

```
In [80]: > #importing packages
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sb
import matplotlib.patches
import numpy as np
from pandas import set_option
```

```
In [2]: > #Loading data
file = 'C:\\Users\\OWNER\\Documents\\WORKSPACE\\heart.csv'
ds= pd.read_csv(file)
```

#### DATA EXPLORATORY ANALYSIS

```
In [4]: > ds.head(11)
```

```
Out[4]:
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target
0	63	1	3	145	233	1	0	150	0	2.3	0	0	1	1
1	37	1	2	130	250	0	1	187	0	3.5	0	0	2	1
2	41	0	1	130	204	0	0	172	0	1.4	2	0	2	1
3	56	1	1	120	236	0	1	178	0	0.8	2	0	2	1
4	57	0	0	120	354	0	1	163	1	0.6	2	0	2	1
5	57	1	0	140	192	0	1	148	0	0.4	1	0	1	1
6	56	0	1	140	294	0	0	153	0	1.3	1	0	2	1
7	44	1	1	120	263	0	1	173	0	0.0	2	0	3	1
8	52	1	2	172	199	1	1	162	0	0.5	2	0	3	1
9	57	1	2	150	168	0	1	174	0	1.6	2	0	2	1
10	54	1	0	140	239	0	1	160	0	1.2	2	0	2	1

```
In [5]: > ds.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):
#   Column      Non-Null Count  Dtype
---  -
0   age         303 non-null    int64
1   sex         303 non-null    int64
2   cp          303 non-null    int64
3   trestbps    303 non-null    int64
4   chol        303 non-null    int64
5   fbs         303 non-null    int64
6   restecg     303 non-null    int64
7   thalach     303 non-null    int64
8   exang       303 non-null    int64
9   oldpeak     303 non-null    float64
10  slope       303 non-null    int64
11  ca          303 non-null    int64
12  thal        303 non-null    int64
13  target      303 non-null    int64
dtypes: float64(1), int64(13)
memory usage: 33.2 KB
```

There are no missing values in the dataset.

```
In [120]: > ds.columns
```

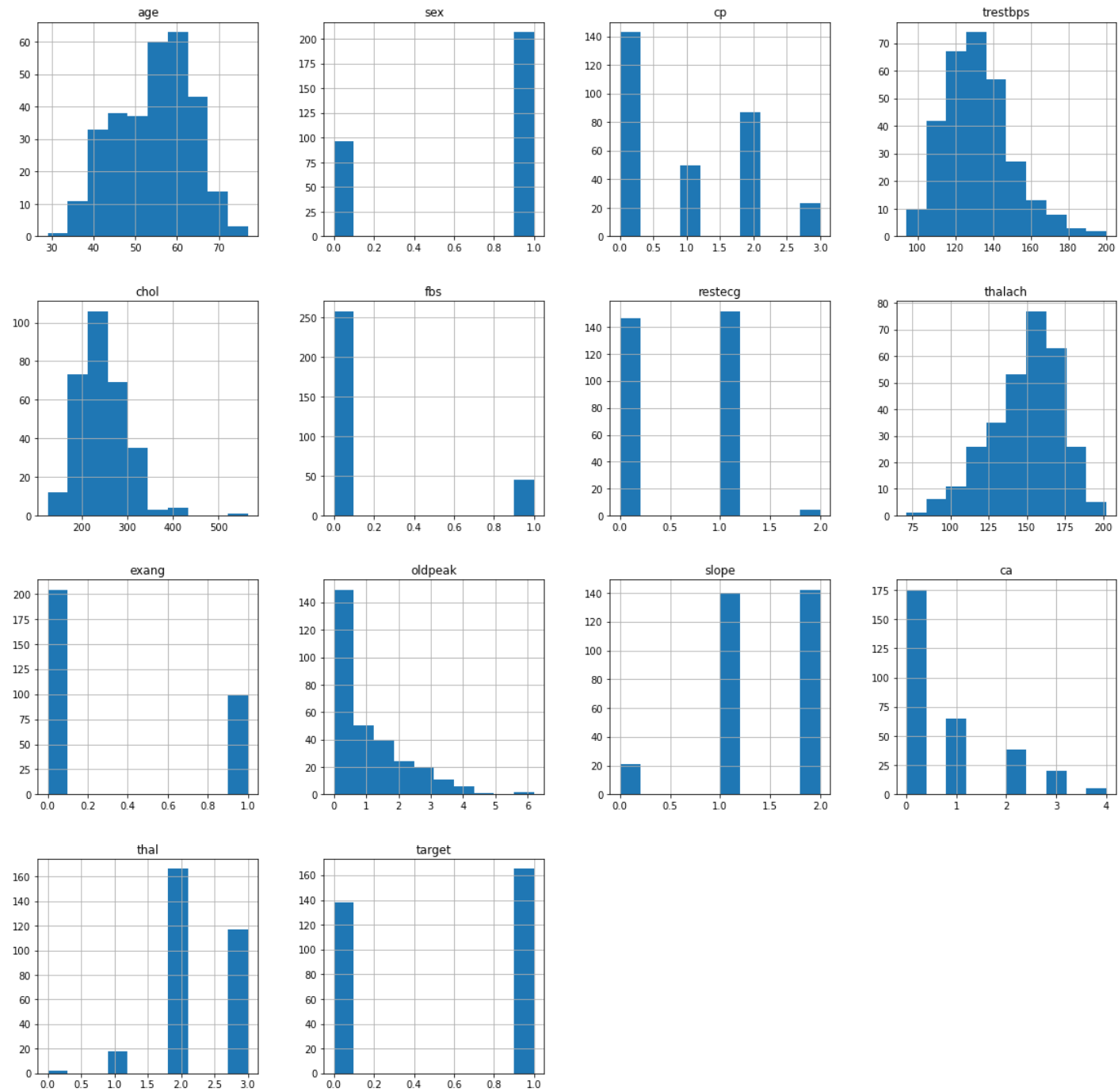
```
Out[120]: Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach', 'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'], dtype='object')
```

```
In [6]: ds.describe()

Out[6]:
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca
count	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000
mean	54.366337	0.683168	0.966997	131.623762	246.264026	0.148515	0.528053	149.646865	0.326733	1.039604	1.399340	0.729373
std	9.082101	0.466011	1.032052	17.538143	51.830751	0.356198	0.525860	22.905161	0.469794	1.161075	0.616226	1.022606
min	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	0.000000	71.000000	0.000000	0.000000	0.000000	0.000000
25%	47.500000	0.000000	0.000000	120.000000	211.000000	0.000000	0.000000	133.500000	0.000000	0.000000	1.000000	0.000000
50%	55.000000	1.000000	1.000000	130.000000	240.000000	0.000000	1.000000	153.000000	0.000000	0.800000	1.000000	0.000000
75%	61.000000	1.000000	2.000000	140.000000	274.500000	0.000000	1.000000	166.000000	1.000000	1.600000	2.000000	1.000000
max	77.000000	1.000000	3.000000	200.000000	564.000000	1.000000	2.000000	202.000000	1.000000	6.200000	2.000000	4.000000

```
In [7]: ds.hist()
plt.gcf().set_size_inches(20,20)
plt.show()
```



There is a poor distribution across the features as regards the scale but due to the fact we won't be building a model that won't be addressed right now.

