

```
In [1]: import pandas as pd
df = pd.read_csv(r"C:/Users/Administrator/Downloads/Heart Disease UCI.csv")
df
```

Out[1]:

	age	sex	cp	trestbps	chol	fbps	restecg	thalach	exang	oldpeak	slope	ca	tl
0	69	1	0	160	234	1	2	131	0	0.1	1	1	
1	69	0	0	140	239	0	0	151	0	1.8	0	2	
2	66	0	0	150	226	0	0	114	0	2.6	2	0	
3	65	1	0	138	282	1	2	174	0	1.4	1	1	
4	64	1	0	110	211	0	2	144	1	1.8	1	0	
...	...	...	...	...	...	...	...	...	...	...	...	...	
292	40	1	3	152	223	0	0	181	0	0.0	0	0	
293	39	1	3	118	219	0	0	140	0	1.2	1	0	
294	35	1	3	120	198	0	0	130	1	1.6	1	0	
295	35	0	3	138	183	0	0	182	0	1.4	0	0	
296	35	1	3	126	282	0	2	156	1	0.0	0	0	

297 rows × 14 columns

```
In [53]: # To Display 10 rows
df.head(10)
```

```
# check data types
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 297 entries, 0 to 296
Data columns (total 16 columns):
 #   Column      Non-Null Count  Dtype  
 --- 
 0   age         297 non-null    int64  
 1   sex         297 non-null    int64  
 2   cp          297 non-null    int64  
 3   trestbps    297 non-null    int64  
 4   chol        297 non-null    int64  
 5   fbs         297 non-null    int64  
 6   restecg     297 non-null    int64  
 7   thalach     297 non-null    int64  
 8   exang       297 non-null    int64  
 9   oldpeak     297 non-null    float64 
 10  slope        297 non-null    int64  
 11  ca          297 non-null    int64  
 12  thal        297 non-null    int64  
 13  condition    297 non-null    int64  
 14  risk_score   297 non-null    float64 
 15  risk_level   297 non-null    object  
dtypes: float64(2), int64(13), object(1)
memory usage: 37.3+ KB
```

```
In [54]: # missing values
df.isnull().sum()
```

```
Out[54]: 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
          condition 0
          risk_score 0
          risk_level 0
          dtype: int64
```

```
In [5]: # Display Statistics
df.describe()
```

	age	sex	cp	trestbps	chol	fbs	restecg
<b>count</b>	297.000000	297.000000	297.000000	297.000000	297.000000	297.000000	297.000000
<b>mean</b>	54.542088	0.676768	2.158249	131.693603	247.350168	0.144781	0.994667
<b>std</b>	9.049736	0.468500	0.964859	17.762806	51.997583	0.352474	0.994938
<b>min</b>	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	0.000000
<b>25%</b>	48.000000	0.000000	2.000000	120.000000	211.000000	0.000000	0.000000
<b>50%</b>	56.000000	1.000000	2.000000	130.000000	243.000000	0.000000	1.000000
<b>75%</b>	61.000000	1.000000	3.000000	140.000000	276.000000	0.000000	2.000000
<b>max</b>	77.000000	1.000000	3.000000	200.000000	564.000000	1.000000	2.000000

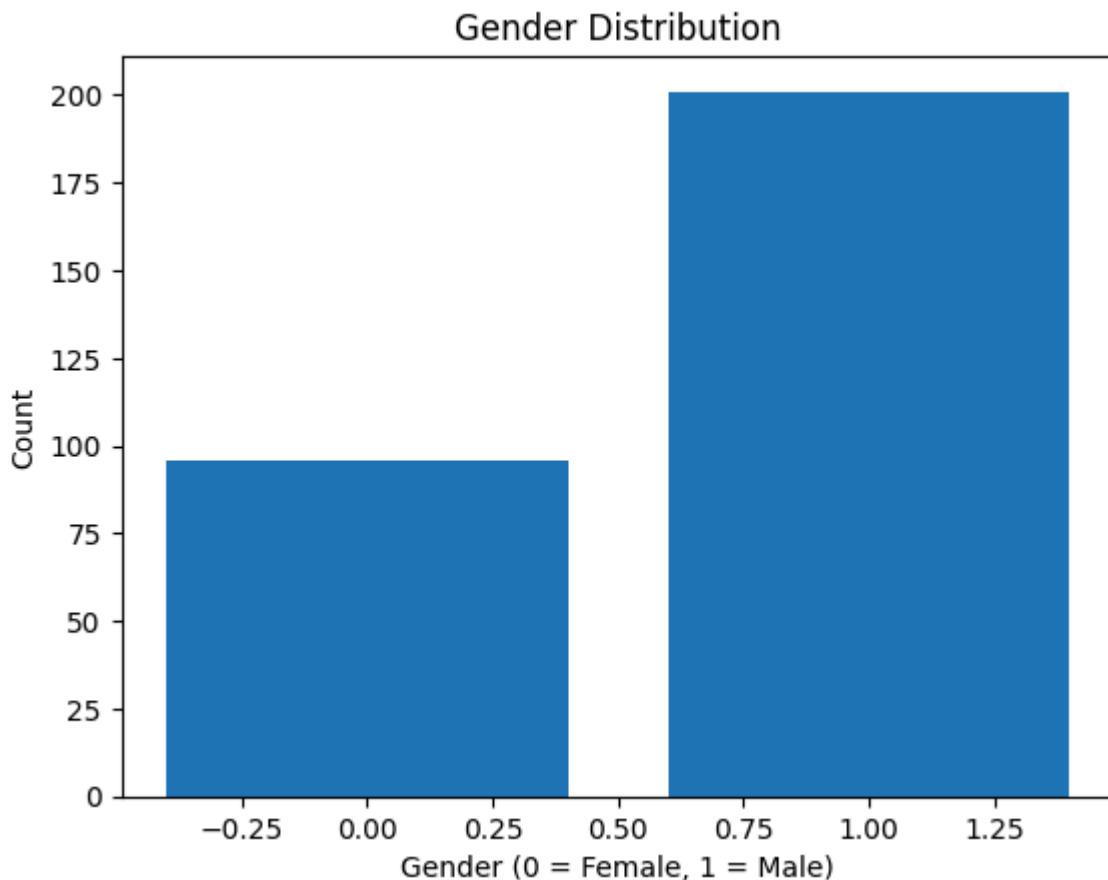
```
In [51]: # 2.Gender Distribution Analysis
```

```
#Count number of males and females
import numpy as np
import matplotlib.pyplot as plt
gender_count = df['sex'].value_counts()
print(gender_count)

#Calculate percentage distribution using NumPy
gender_percentage = (gender_count.values / np.sum(gender_count.values)) * 100
print(gender_percentage)

plt.bar(gender_count.index, gender_count.values)
plt.xlabel("Gender (0 = Female, 1 = Male)")
plt.ylabel("Count")
plt.title("Gender Distribution")
plt.show()
```

```
sex
1    201
0     96
Name: count, dtype: int64
[67.67676768 32.32323232]
```

