

206 people are male and 96 people are Female

[40]:
```
sns.countplot(x='sex', data=df)
plt.xticks([1,0], ['Male','Female'])
plt.show()
```

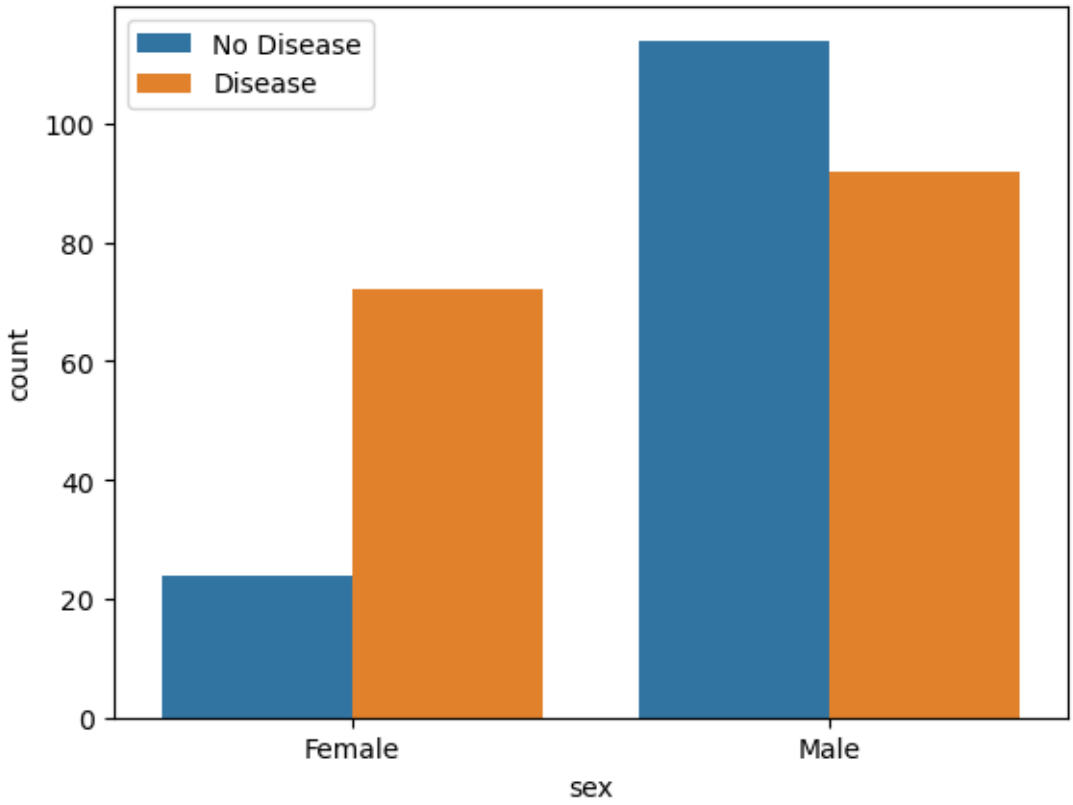


Question 3: Find gender distribution According to the target variable

[32]:
```
df.columns
```

[32]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[35]:
```
sns.countplot(x='sex', hue='target', data=df)
plt.xticks([1,0],['Male','Female'])
plt.legend(labels=['No Disease','Disease'])
plt.show()
```