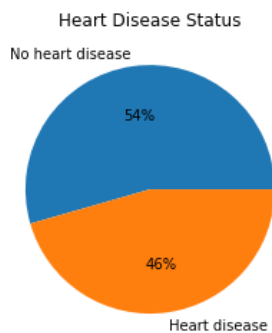
# UNIVARIATE ANALYSIS

Analysis of the key features to uncover more insight about the data.

```
In [8]: print(ds['target'].value_counts())
print(ds['target'].value_counts(normalize=True))

1    165
0    138
Name: target, dtype: int64
1    0.544554
0    0.455446
Name: target, dtype: float64
```

```
In [9]: %matplotlib inline
x=(ds['target'].value_counts())
labels = ['No heart disease', 'Heart disease']
fig, ax = plt.subplots()
ax.pie(x, labels = labels, autopct='%0f%%')
ax.set_title('Heart Disease Status')
plt.show()
```

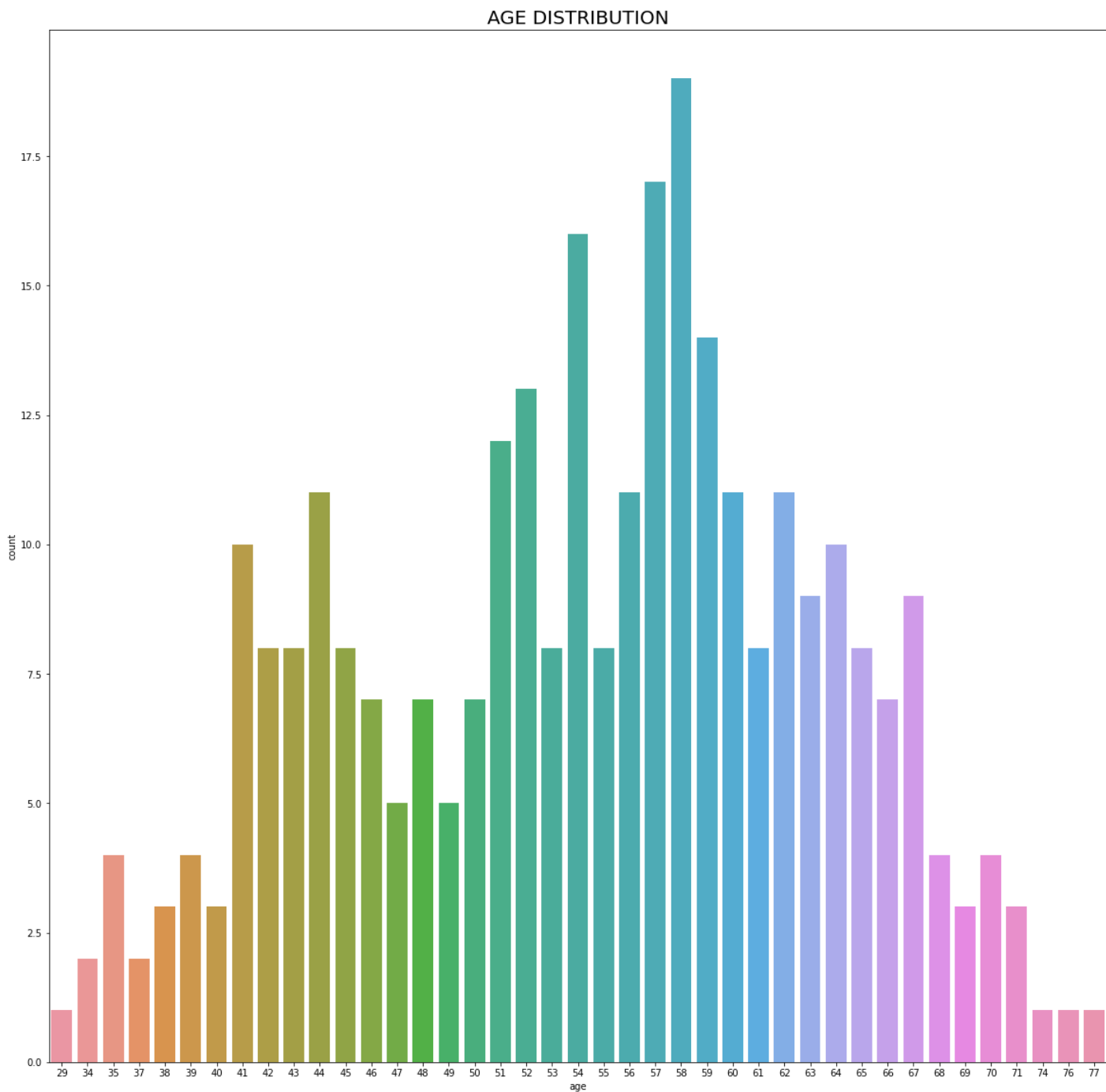


54% of the patients show no signs of heart disease and whilst 46% of the patients show signs of heart disease.

```
In [10]: print(ds['age'].value_counts(normalize=True))
```

```
58    0.062706
57    0.056106
54    0.052805
59    0.046205
52    0.042904
51    0.039604
62    0.036304
60    0.036304
44    0.036304
56    0.036304
41    0.033003
64    0.033003
63    0.029703
67    0.029703
65    0.026403
55    0.026403
61    0.026403
53    0.026403
45    0.026403
43    0.026403
42    0.026403
50    0.023102
66    0.023102
48    0.023102
46    0.023102
49    0.016502
47    0.016502
70    0.013201
39    0.013201
68    0.013201
35    0.013201
69    0.009901
40    0.009901
38    0.009901
71    0.009901
37    0.006601
34    0.006601
76    0.003300
29    0.003300
74    0.003300
77    0.003300
Name: age, dtype: float64
```

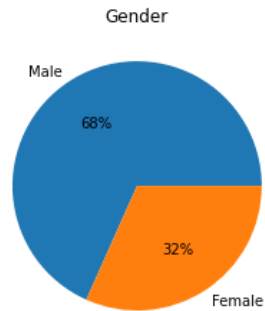
```
In [11]: ▶ plt.figure(figsize = (20,20))
sb.countplot(data = ds, x='age')
plt.title('AGE DISTRIBUTION', fontsize = 20)
plt.show()
```



Most of the patients are within the ages of 54,57 and 58 years, the youngest patient is 29 years old whilst the oldest is 77. from the data set we can also see that the average age of the patients is 54 years which gives a clear indication that most of the patients are within the bracket of the elderly.

```
In [12]: ▶ print (ds['sex'].value_counts())
1      207
0       96
Name: sex, dtype: int64
```

```
In [13]: x=(ds['sex'].value_counts())
labels = ['Male', 'Female']
fig, ax = plt.subplots()
ax.pie(x, labels = labels, autopct='%.0f%%')
ax.set_title('Gender')
plt.show()
```

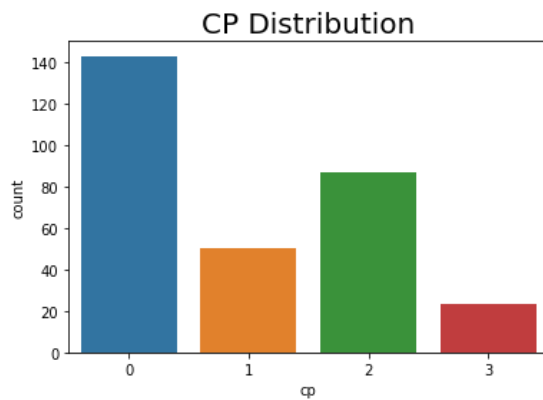


68% of the patients are male whilst 32% are female which give us information that more of the patients are male.

```
In [14]: print(ds['cp'].value_counts())
print(ds['cp'].value_counts(normalize=True))
```

```
0    143
2     87
1     50
3     23
Name: cp, dtype: int64
0    0.471947
2    0.287129
1    0.165017
3    0.075908
Name: cp, dtype: float64
```

```
In [15]: sb.countplot(data = ds, x='cp')
plt.title('CP Distribution', fontsize = 20)
plt.show()
```