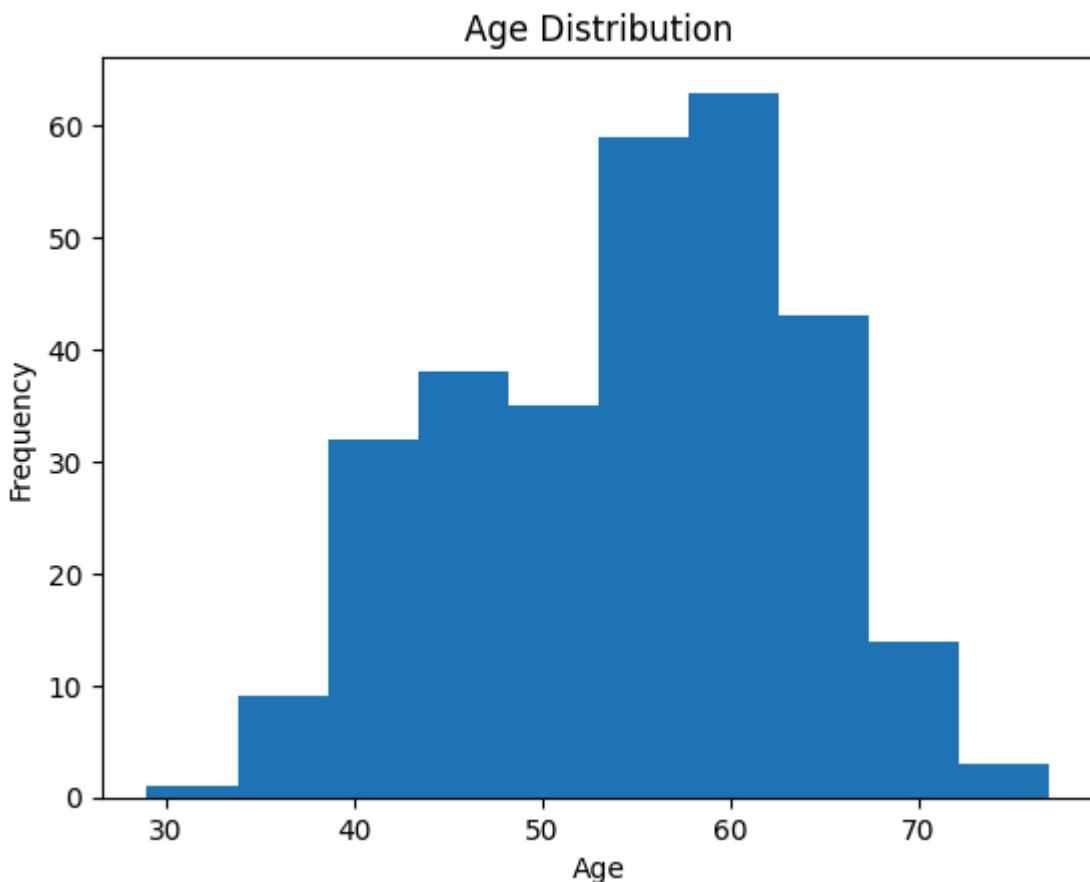


```
In [52]: # 3.Age Analysis
```

```
# to find age
print("Minimum Age : ", df['age'].min())
print("Maximum Age : ", df['age'].max())
print("Mean Age    : ", df['age'].mean())
print("Median Age   : ", df['age'].median())

import matplotlib.pyplot as plt
plt.hist(df['age'], bins=10)
plt.xlabel("Age")
plt.ylabel("Frequency")
plt.title("Age Distribution")
plt.show()
```

```
Minimum Age : 29
Maximum Age : 77
Mean Age    : 54.54208754208754
Median Age   : 56.0
```



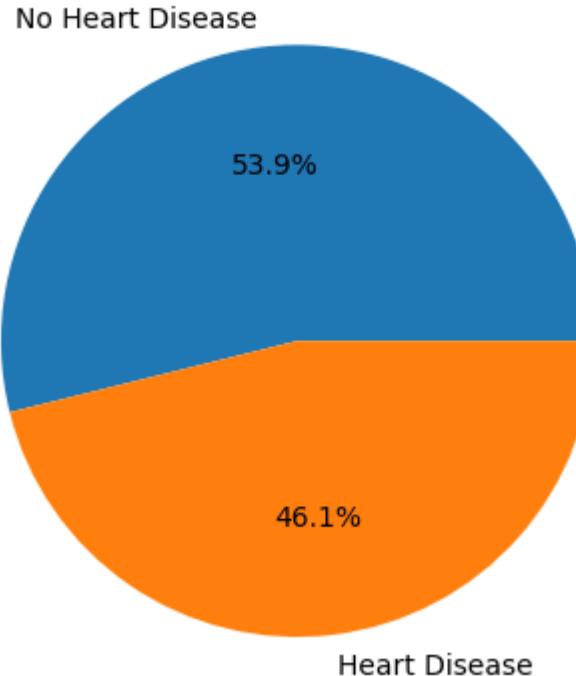
```
In [20]: # 4.Target Variable Analysis

#Count number of patients with and without heart disease
df['condition'].value_counts()

#Plot pie chart
import matplotlib.pyplot as plt
counts = df['condition'].value_counts()
plt.figure()
plt.pie(counts, labels=['No Heart Disease', 'Heart Disease'], autopct='%1.1f%%')
plt.title('Heart Disease Distribution')
plt.show()

#Calculate disease percentage
(df['condition'].value_counts(normalize=True) * 100)
```