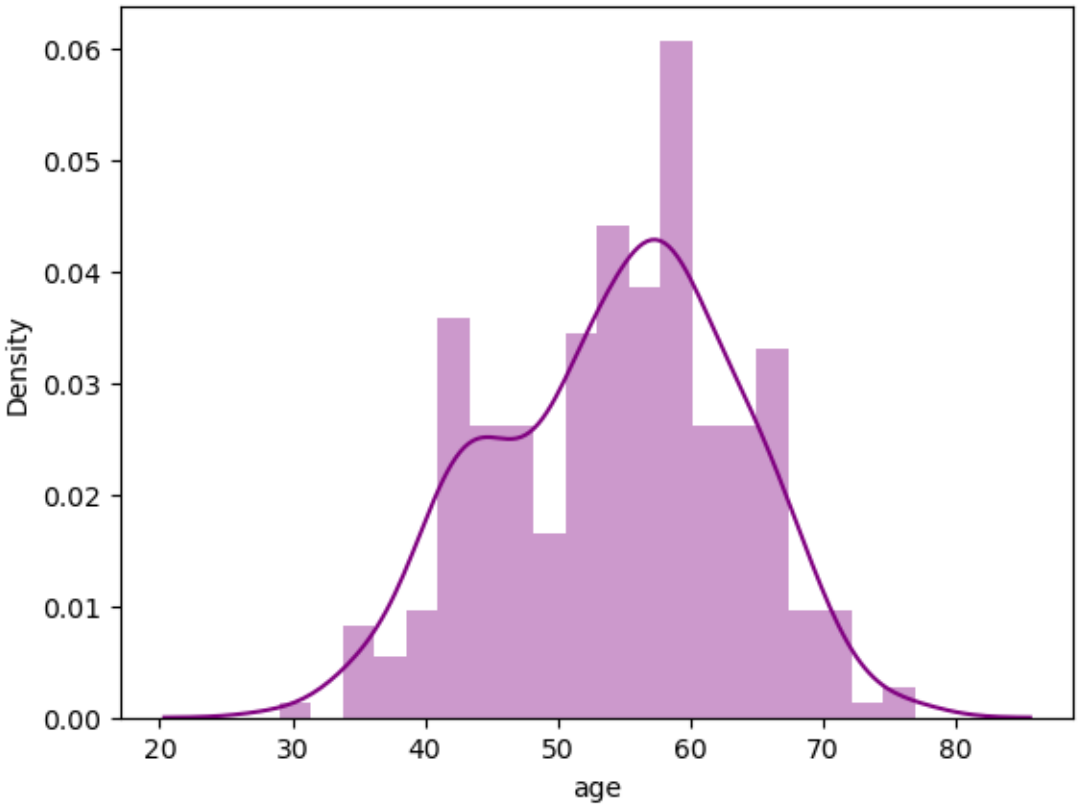
Question 4: Check Age Distribution in the Dataset

[44]:
```
df.columns
```

[44]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[48]:
```
sns.distplot(df.age, bins=20,color='purple')
plt.show()
```

```
C:\Users\Josphat\AppData\Local\Temp\ipykernel_15108\415452091.py:1: UserWarning:

`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either `displot` (a figure-level function with
similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see
https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751

  sns.distplot(df.age, bins=20,color='purple')
```



Question 5: Check Chest pain type

* Value 0: typical angina
* Value 1: atypical angina
* Value 2: non-anginal pain
* Value 3: asymptomatic