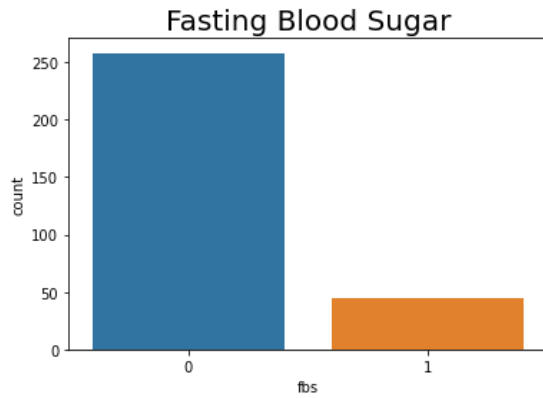


47% (143) of the patients are asymptomatic 28% (87) of the patients are atypical angina 16% (50) of the patients experience non-anginal pain 7%(23) of the patients experience typical angina. From the above we can see that most of the patients show no symptoms of chest related pain

```
In [16]: print(ds['fbs'].value_counts())
print(ds['fbs'].value_counts(normalize=True))
```

```
0    258
1     45
Name: fbs, dtype: int64
0    0.851485
1    0.148515
Name: fbs, dtype: float64
```

```
In [17]: sb.countplot(data = ds, x='fbs')
plt.title('Fasting Blood Sugar', fontsize = 20)
plt.show()
```

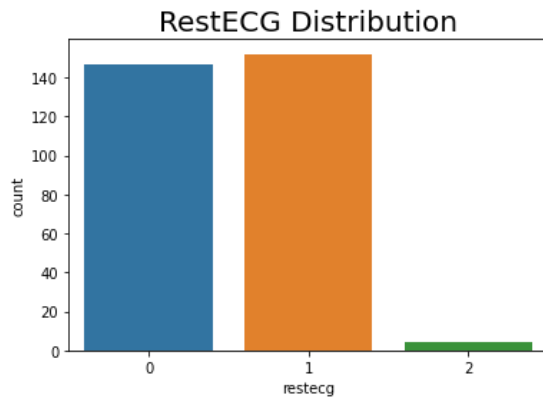


85% of the patients have a blood sugar level that is less than 120 mg/dl

```
In [18]: print (ds['restecg'].value_counts())
print(ds['restecg'].value_counts(normalize=True))

1    152
0    147
2      4
Name: restecg, dtype: int64
1    0.501650
0    0.485149
2    0.013201
Name: restecg, dtype: float64
```

```
In [19]: sb.countplot(data = ds, x='restecg')
plt.title('RestECG Distribution', fontsize = 20)
plt.show()
```



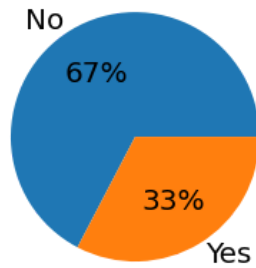
50% of the patients suffer from a high hypertrophy whilst 48% have a normal heart condition.

```
In [20]: print (ds['exang'].value_counts())

0    204
1     99
Name: exang, dtype: int64
```

```
In [34]: x = (ds['exang'].value_counts())
#plt.rcParams.update({'font.size': 20})
labels = ['No', 'Yes']
fig, ax = plt.subplots()
ax.pie(x, labels = labels, autopct='%0f%%')
ax.set_title('Exercise Induced Heart Pain')
plt.show()
```

## Exercise Induced Heart Pain

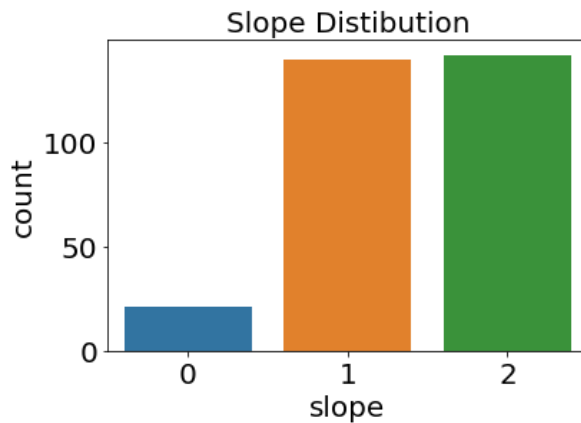


67% of the patients do not suffer from exercise induced heart pain whilst 33% of the patients do.

```
In [22]: print(ds['slope'].value_counts())
print(ds['slope'].value_counts(normalize=True))
```

```
2    142
1    140
0     21
Name: slope, dtype: int64
2    0.468647
1    0.462046
0    0.069307
Name: slope, dtype: float64
```

```
In [23]: sb.countplot(data = ds, x='slope')
plt.title('Slope Distribution', fontsize = 20)
plt.show()
```

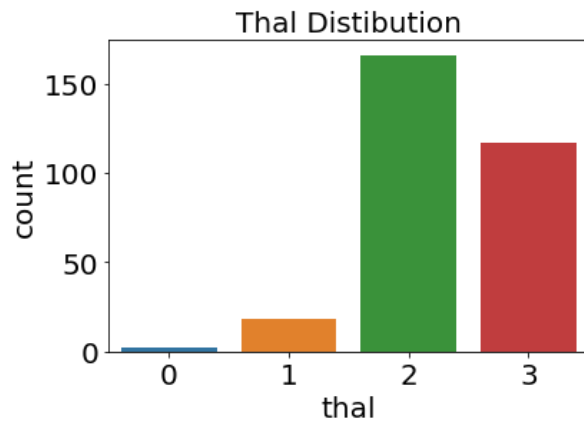


From the above chart, 46% of the patients show a flat sloping st segment, 46% of the patients also show an upsloping st segment and 6% of the patients show a downsloping st segment.

```
In [24]: print(ds['thal'].value_counts())
print(ds['thal'].value_counts(normalize=True))
```

```
2    166
3    117
1     18
0       2
Name: thal, dtype: int64
2    0.547855
3    0.386139
1    0.059406
0    0.006601
Name: thal, dtype: float64
```

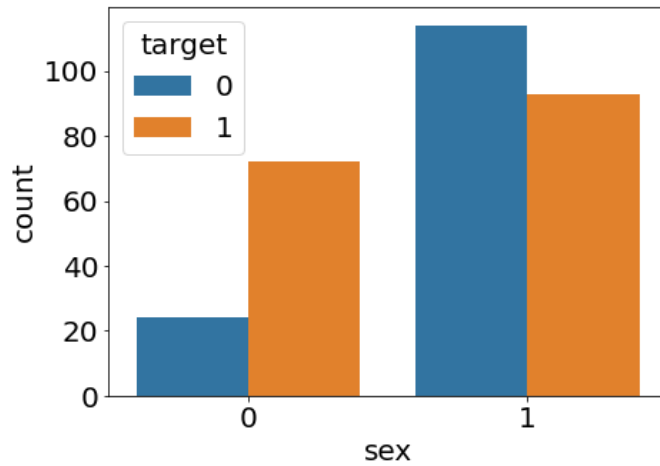
```
In [25]: ▶ sb.countplot(data = ds, x='thal')
plt.title('Thal Distribution', fontsize = 20)
plt.show()
```



from the above chart, 54% of the patients have fixed defect thal, 38% of the patients have a reversible defect thal whilst 5% of the patients have a normal thal.