## Heart Disease Distribution



```
Out[20]: condition
0    53.872054
1    46.127946
Name: proportion, dtype: float64
```

```
In [34]: # 5. Correlation Between Age and Cholesterol

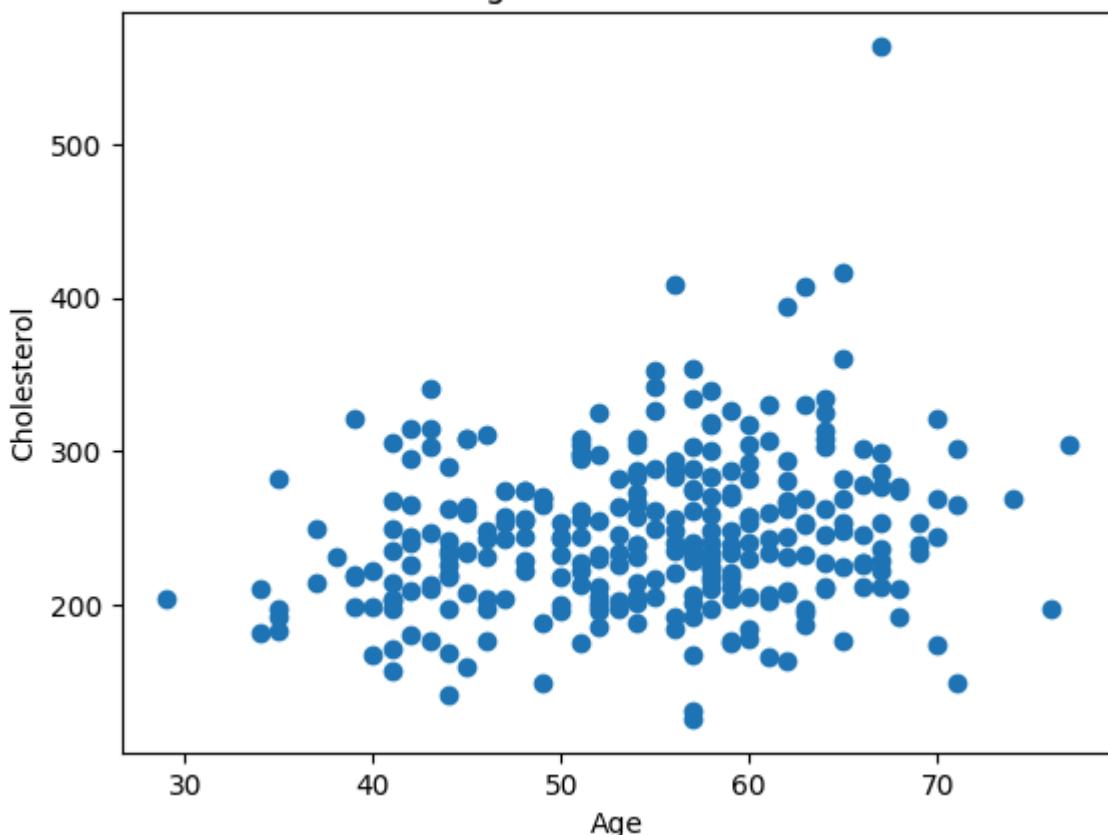
#Calculate correlation using df.corr()
correlation_matrix = df.corr()
print("Correlation Matrix:\n",correlation_matrix)

#scatter plot (Age vs Cholesterol)
df.corr().loc['age', 'chol']
import matplotlib.pyplot as plt
plt.figure()
plt.scatter(df['age'], df['chol'])
plt.xlabel("Age")
plt.ylabel("Cholesterol")
plt.title("Age vs Cholesterol")
plt.show()
```

## Correlation Matrix:

	age	sex	cp	trestbps	chol	fbs	\
age	1.000000	-0.092399	0.110471	0.290476	0.202644	0.132062	
sex	-0.092399	1.000000	0.008908	-0.066340	-0.198089	0.038850	
cp	0.110471	0.008908	1.000000	-0.036980	0.072088	-0.057663	
trestbps	0.290476	-0.066340	-0.036980	1.000000	0.131536	0.180860	
chol	0.202644	-0.198089	0.072088	0.131536	1.000000	0.012708	
fbs	0.132062	0.038850	-0.057663	0.180860	0.012708	1.000000	
restecg	0.149917	0.033897	0.063905	0.149242	0.165046	0.068831	
thalach	-0.394563	-0.060496	-0.339308	-0.049108	-0.000075	-0.007842	
exang	0.096489	0.143581	0.377525	0.066691	0.059339	-0.000893	
oldpeak	0.197123	0.106567	0.203244	0.191243	0.038596	0.008311	
slope	0.159405	0.033345	0.151079	0.121172	-0.009215	0.047819	
ca	0.362210	0.091925	0.235644	0.097954	0.115945	0.152086	
thal	0.120795	0.370556	0.266275	0.130612	0.023441	0.051038	
condition	0.227075	0.278467	0.408945	0.153490	0.080285	0.003167	
							\
	restecg	thalach	exang	oldpeak	slope	ca	\
age	0.149917	-0.394563	0.096489	0.197123	0.159405	0.362210	
sex	0.033897	-0.060496	0.143581	0.106567	0.033345	0.091925	
cp	0.063905	-0.339308	0.377525	0.203244	0.151079	0.235644	
trestbps	0.149242	-0.049108	0.066691	0.191243	0.121172	0.097954	
chol	0.165046	-0.000075	0.059339	0.038596	-0.009215	0.115945	
fbs	0.068831	-0.007842	-0.000893	0.008311	0.047819	0.152086	
restecg	1.000000	-0.072290	0.081874	0.113726	0.135141	0.129021	
thalach	-0.072290	1.000000	-0.384368	-0.347640	-0.389307	-0.268727	
exang	0.081874	-0.384368	1.000000	0.289310	0.250572	0.148232	
oldpeak	0.113726	-0.347640	0.289310	1.000000	0.579037	0.294452	
slope	0.135141	-0.389307	0.250572	0.579037	1.000000	0.109761	
ca	0.129021	-0.268727	0.148232	0.294452	0.109761	1.000000	
thal	0.013612	-0.258386	0.323268	0.336809	0.260096	0.248825	
condition	0.166343	-0.423817	0.421355	0.424052	0.333049	0.463189	
							\
	thal	condition					
age	0.120795	0.227075					
sex	0.370556	0.278467					
cp	0.266275	0.408945					
trestbps	0.130612	0.153490					
chol	0.023441	0.080285					
fbs	0.051038	0.003167					
restecg	0.013612	0.166343					
thalach	-0.258386	-0.423817					
exang	0.323268	0.421355					
oldpeak	0.336809	0.424052					
slope	0.260096	0.333049					
ca	0.248825	0.463189					
thal	1.000000	0.520516					
condition	0.520516	1.000000					