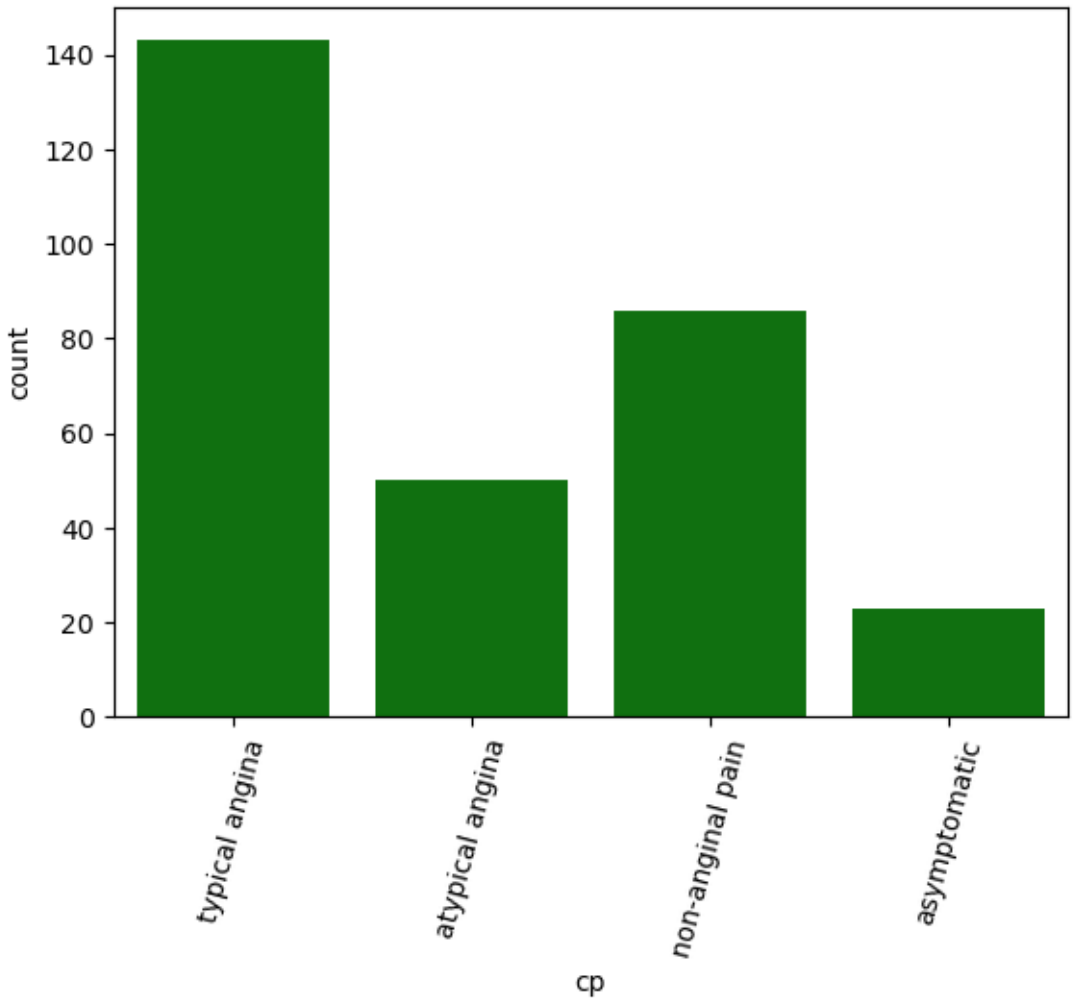
[49]:
```
df.columns
```

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

[54]:
```
df.cp.value_counts()
```

[54]:
```
cp
0    143
2     86
1     50
3     23
Name: count, dtype: int64
```

[56]:
```
sns.countplot(x='cp', data=df, color='green')
plt.xticks([0,1,2,3],['typical angina','atypical angina','non-anginal␣
 ↪pain','asymptomatic'])
plt.xticks(rotation=75)
plt.show()
```