#### BIVARIATE ANALYSIS

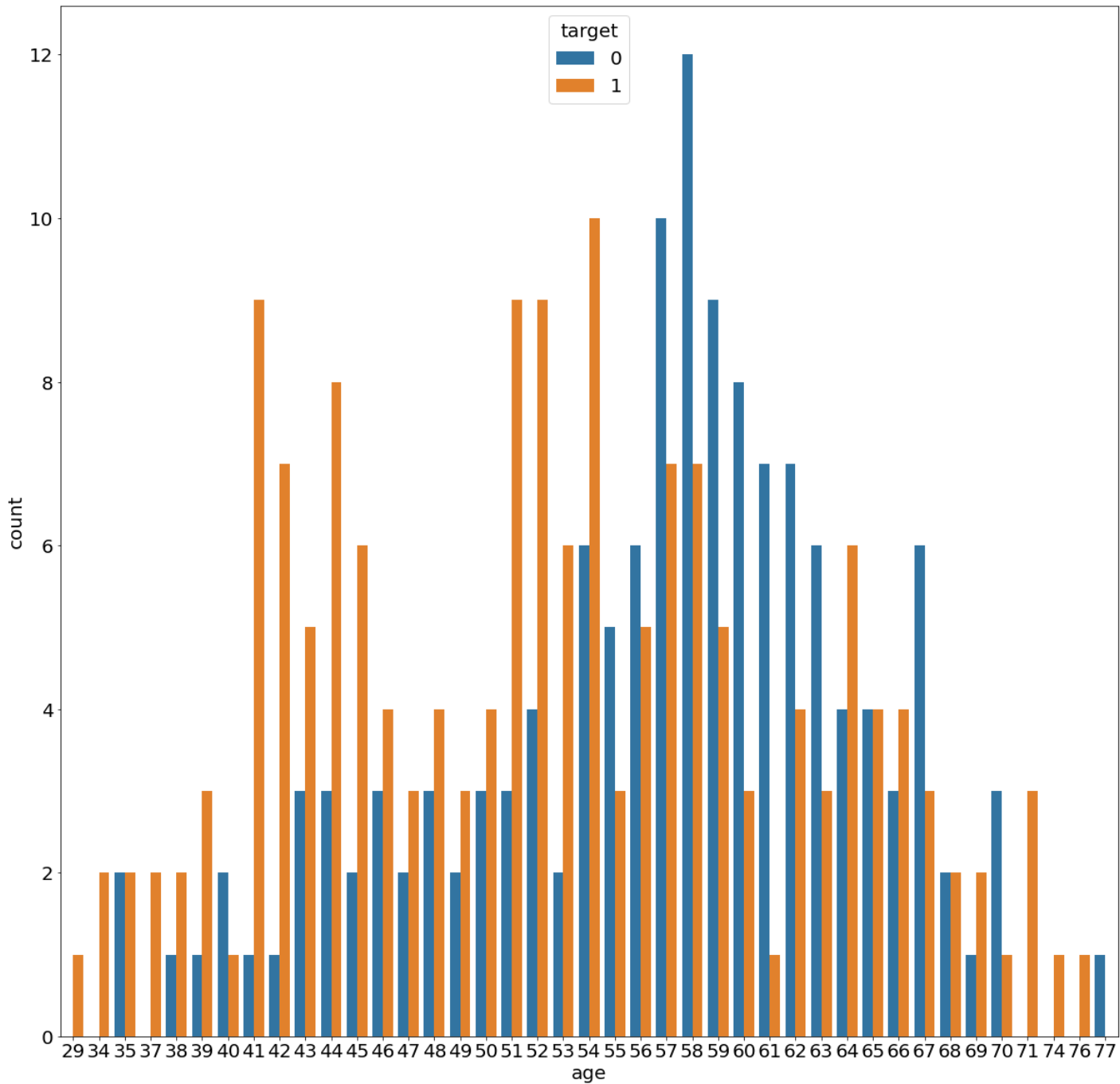
```
In [44]: ▶ plt.figure(figsize = (7,5))
sb.countplot(x="sex", hue="target", data=ds)
plt.show()
```



From the chart we can see that more of the male patients tend to suffer from hear disease than the female patients.

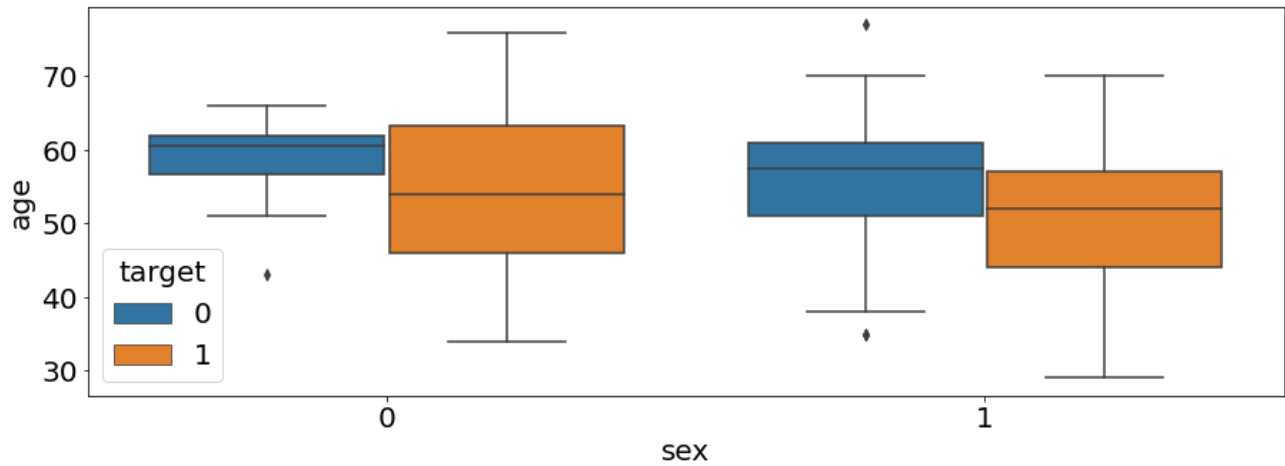


```
In [48]: ▶ plt.figure(figsize = (20,20))
sb.countplot(x="age", hue="target", data=ds)
plt.show()
```



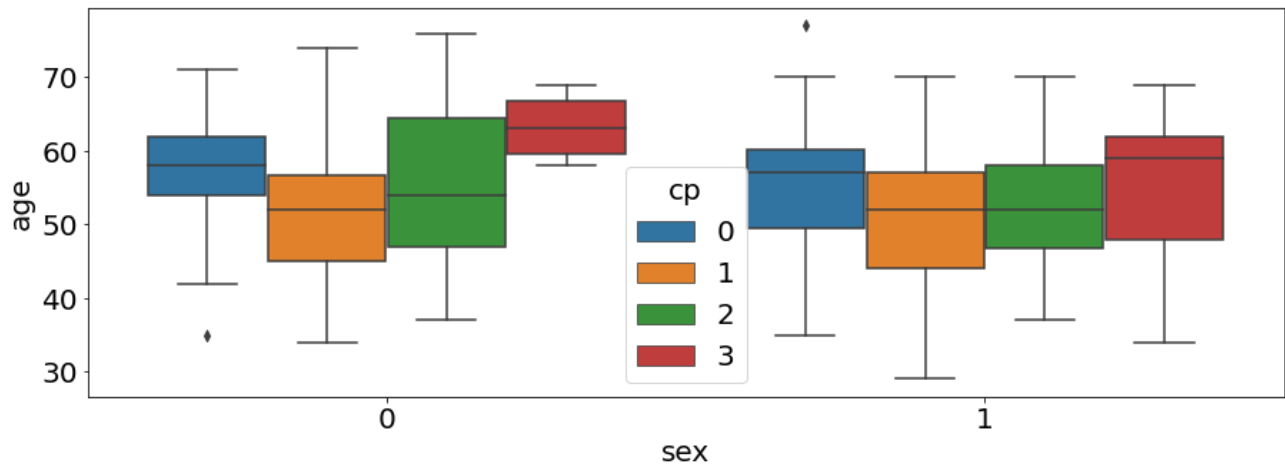
from the chart we can see that people within the ages of 57, 58, 59, 60, had the highest occurrence of heart disease. the age group of 29,34,37,71,74 and 76 had no cases of heart disease

```
In [57]: ▶ plt.figure(figsize = (15,5))
sb.boxplot(x="sex", y="age", hue="target", data=ds);
```