### Age vs Cholesterol



```
In [35]: # 6.Chest Pain Type vs Disease
```

```
#Group by cp and calculate disease rate
cp_disease_rate = df.groupby('cp')['condition'].mean() * 100
print("Disease Rate (%) by Chest Pain Type:\n")
print(cp_disease_rate)

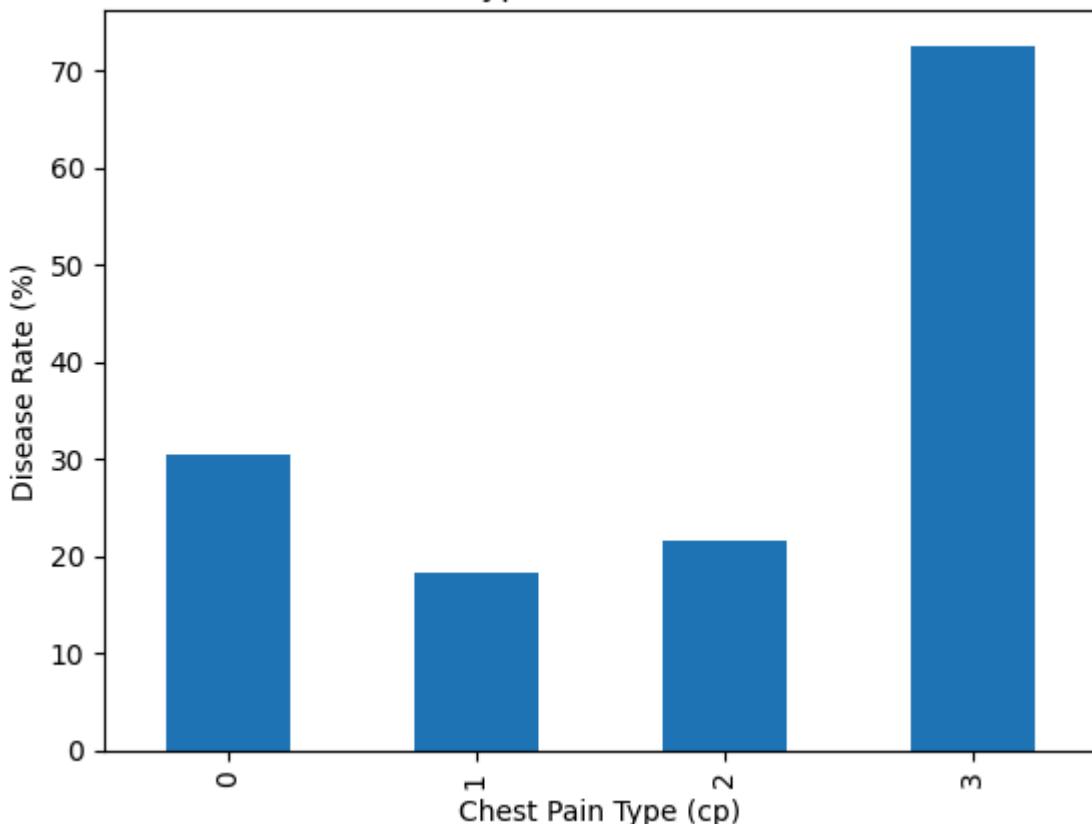
#bar chart
cp_disease_rate.plot(kind='bar')
plt.xlabel("Chest Pain Type (cp)")
plt.ylabel("Disease Rate (%)")
plt.title("Chest Pain Type vs Heart Disease Rate")
plt.show()

#Identify which chest pain type is most risky
most_risky_cp = cp_disease_rate.idxmax()
print("\nMost risky chest pain type:", most_risky_cp)
```

Disease Rate (%) by Chest Pain Type:

```
cp
0    30.434783
1    18.367347
2    21.686747
3    72.535211
Name: condition, dtype: float64
```

### Chest Pain Type vs Heart Disease Rate



Most risky chest pain type: 3

```
In [36]: # 7.Average Cholesterol by Gender

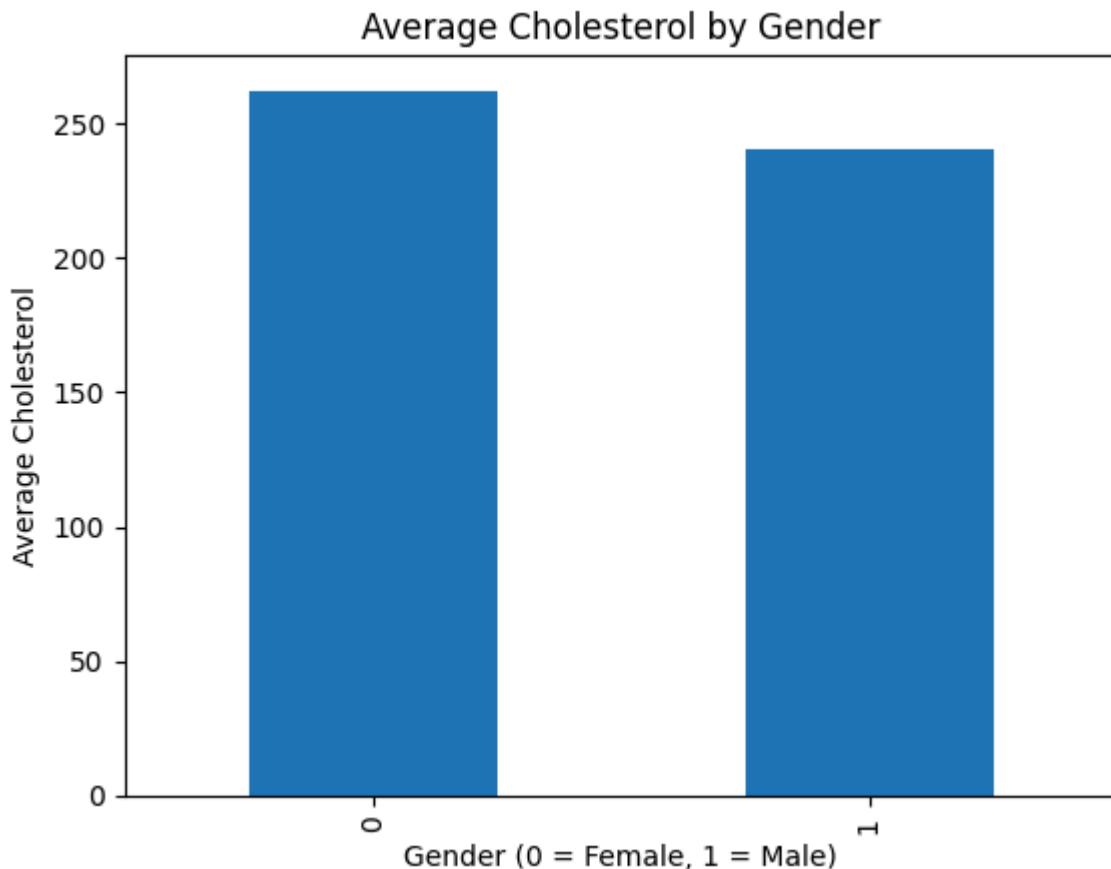
#Group by sex
avg_chol_gender = df.groupby('sex')['chol'].mean()

#Calculate mean cholesterol
print("Average Cholesterol by Gender:\n")
print(avg_chol_gender)

#bar plot
avg_chol_gender.plot(kind='bar')
plt.xlabel("Gender (0 = Female, 1 = Male)")
plt.ylabel("Average Cholesterol")
plt.title("Average Cholesterol by Gender")
plt.show()
```