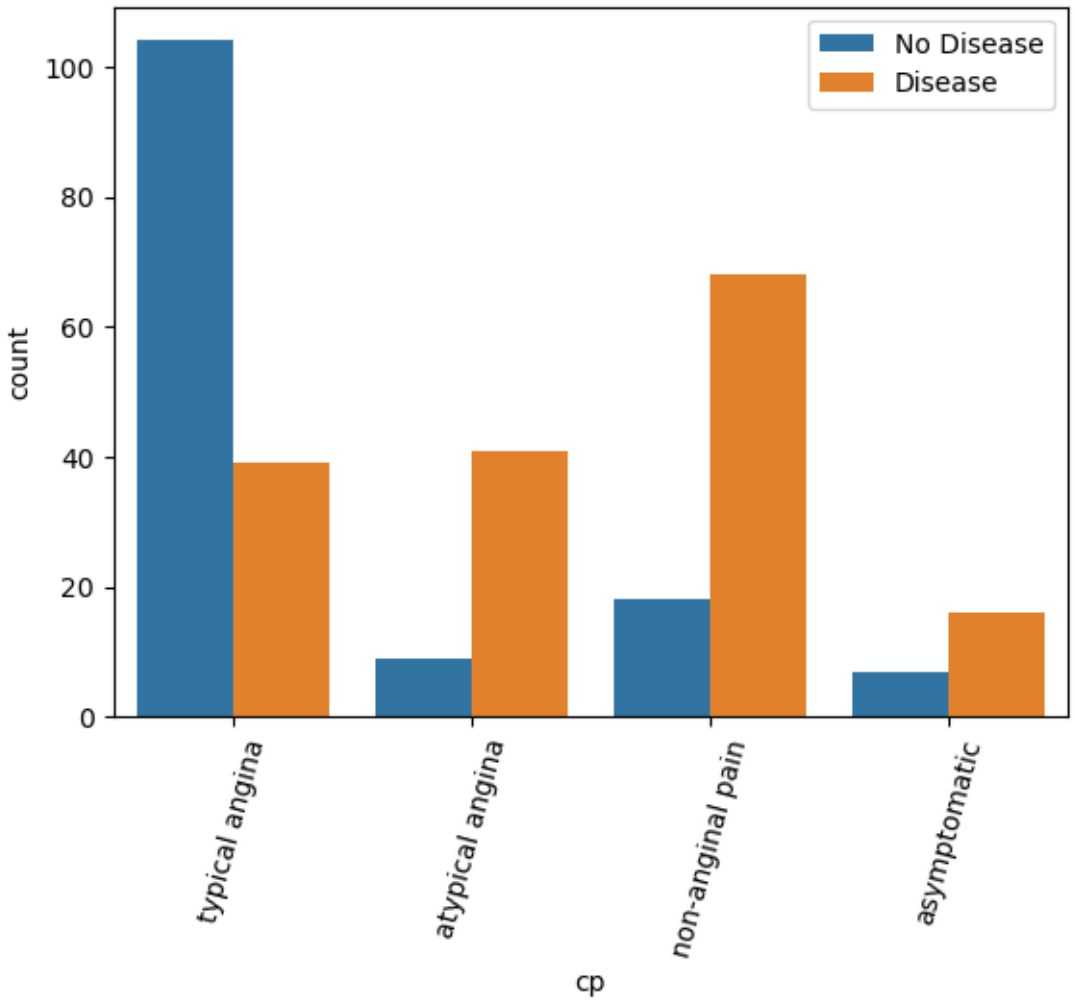
Question 6: Show chest pain Distribution as per Target variabe

[58]:
```
df.columns
```

[58]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[61]:
```
sns.countplot(x='cp', hue='target',data=df)
```

```
plt.xticks([0,1,2,3],['typical angina','atypical angina','non-anginal␣
 ↪pain','asymptomatic'],rotation=75)
plt.legend(labels=['No Disease','Disease'])
plt.show()
```



[68]:
```
df[['cp','target']].value_counts().sort_values(ascending=False)
```

[68]:
```
cp  target
0   0         104
2   1          68
1   1          41
0   1          39
2   0          18
3   1          16
1   0           9
```

```
3   0           7
Name: count, dtype: int64
```

Question 7: Show Fasting Blood Sugar Distribution according to Target variable.