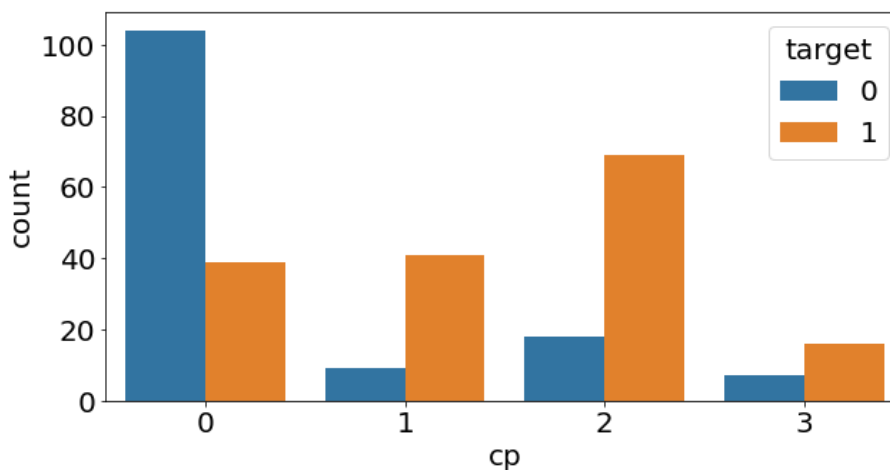
from the chart we can see that men between the ages from 55-60 have a higher occurrence of heart disease whilst women between the ages of 45-60 had lower occurrence of heart disease

```
In [58]: ▶ plt.figure(figsize = (15,5))
sb.boxplot(x="sex", y="age", hue="cp", data=ds);
```



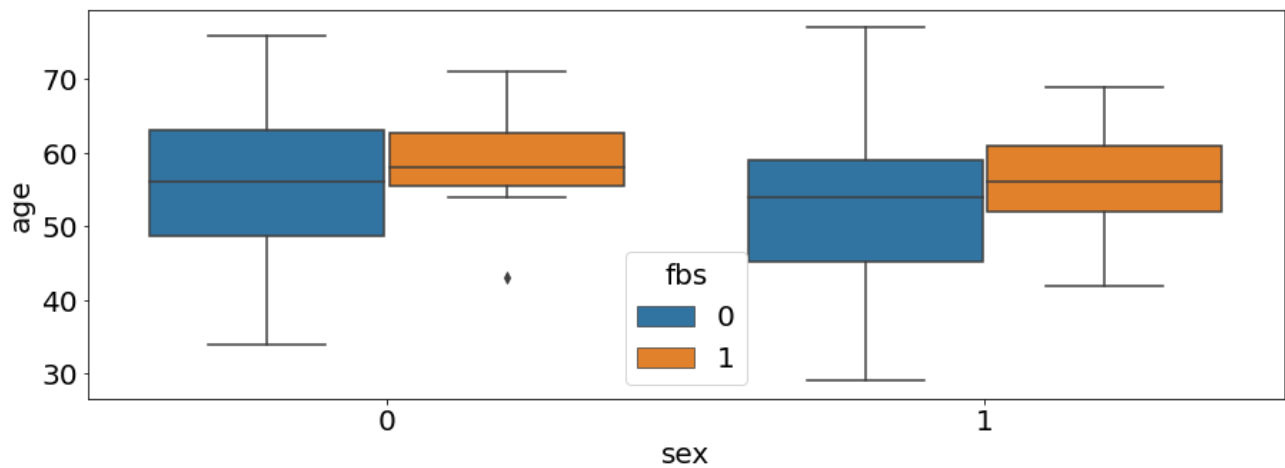
This plot shows us sex and age related chest pain cases from the patients male patients between the ages of 50-60 were asymptomatic female patients between the ages of 55-62 were also asymptomatic male patients between the ages of 45-58 were atypical angina female patients between the ages of 45-58 were atypical angina also male patients between the ages of 48-60 had non-anginal whilst female patients between the ages of 47-64 had non-anginal also male patients between the ages of 49-62 had typical angina whilst female patients between the ages of 62-65 had typical angina chest pain

```
In [59]: ▶ plt.figure(figsize = (10,5))
sb.countplot(x="cp", hue="target", data=ds)
plt.show()
```



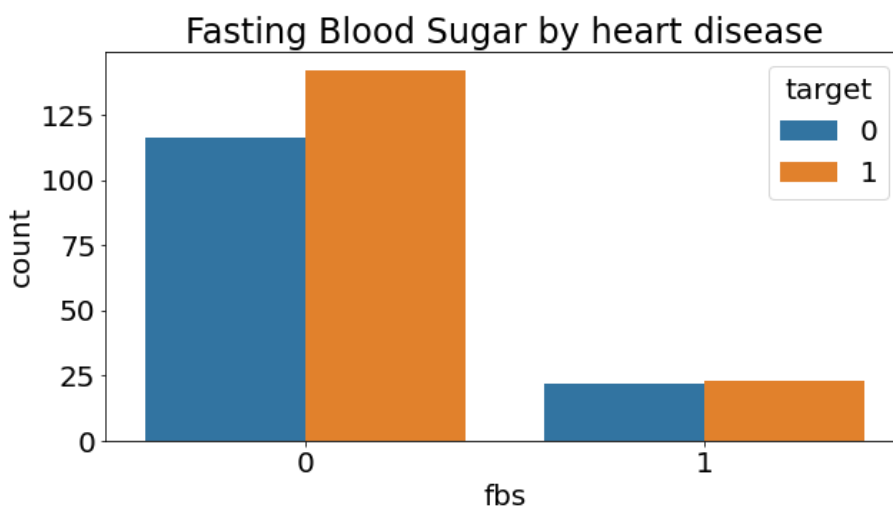
from the above chart we can see that even though the patients were asymptomatic, they were still liable to suffer from heart disease.

```
In [66]: ▶ plt.figure(figsize = (15,5))
sb.boxplot(x="sex", y="age", hue="fbs", data=ds);
```