Average Cholesterol by Gender:

```
sex
0    262.229167
1    240.243781
Name: chol, dtype: float64
```



```
In [38]: # 8. Resting Blood Pressure Analysis

# Average BP
avg_bp = df['trestbps'].mean()
print("Average Resting Blood Pressure:", avg_bp)

# Patients with BP > 140
high_bp = df[df['trestbps'] > 140]
print("Number of patients with BP > 140:", high_bp.shape[0])

# Compare disease presence in high BP group
high_bp_disease_rate = high_bp['condition'].value_counts(normalize=True) * 100
print("\nDisease presence in high BP group (%):")
print(high_bp_disease_rate)
```

Average Resting Blood Pressure: 131.69360269360268  
 Number of patients with BP > 140: 66

Disease presence in high BP group (%):  
 condition  
 1 59.090909  
 0 40.909091  
 Name: proportion, dtype: float64

```
In [39]: # 9. Maximum Heart Rate vs Disease

# Average maximum heart rate for disease and non-disease patients
avg_thalach = df.groupby('condition')['thalach'].mean()

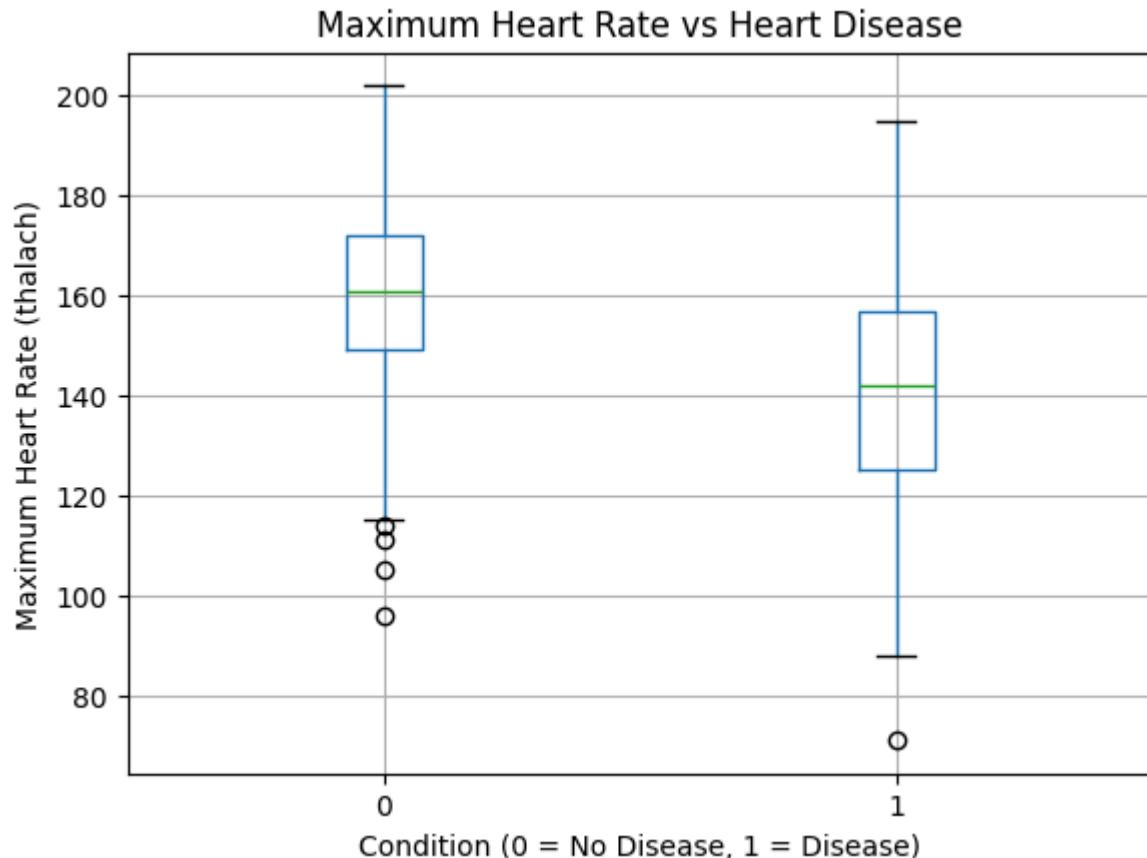
print("Average Maximum Heart Rate (thalach):\n")
print("No Heart Disease (0):", avg_thalach[0])
print("Heart Disease (1):", avg_thalach[1])
```

```
# Boxplot
df.boxplot(column='thalach', by='condition')
plt.xlabel("Condition (0 = No Disease, 1 = Disease)")
plt.ylabel("Maximum Heart Rate (thalach)")
plt.title("Maximum Heart Rate vs Heart Disease")
plt.suptitle("") # remove automatic subtitle
plt.show()
```

Average Maximum Heart Rate (thalach):