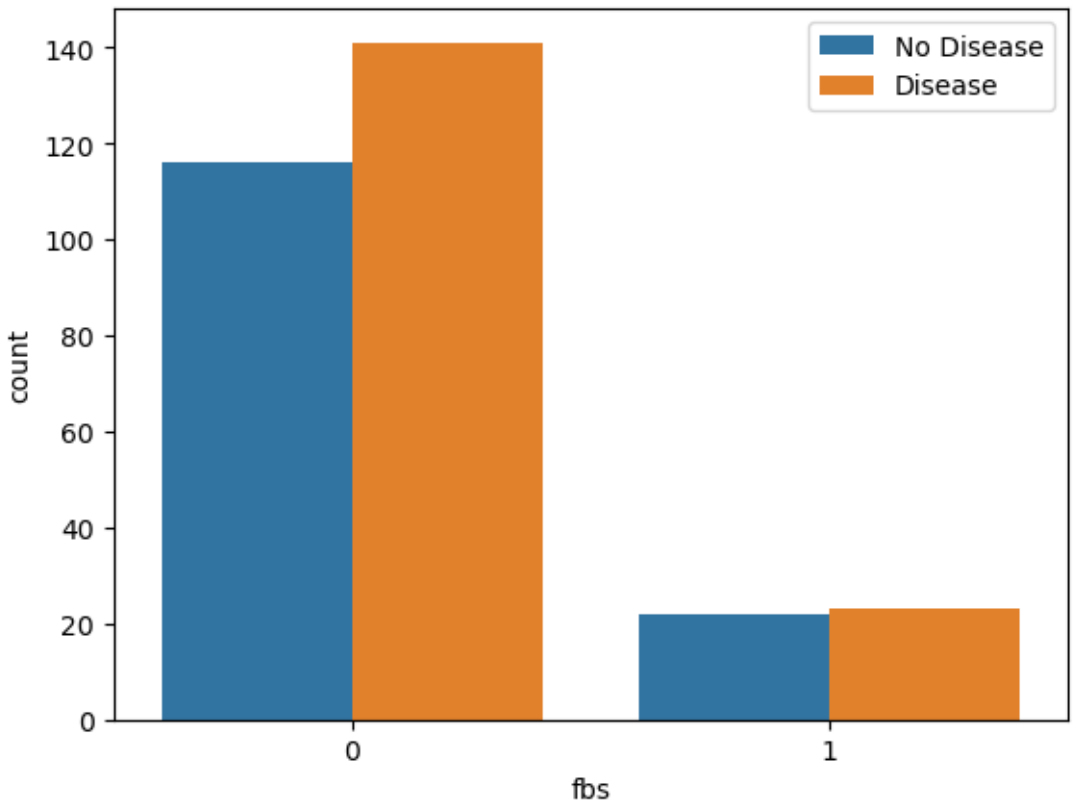
[69]:
```
df.columns
```

[69]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[70]:
```
df[['fbs','target']].value_counts().sort_values(ascending=False)
```

[70]:
```
fbs  target
0    1         141
     0         116
1    1          23
     0          22
Name: count, dtype: int64
```

[73]:
```
sns.countplot(x='fbs', hue='target',data=df)
plt.legend(labels=['No Disease','Disease'])
plt.show()
```