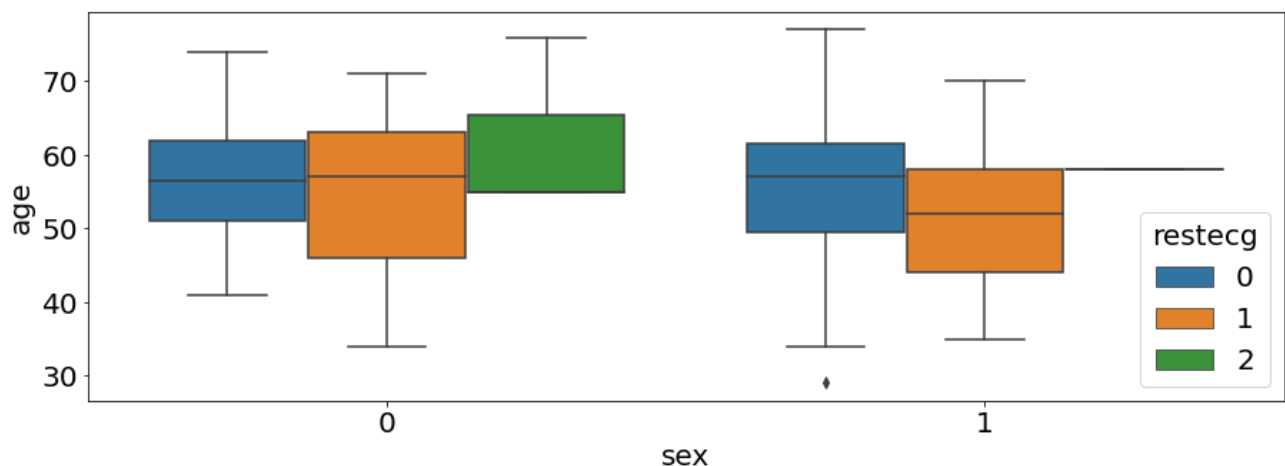
In [ ]: ▶ This chart shows a boxplot of blood sugar leve between male and female patients also taking into consideration their ages. we can see that male between the ages of 48-60 ten to have a high blood sugar level whilst females between the the ages of 50-63 also tend to have a high level.

```
In [69]: ▶ plt.figure(figsize = (10,5))
sb.countplot(x="fbs", hue="target", data=ds)
plt.title('Fasting Blood Sugar by heart disease')
plt.show()
```



This chart shows the Blood Sugar level of pateints in comparison to heart disease we can see that those with sugar levels below 120 mg/dl have a higher chance with no heart disease and also heart disease this might also be a clear indication that blood sugar level alone might not give the clearest indication of heart disease.

```
In [72]: ▶ plt.figure(figsize = (15,5))
sb.boxplot(x="sex", y="age", hue="restecg", data=ds);
```



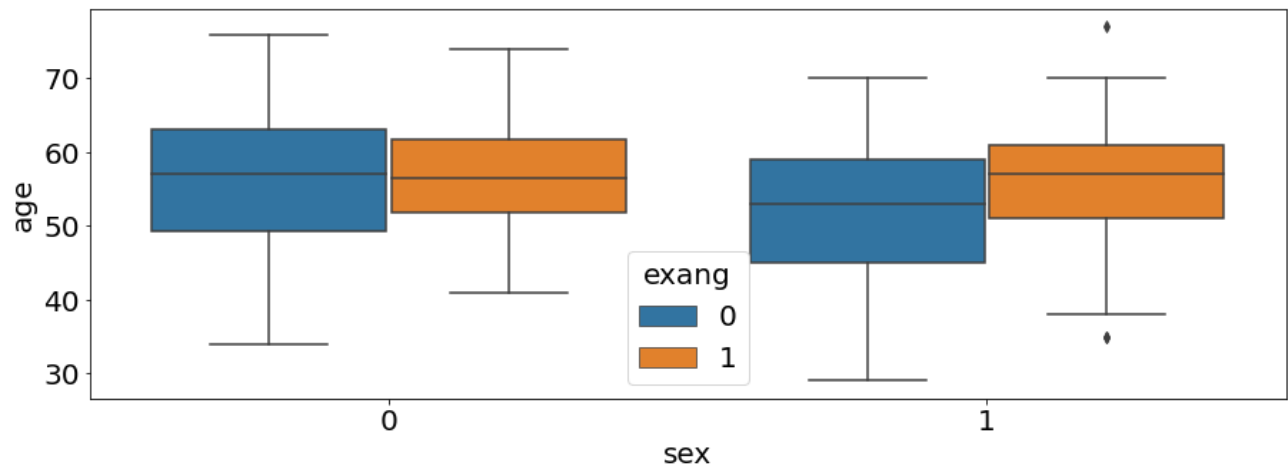
This chart shows the Resting ECG of both Female and Male and their ages. Females within the ages of 52-62 having a definite left ventricular hypertrophy whilst Males within the ages of 50-60 having a definite left ventricular hypertrophy Females within the ages of 47-64 having a normal resting ECG whilst Males within the ages of 45-56 having a normal resting ECG Females within the ages of 55-65 having a ST-T wave abnormality

```
In [74]: ▶ plt.figure(figsize = (10,5))
sb.countplot(x="restecg", hue="target", data=ds)
plt.title('Resting ECG by heart disease')
plt.show()
```



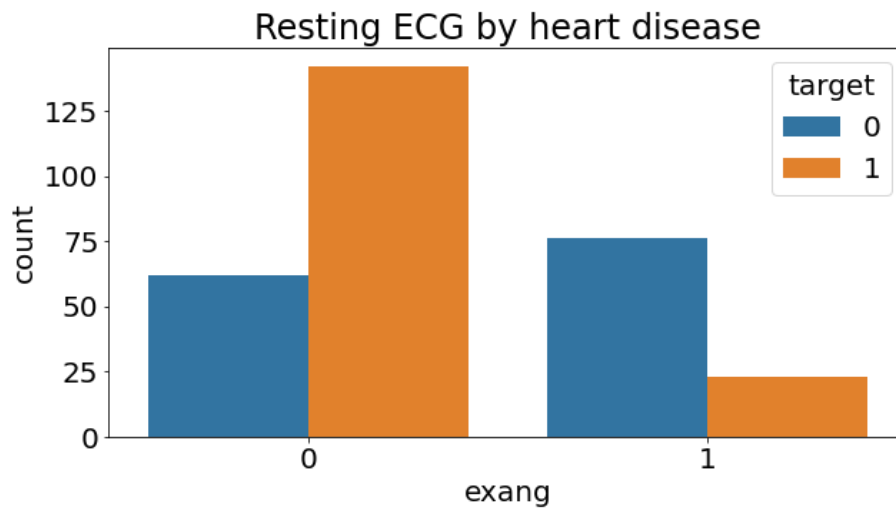
This chart compares The resting ECG to heart disease. From the chart we can see that patients with a definite left ventricular hypertrophy are at a higher risk of heart disease whilst patients with a normal resting ECG having a higher rate of no heart disease.

```
In [75]: ▶ plt.figure(figsize = (15,5))
sb.boxplot(x="sex", y="age", hue="exang", data=ds);
```



