

```
In [1]: # Importing all necessary libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.decomposition import PCA
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
from sklearn.preprocessing import StandardScaler
from sklearn.feature_selection import SelectKBest, f_classif
```

```
In [2]: # Loading the dataset
df = pd.read_excel("C:/Users/Ansh/Desktop/Main Flow/heart.xlsx")

# Convert inf values to NaN
df.replace([np.inf, -np.inf], np.nan, inplace=True)
```

```
In [3]: # Display the first few rows of the dataset
print("Dataset Overview:")
print(df.head())
```

Dataset Overview:

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

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

```
In [4]: # Display column names
print("Column Names:")
print(df.columns.values)
```

Column Names:

```
['age' 'sex' 'cp' 'trestbps' 'chol' 'fbs' 'restecg' 'thalach' 'exang'
 'oldpeak' 'slope' 'ca' 'thal' 'target']
```

```
In [5]: # Checking for null values
print("Null Values in Each Column:")
print(df.isna().sum())
```

Null Values in Each Column:

```
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
```

```
In [6]: # Data type and basic info
print("Dataset Info:")
df.info()
```

Dataset 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
```

```
In [7]: # Descriptive statistics of the dataset
print("Statistical Summary:")
print(df.describe())
```

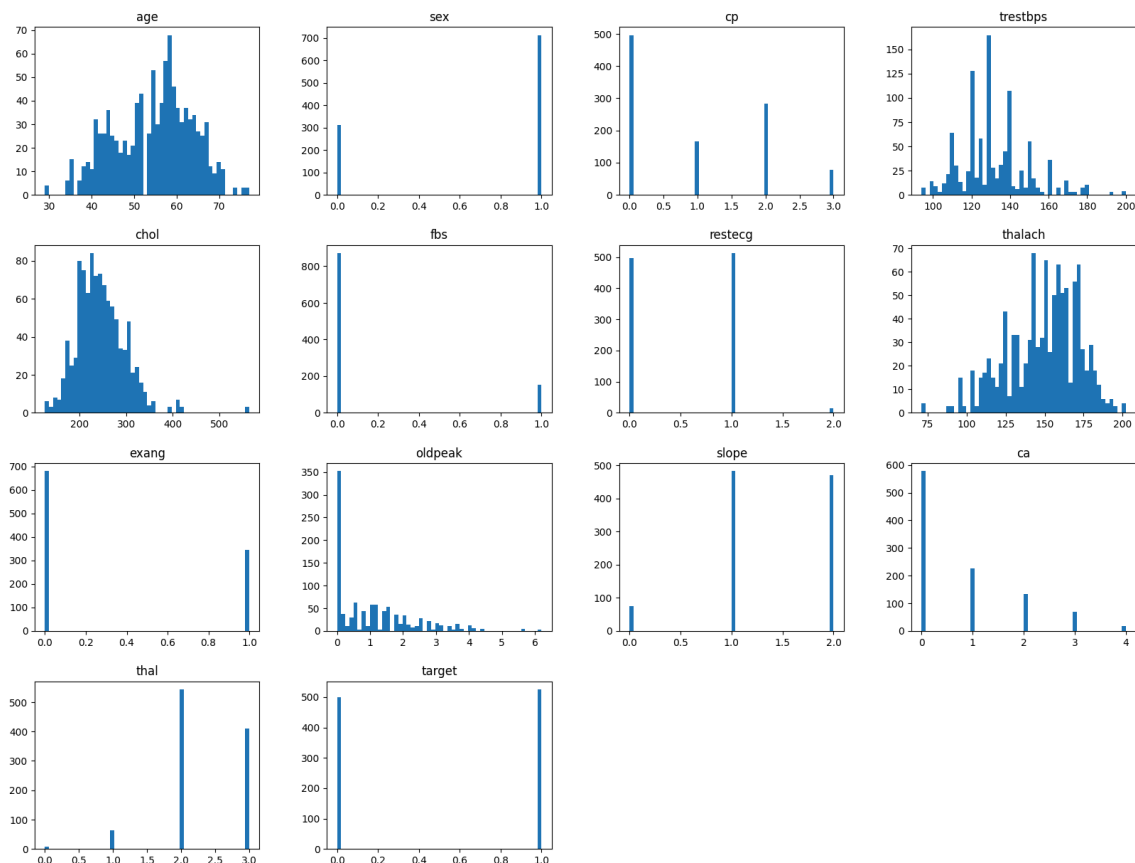
Statistical Summary:

	age	sex	cp	trestbps	chol \
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	54.434146	0.695610	0.942439	131.611707	246.000000
std	9.072290	0.460373	1.029641	17.516718	51.59251
min	29.000000	0.000000	0.000000	94.000000	126.000000
25%	48.000000	0.000000	0.000000	120.000000	211.000000
50%	56.000000	1.000000	1.000000	130.000000	240.000000
75%	61.000000	1.000000	2.000000	140.000000	275.000000
max	77.000000	1.000000	3.000000	200.000000	564.000000

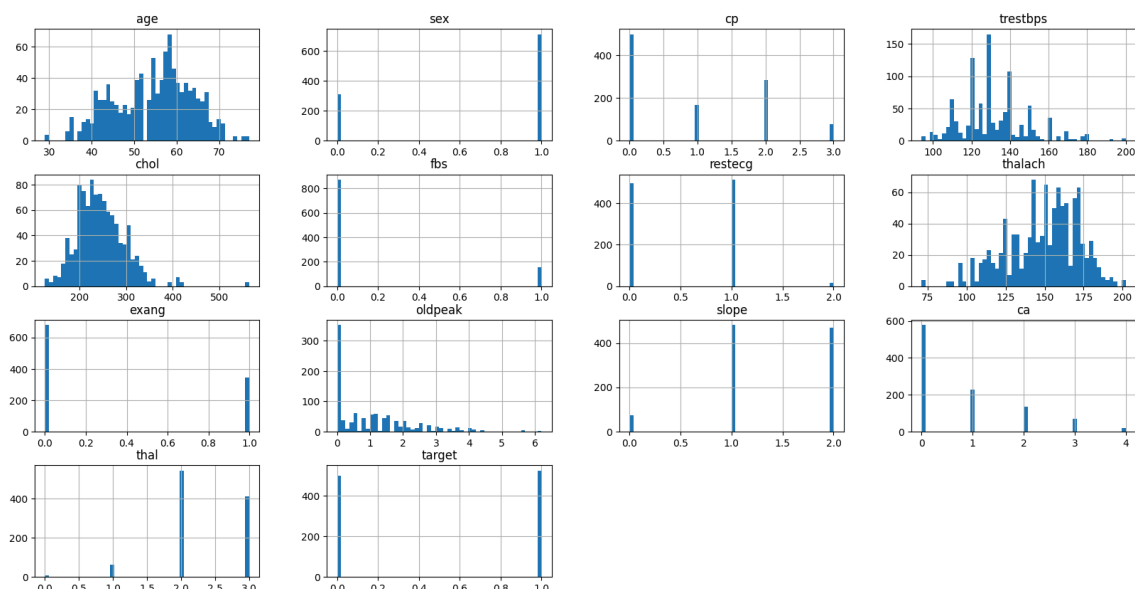
	fbs	restecg	thalach	exang	oldpeak \
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	0.149268	0.529756	149.114146	0.336585	1.071512
std	0.356527	0.527878	23.005724	0.472772	1.175053
min	0.000000	0.000000	71.000000	0.000000	0.000000
25%	0.000000	0.000000	132.000000	0.000000	0.000000
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