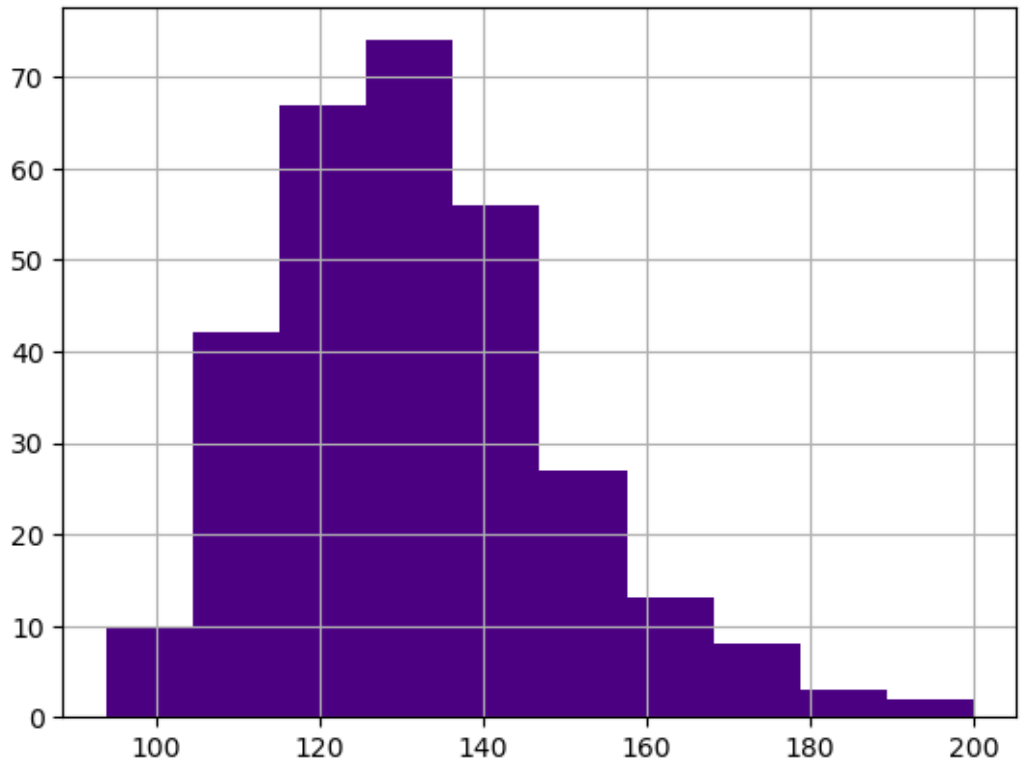


Question 8: Check Resting Blood Pressure Distribution

[74]:
```
df.columns
```

[74]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[97]:
```
df.trestbps.hist(color='indigo')
plt.show()
```

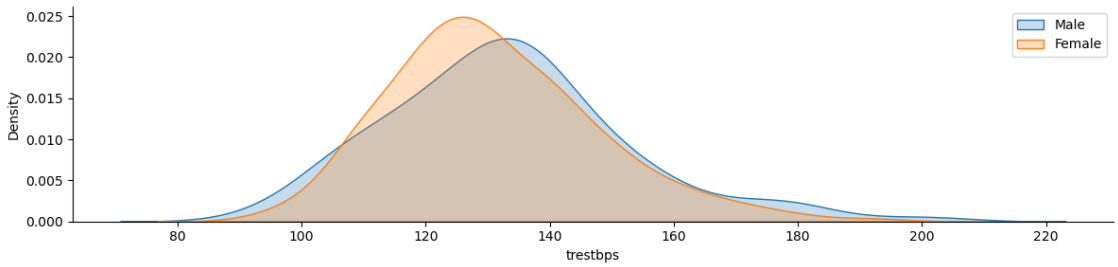


Question 9: compare Resting Blood Pressure as per sex column

[102]:
```
g=sns.FacetGrid(df, hue='sex',aspect=4)
g.map(sns.kdeplot,'trestbps',shade=True)
plt.legend(labels=['Male','Female'])
plt.show()
```

```
C:\Users\Josphat\anaconda3\new anaconda\Lib\site-
packages\seaborn\axisgrid.py:854: FutureWarning:

`shade` is now deprecated in favor of `fill`; setting `fill=True`.
```

```
This will become an error in seaborn v0.14.0; please update your code.

  func(*plot_args, **plot_kwargs)
C:\Users\Josphat\anaconda3\new anaconda\Lib\site-
packages\seaborn\axisgrid.py:854: FutureWarning:

`shade` is now deprecated in favor of `fill`; setting `fill=True`.
This will become an error in seaborn v0.14.0; please update your code.

  func(*plot_args, **plot_kwargs)
```