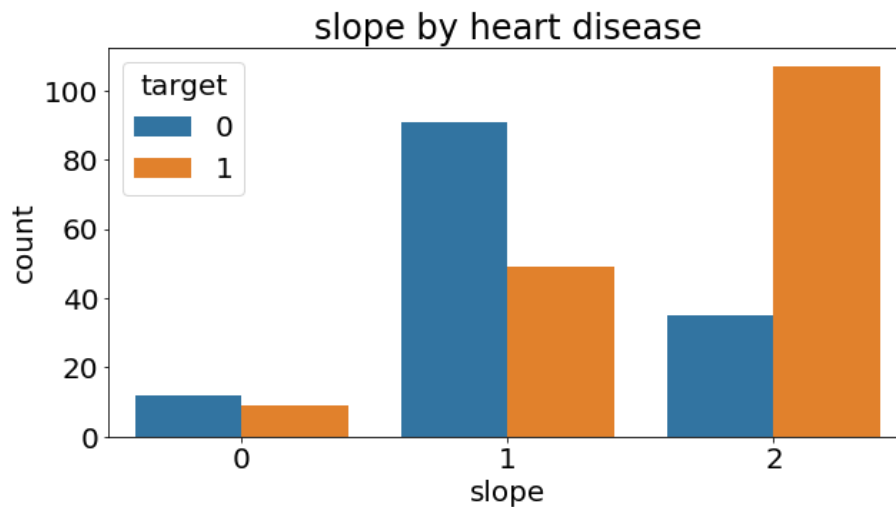
This chart shows Exercise related pain in the chest comparing it to male and female and thier ages Females withing the ages of 50-64 expereined exercise Induced Angina Males withing the ages of 46-60 experienced exercise Induced Angina

```
In [76]: ▶ plt.figure(figsize = (10,5))
sb.countplot(x="exang", hue="target", data=ds)
plt.title('Resting ECG by heart disease')
plt.show()
```



This chart shows how Exercise Induced Angina compares to heart disease from the chart we can see that the likelihood of no pain in the chest area during exercise , the less likely you are to have heart disease whilst the more you feel pain during exercise the higher the chance of heart disease.

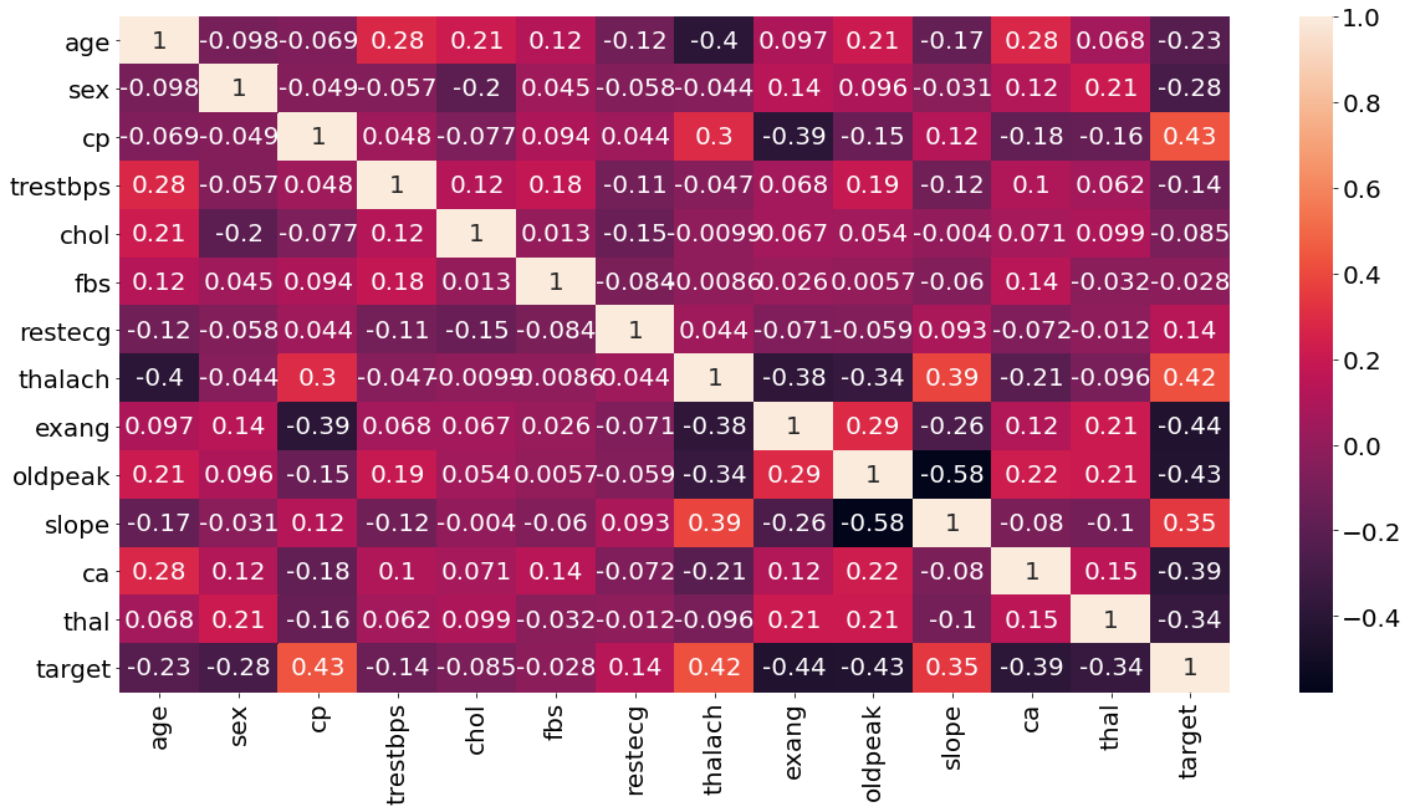
```
In [77]: ▶ plt.figure(figsize = (10,5))
sb.countplot(x="slope", hue="target", data=ds)
plt.title('slope by heart disease')
plt.show()
```



This chart shows the slope (the slope of the peak exercise ST segment) as it compares to heart disease a flatslope shows a high tendency of heart disease whilst an upslope shows a high tendency of no heart disease.

correlation Heat Map

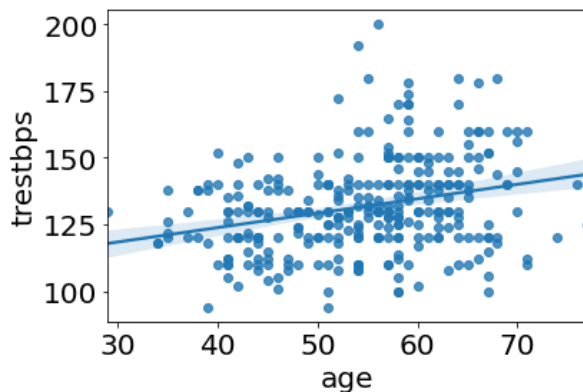
```
In [82]: plt.figure(figsize = (20,10))
sb.heatmap(ds.corr(), annot = True)
set_option('display.width', 1000)
```



from the above heat map, there seems to be slight correlation between a few features. We will try to explore that further to tease out more insights.

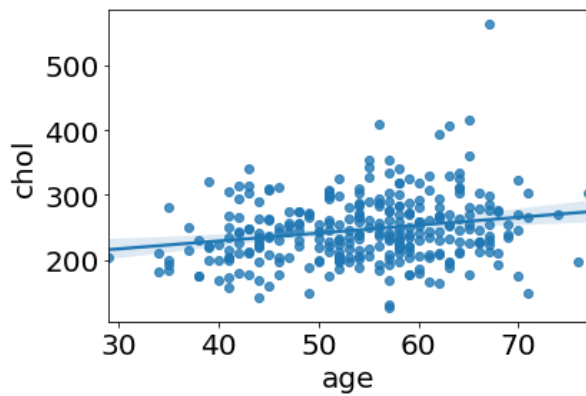
## Linear regression analysis

```
In [90]: sb.regplot(x="age", y="trestbps", data=ds);
```