```
In [8]: # Visualizing distributions of numerical features
df.hist(bins=50, grid=False, figsize=(20,15))
plt.show()
```



```
In [9]: # Visualizing distributions of numerical features
df.hist(bins=50, grid=True, figsize=(20,10))
plt.show()
```



```
In [10]: # Exploring specific questions:
questions = [
    "1. How many people have heart disease and how many don't?",
    "2. Which sex has the most heart disease?",
    "3. Which sex has which type of chest pain most?",
    "4. Which chest pain type is most prone to heart disease?",
    "5. Distribution of age and its relation to heart disease",
    "6. Maximum heart rate and its impact on heart disease",
    "7. Relation between resting blood pressure and heart disease"
]
```

```
In [11]: # 1. How many people have heart disease and how many don't?
print(questions[0])
print(df['target'].value_counts())

# Plotting the count of heart disease cases
df['target'].value_counts().plot(kind='bar', color=["orchid", "salmon"])
plt.title("Heart Disease Distribution")
plt.xlabel("0: No Heart Disease, 1: Heart Disease")
plt.ylabel("Count")
plt.show()
```

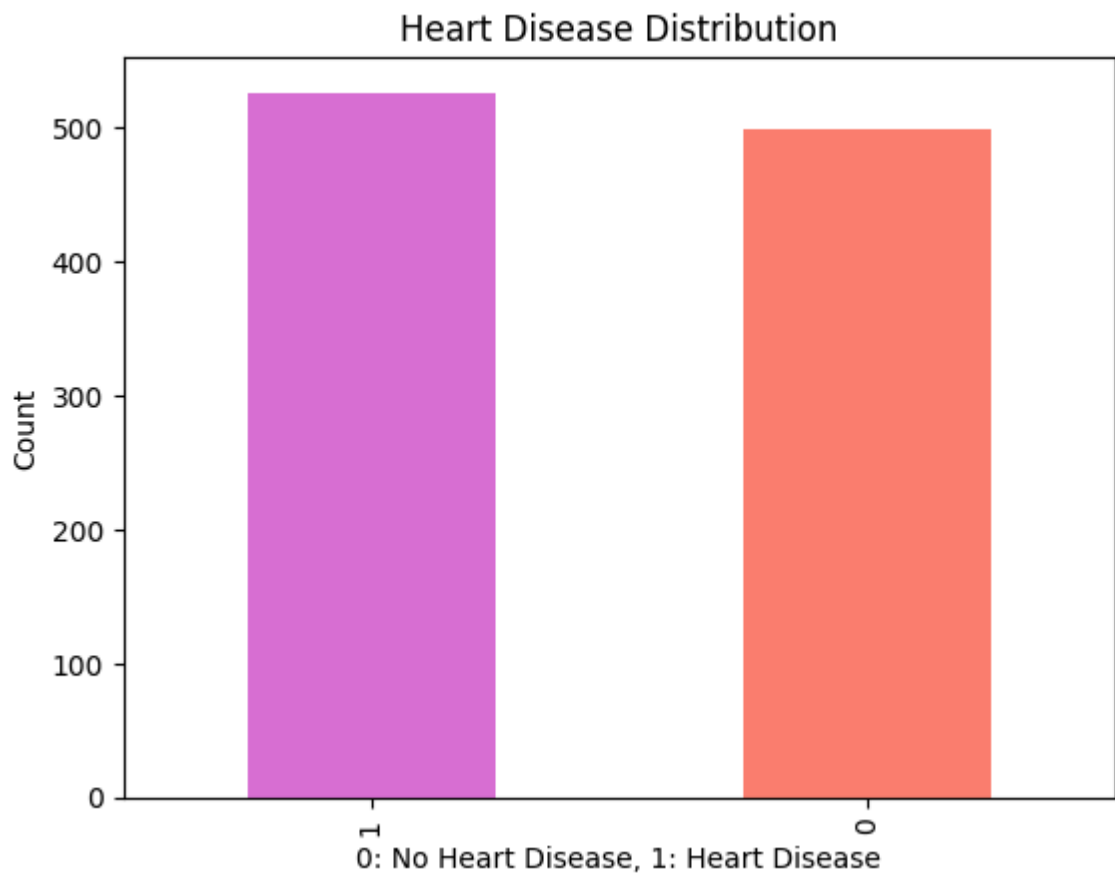
1. How many people have heart disease and how many don't?

target

1 526

0 499

Name: count, dtype: int64

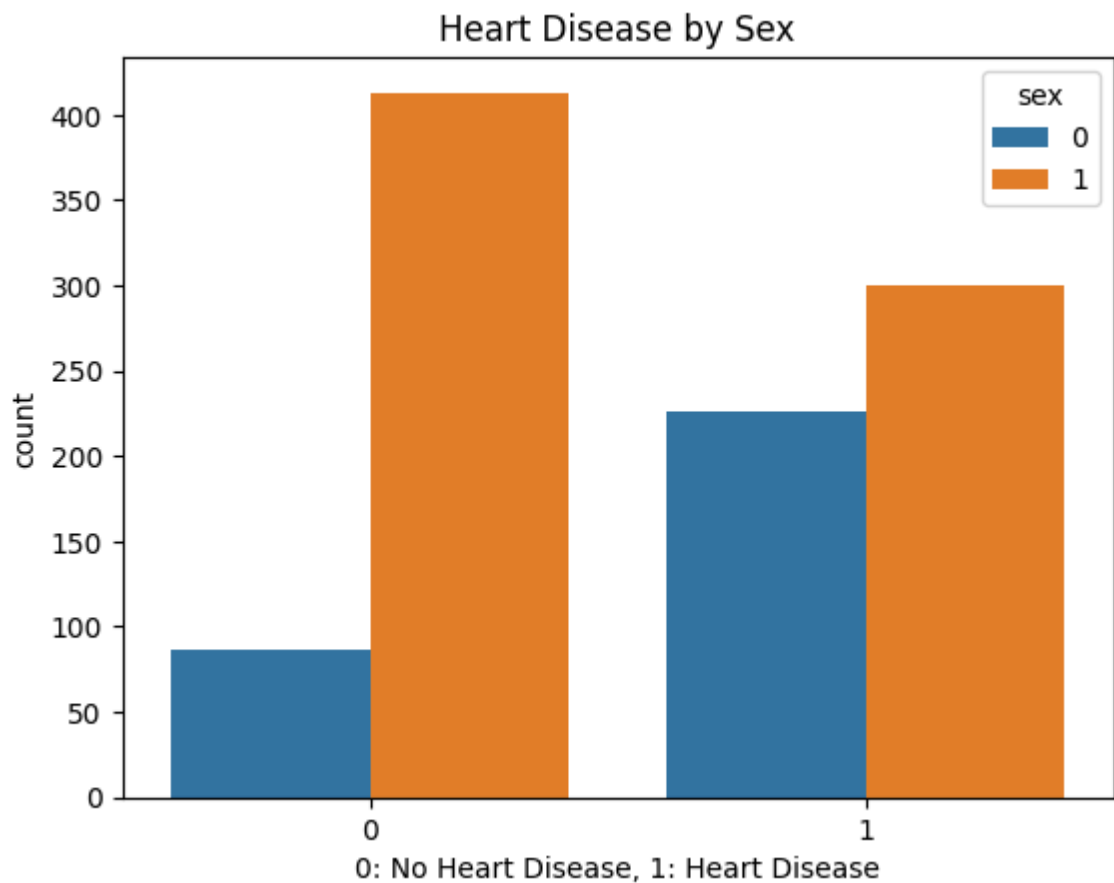