No Heart Disease (0): 158.58125

Heart Disease (1): 139.1094890510949



In [41]: `# 10. Exercise Induced Angina Impact`

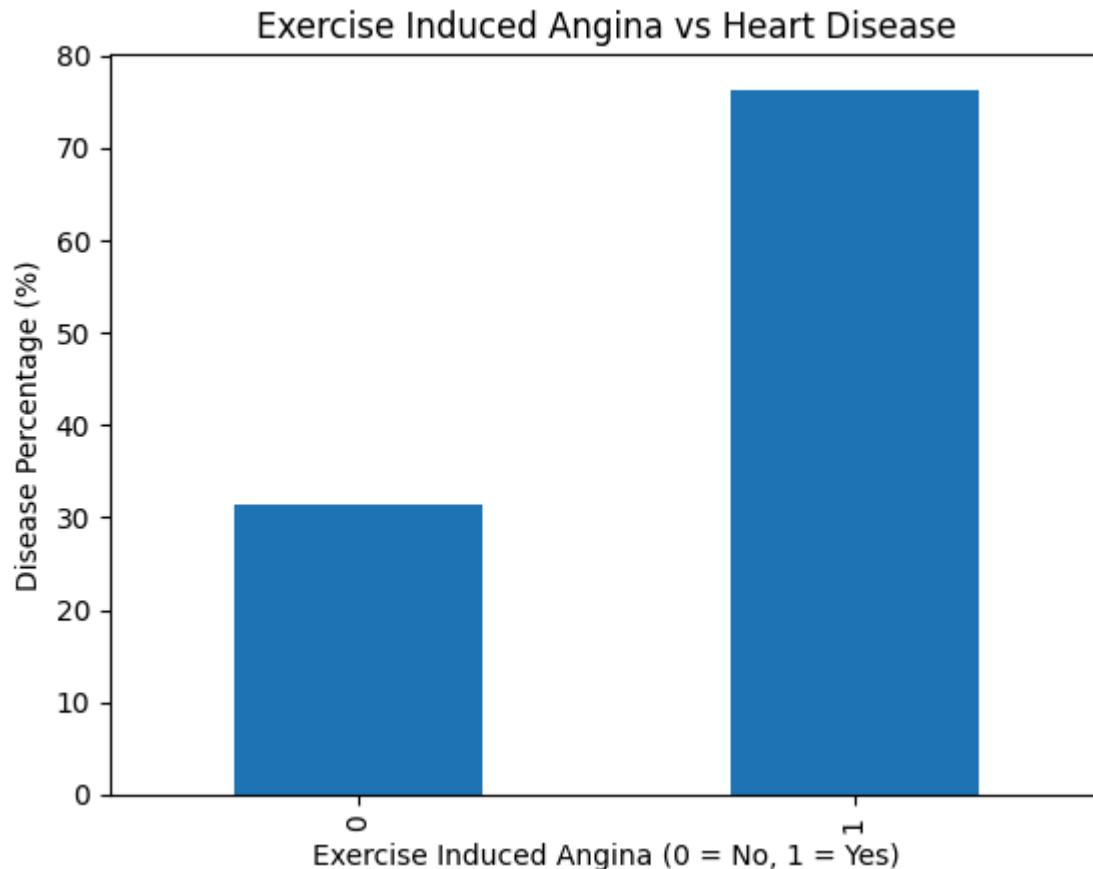
```
#Calculate disease percentage in:
#exang = 0
#exang = 1

exang_disease_rate = df.groupby('exang')['condition'].mean() * 100
print("Disease Percentage by Exercise Induced Angina:\n")
print("exang = 0 (No Angina):", exang_disease_rate[0], "%")
print("exang = 1 (Angina):", exang_disease_rate[1], "%")

#bar chart
exang_disease_rate.plot(kind='bar')
plt.xlabel("Exercise Induced Angina (0 = No, 1 = Yes)")
plt.ylabel("Disease Percentage (%)")
plt.title("Exercise Induced Angina vs Heart Disease")
plt.show()
```

## Disease Percentage by Exercise Induced Angina:

```
exang = 0 (No Angina): 31.5 %
exang = 1 (Angina): 76.28865979381443 %
```



In [42]: # 11. ST Depression (oldpeak) Analysis

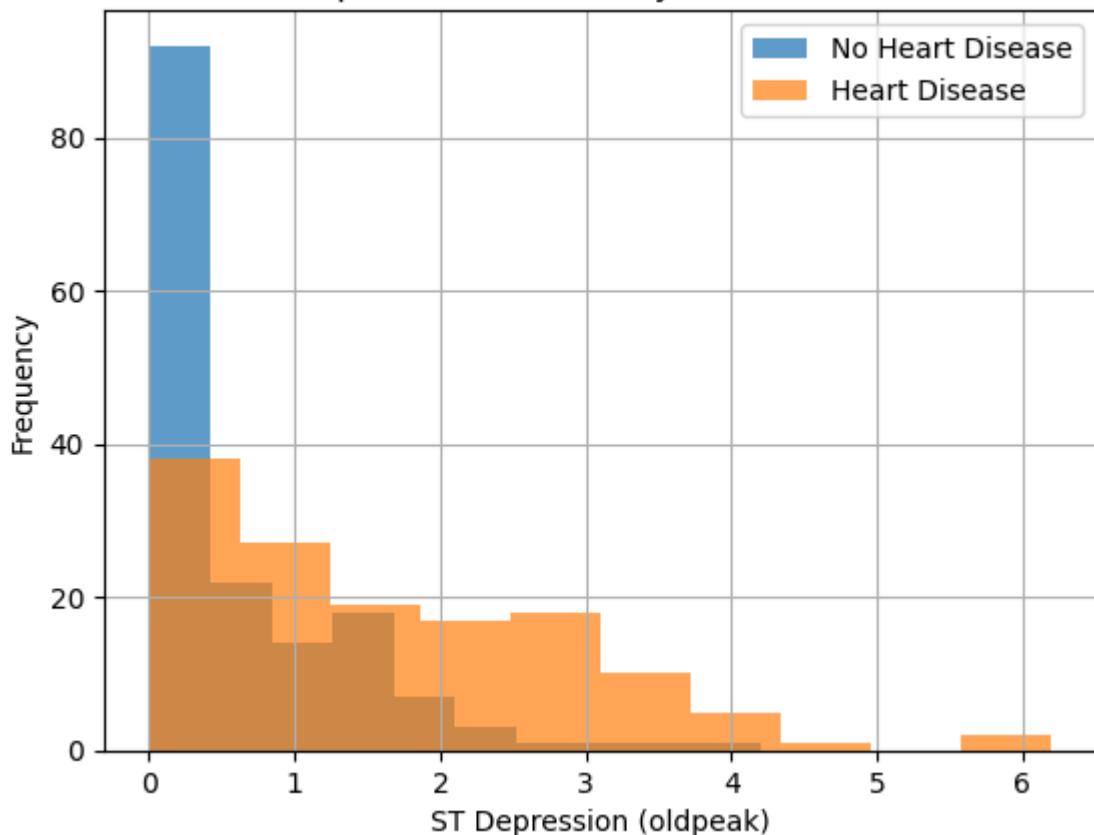
```
#Calculate mean oldpeak by target
mean_oldpeak = df.groupby('condition')['oldpeak'].mean()
print("Mean ST Depression (oldpeak):\n")
print("No Heart Disease (0):", mean_oldpeak[0])
print("Heart Disease (1):", mean_oldpeak[1])

#histogram
df[df['condition'] == 0]['oldpeak'].hist(alpha=0.7)
df[df['condition'] == 1]['oldpeak'].hist(alpha=0.7)
plt.xlabel("ST Depression (oldpeak)")
plt.ylabel("Frequency")
plt.title("Oldpeak Distribution by Heart Disease")
plt.legend(["No Heart Disease", "Heart Disease"])
plt.show()
```

Mean ST Depression (oldpeak):

```
No Heart Disease (0): 0.59875
Heart Disease (1): 1.5890510948905108
```