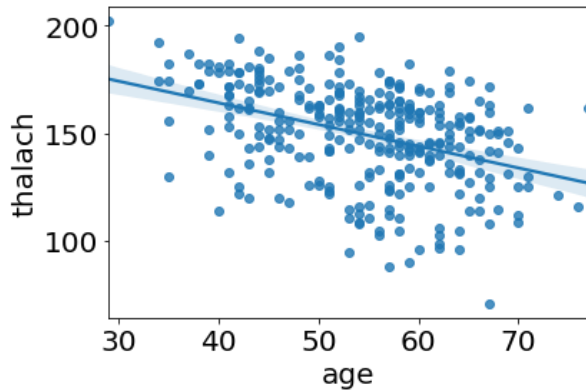
there is a positive correlation between age and Resting Blood Pressure. The older the age the higher the Resting Blood Pressure.

```
In [91]: sb.regplot(x="age", y="chol", data=ds);
```



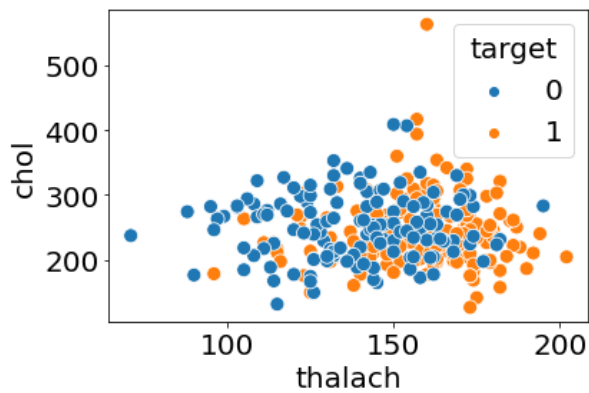
there is a positive linear correlation between age and cholesterol the older the age, the higher the cholesterol level.

```
In [92]: sb.regplot(x="age", y="thalach", data=ds);
```



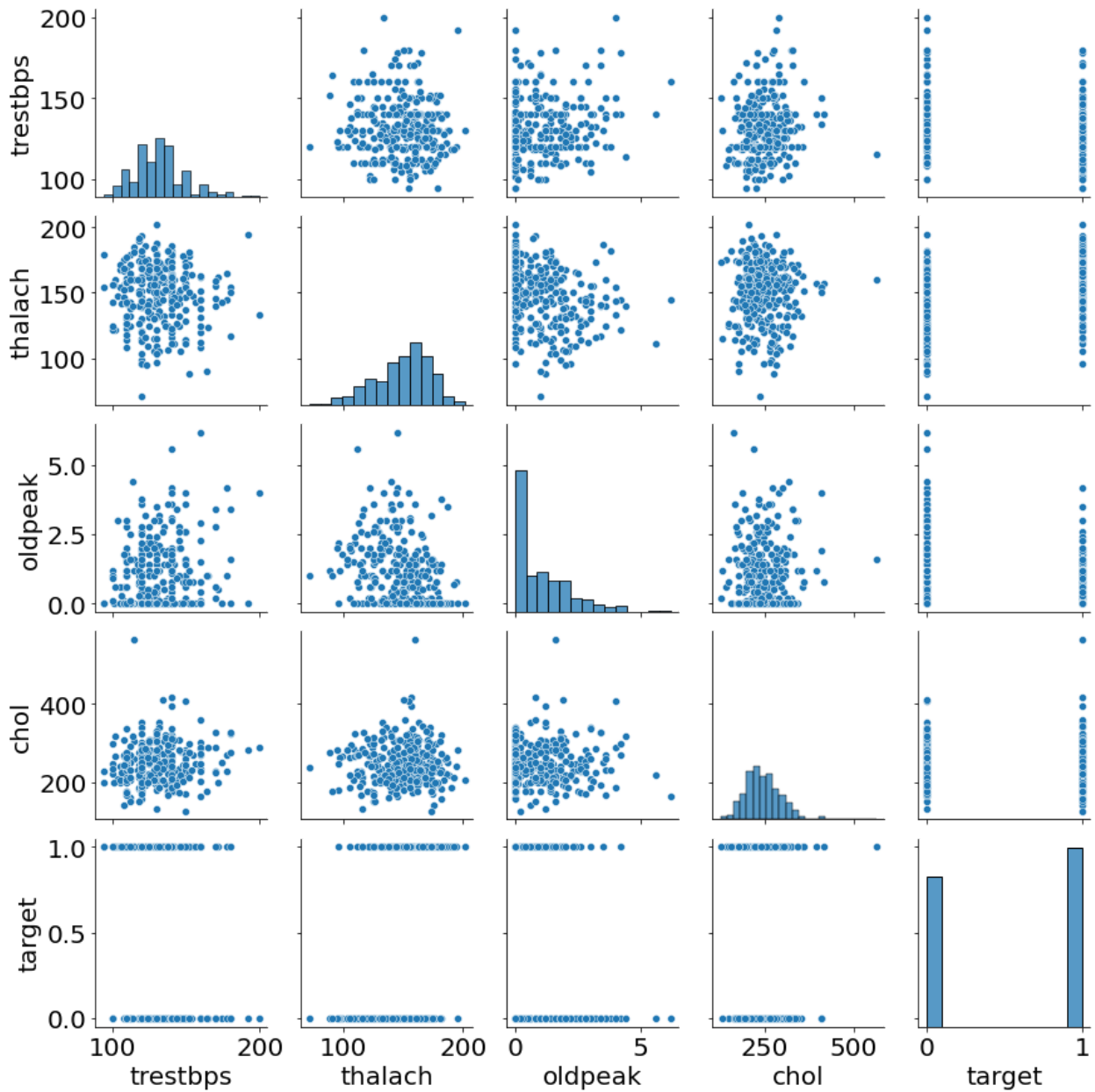
There is a negative linear relationship between age and maximum heart rate achieved by patients, as heart rate increases age decreases.

```
In [100]: sb.scatterplot(x =ds.thalach, # X-axis  
                        y = ds.chol, # Y-axis  
                        hue=ds.target,  
                        s=100);
```

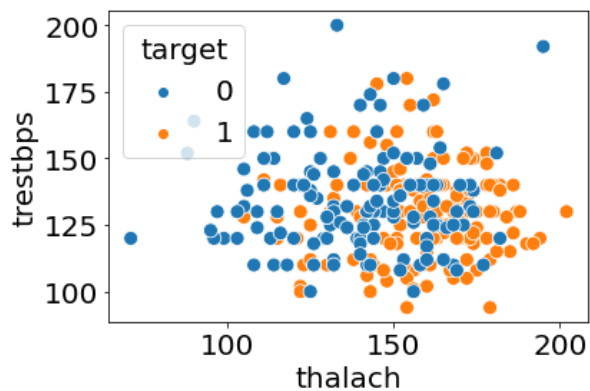


from the scatter plot we can see that as cholesterol and heart rate rises the likelihood of heart disease also does.

```
In [121]: sb.pairplot(ds[['trestbps', 'thalach', 'oldpeak', 'chol', 'target']])
plt.show()
```



```
In [114]: sb.scatterplot(x =ds.thalach, # X-axis
y = ds.trestbps, # Y-a
hue=ds.target,
s=100);
```



A high heart rate and also a high blood sugar level has an effect on the rate of heart disease as it can be seen on this chart.



```
In [ ]: ► conclusion
from the above I was able to carry out a
Univariate, Bivariate and Linear regression analysis on the data
features such as age, chol(cholesterol), fbs (fasting blood sugar)
trestbps(resting blood sugar), thalach (maximum heart rate)
restecg (Resting ECG), cp(Chest Pain Type) were the major drivers
in terms of predicting the likelihood of a patient getting heart disease
relationships across these feature enables significant knowledge about
the prediction of heart disease.
```