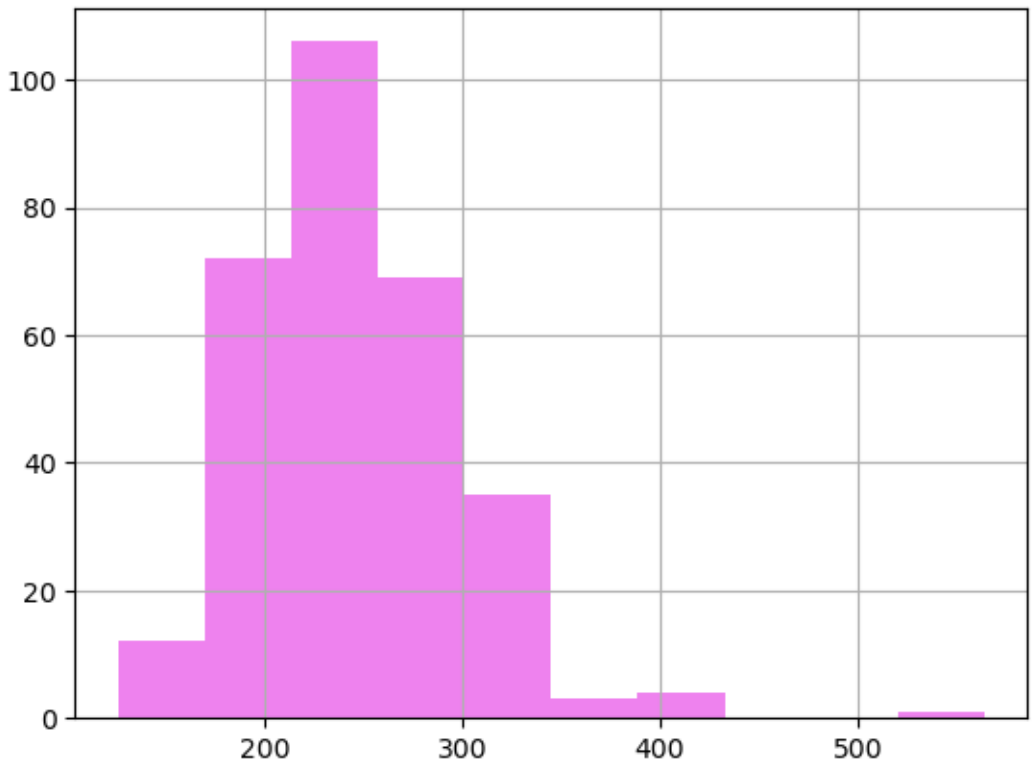


Question 10: Show distribution of Serum Cholestrol

[84]:
```
df.columns
```

[84]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[104]:
```
df.chol.hist(color='violet')
plt.show()
```



Question 11: Plot Continuous Variables

[90]:
```
df.columns
```

[90]:
```
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

[91]:
```
cate_val= []
cont_val= []

for column in df.columns:
    if df[column].nunique() <= 10:
        cate_val.append(column)
    else:
        cont_val.append(column)
```

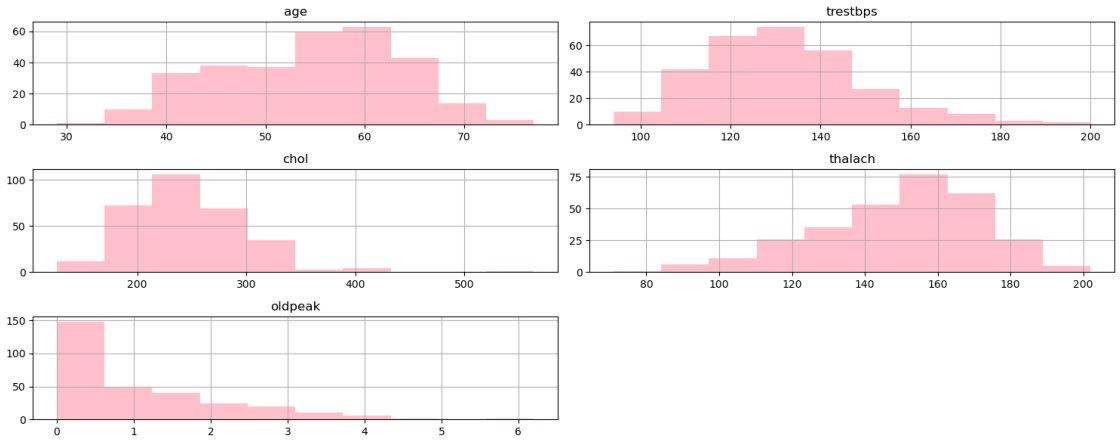
[92]:
```
cate_val
```

[92]:
```
['sex', 'cp', 'fbs', 'restecg', 'exang', 'slope', 'ca', 'thal', 'target']
```

[93]:
```
cont_val
```

[93]:
```
['age', 'trestbps', 'chol', 'thalach', 'oldpeak']
```

[106]:
```
df.hist(cont_val,figsize=(15,6),color='pink')
plt.tight_layout()
plt.show()
```



[ ]:
```

```