### Oldpeak Distribution by Heart Disease



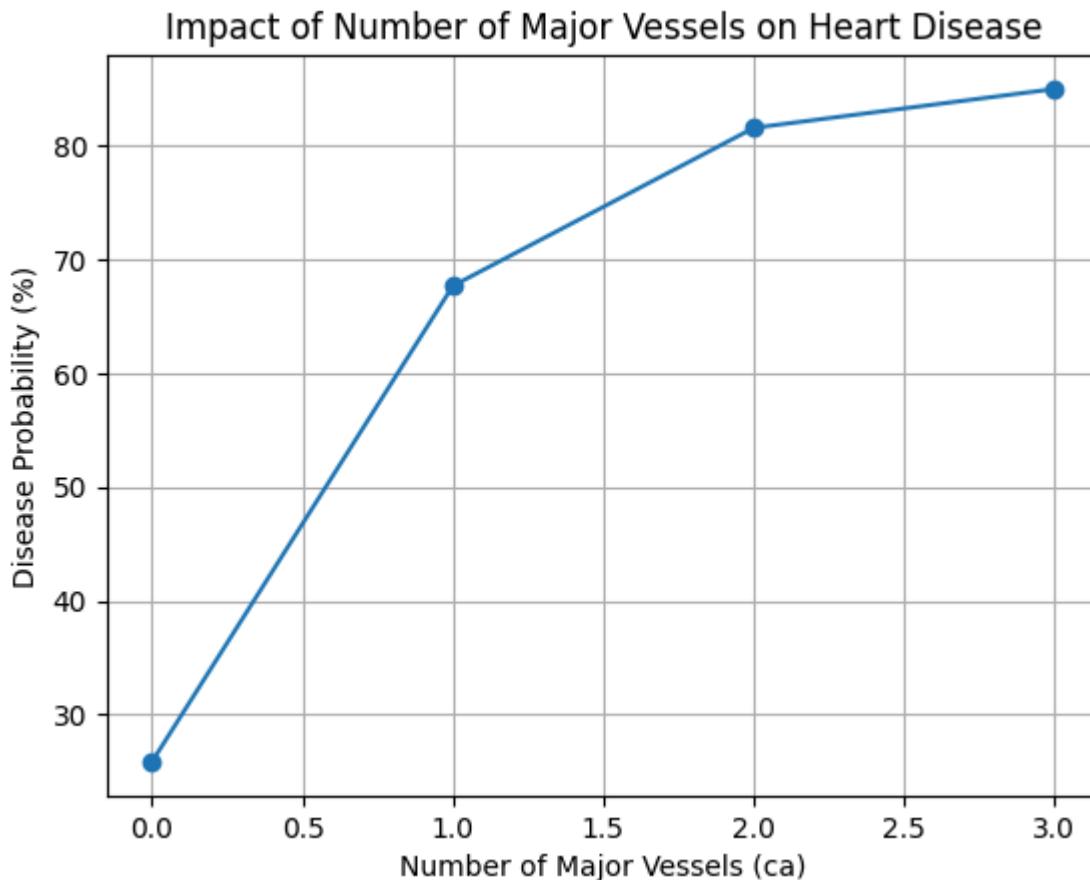
```
In [45]: # 12.Number of Major Vessels (ca) Impact

#Group by ca Calculate disease probability
ca_disease_prob = df.groupby('ca')['condition'].mean() * 100
print("Disease Probability (%) by Number of Major Vessels (ca):\n")
print(ca_disease_prob)

#Plot line chart
ca_disease_prob.plot(marker='o')
plt.xlabel("Number of Major Vessels (ca)")
plt.ylabel("Disease Probability (%)")
plt.title("Impact of Number of Major Vessels on Heart Disease")
plt.grid(True)
plt.show()
```

Disease Probability (%) by Number of Major Vessels (ca):

```
ca
0    25.862069
1    67.692308
2    81.578947
3    85.000000
Name: condition, dtype: float64
```



```
In [47]: # 13.Thalassemia vs Disease

#Cross-tabulate thal and target
thal_ct = pd.crosstab(df['thal'], df['condition'])

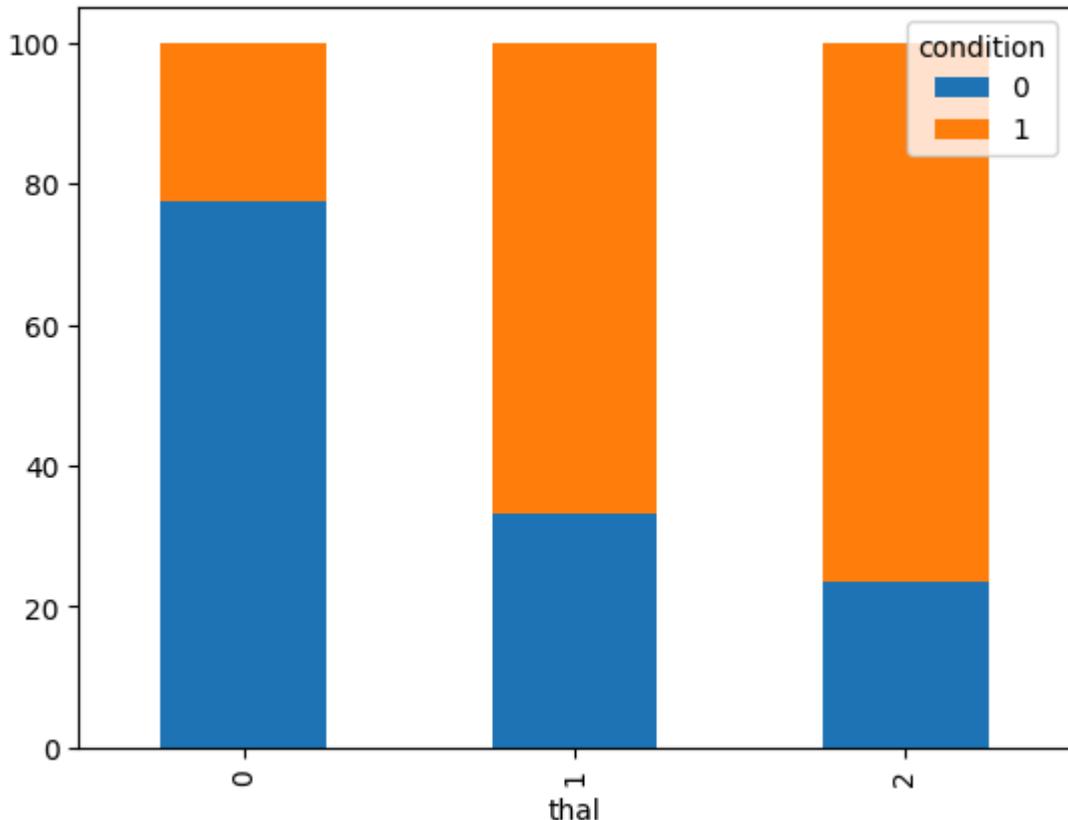
#Convert to percentage
thal_percent = thal_ct.div(thal_ct.sum(axis=1), axis=0) * 100

print("Thalassemia vs Heart Disease (Percentage):\n")
print(thal_percent)

#Plot stacked bar chart
thal_percent.plot(kind='bar', stacked=True)
plt.show()
```

Thalassemia vs Heart Disease (Percentage):

	0	1
thal		
0	77.439024	22.560976
1	33.333333	66.666667
2	23.478261	76.521739



```
In [49]: # 14. Multi-Factor Risk Analysis
```

```
high_risk = df[
    (df['age'] > 50) &
    (df['chol'] > 240) &
    (df['trestbps'] > 140)
]

disease_percentage = high_risk['condition'].mean() * 100
print("Number of high-risk patients:", high_risk.shape[0])
print("Percentage having heart disease:", disease_percentage)
```

Number of high-risk patients: 33  
 Percentage having heart disease: 66.66666666666666

```
In [50]: # 15. Create Risk Score (Custom Analysis)
```

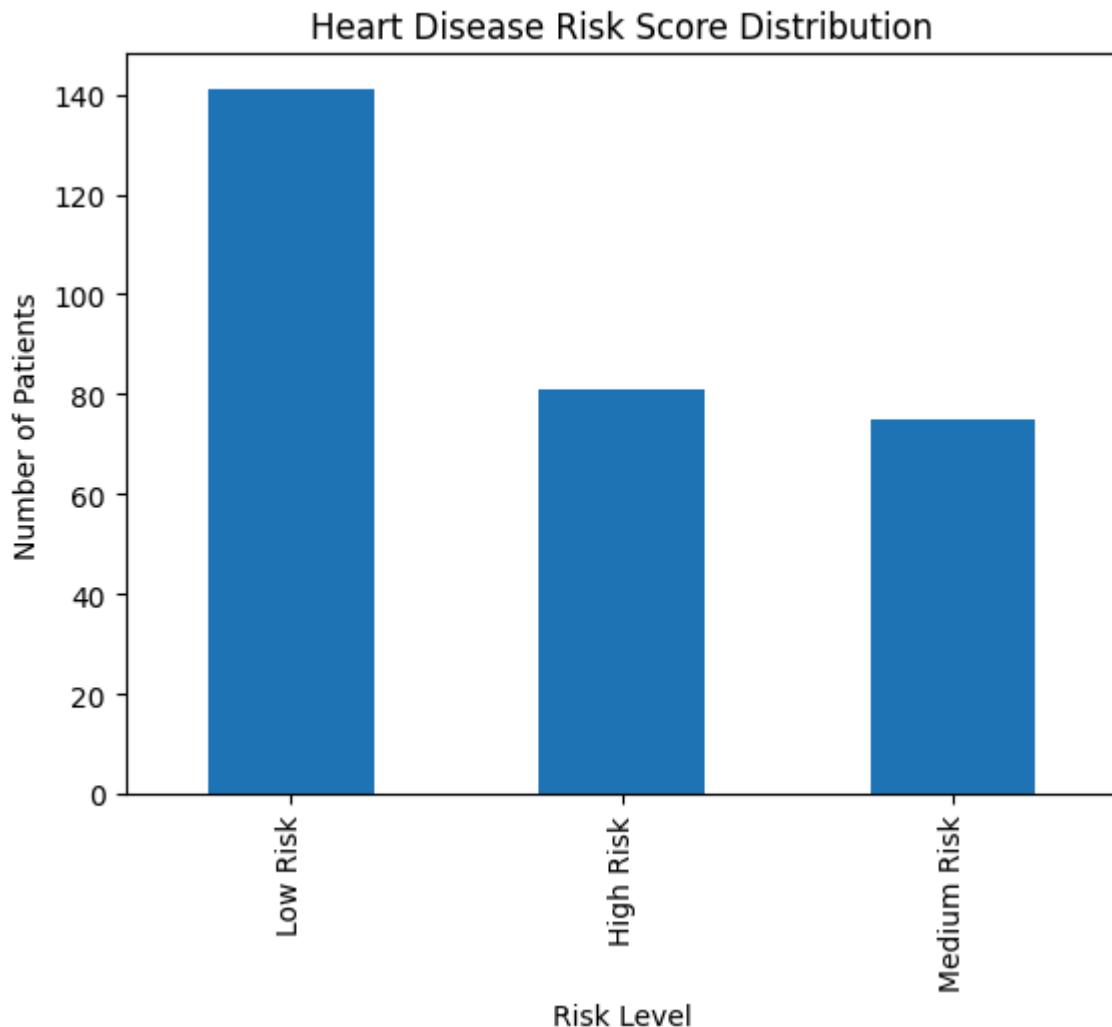
```
df['risk_score'] = (df['chol'] / 200) + (df['trestbps'] / 120) + df['oldpeak']
df['risk_level'] = 'Low Risk'
df.loc[df['risk_score'] >= 3, 'risk_level'] = 'Medium Risk'
df.loc[df['risk_score'] >= 4, 'risk_level'] = 'High Risk'

# Display risk level counts
print("Risk Level Distribution:\n")
print(df['risk_level'].value_counts())

# Visualize distribution
df['risk_level'].value_counts().plot(kind='bar')
plt.xlabel("Risk Level")
plt.ylabel("Number of Patients")
plt.title("Heart Disease Risk Score Distribution")
plt.show()
```

## Risk Level Distribution:

```
risk_level
Low Risk      141
High Risk     81
Medium Risk   75
Name: count, dtype: int64
```



In [ ]: 1. Does cholesterol strongly impact heart disease?

No **not** strongly

Cholesterol shows only a weak to moderate association **with** heart disease

Many patients **with** normal cholesterol still have heart disease **and** vice versa

Cholesterol alone **is not** a strong predictor without other risk factors

Conclusion: Cholesterol contributes to risk but does **not** strongly impact heart disease

2. Is the male population more vulnerable?

Yes

Male patients show a higher prevalence of heart disease compared to females

Gender-based analysis indicates males are more prone to heart disease

Conclusion: The male population **is** more vulnerable to heart disease

3. Does exercise-induced angina significantly increase risk?

Yes significantly

Patients **with** exercise-induced angina (`exang = 1`) have a much higher disease percentage

This feature shows a clear **and** strong separation between diseased **and** non-diseased

Conclusion: Exercise-induced angina **is** a strong indicator of heart disease risk

4.Which feature has the strongest correlation **with** disease

Number of major vessels (ca)

ca shows the strongest positive correlation **with** heart disease

Disease probability increases steadily **as** the number of major vessels increases

Other strong features include

Chest pain type (cp)

ST depression (oldpeak)

Exercise-induced angina (exang)