```
In [12]: # 2. Which sex has the most heart disease?
print(questions[1])
print(pd.crosstab(df.target, df.sex))

sns.countplot(x='target', data=df, hue='sex')
plt.title("Heart Disease by Sex")
plt.xlabel("0: No Heart Disease, 1: Heart Disease")
plt.show()
```

2. Which sex has the most heart disease?

sex	0	1
target		
0	86	413
1	226	300



```
In [13]: # 3. Which sex has which type of chest pain most?
print(questions[2])
print(df['cp'].value_counts())

sns.countplot(x='cp', data=df, hue='sex')
plt.title("Chest Pain Type by Sex")
plt.show()
```

3. Which sex has which type of chest pain most?

cp

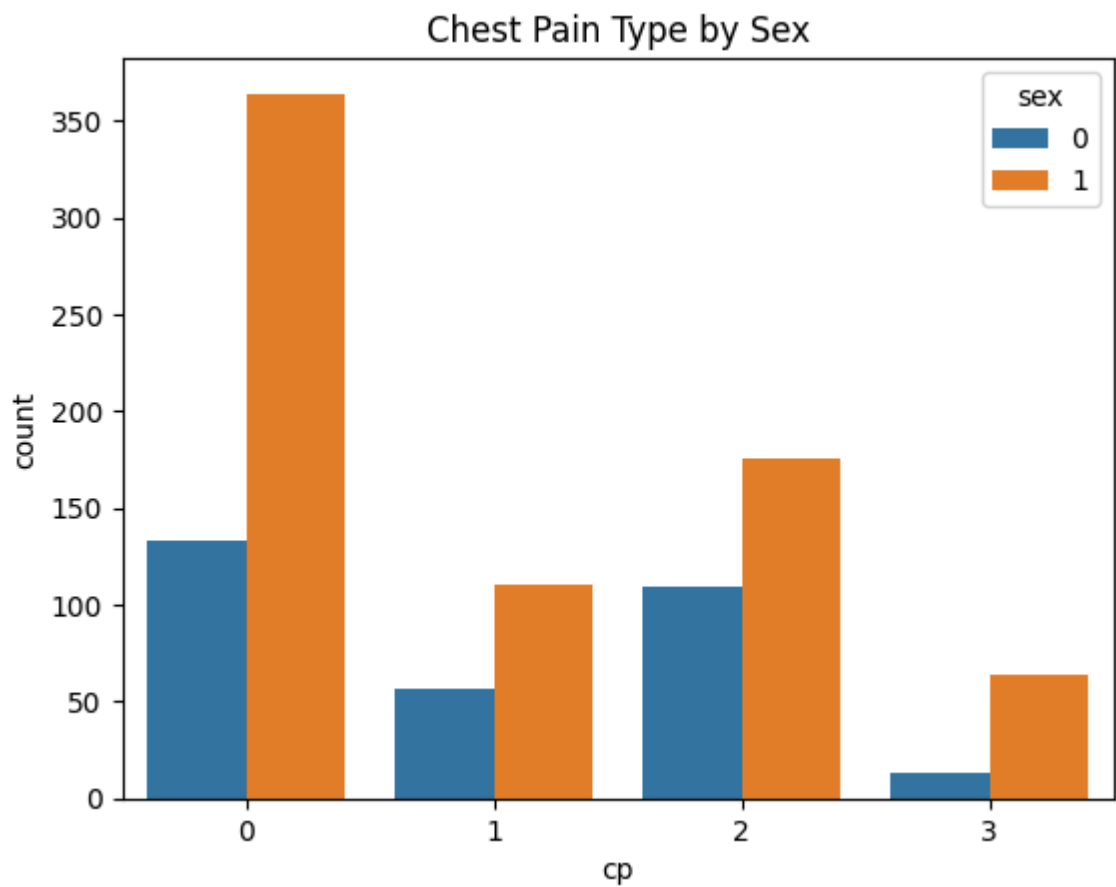
0 497

2 284

1 167

3 77

Name: count, dtype: int64

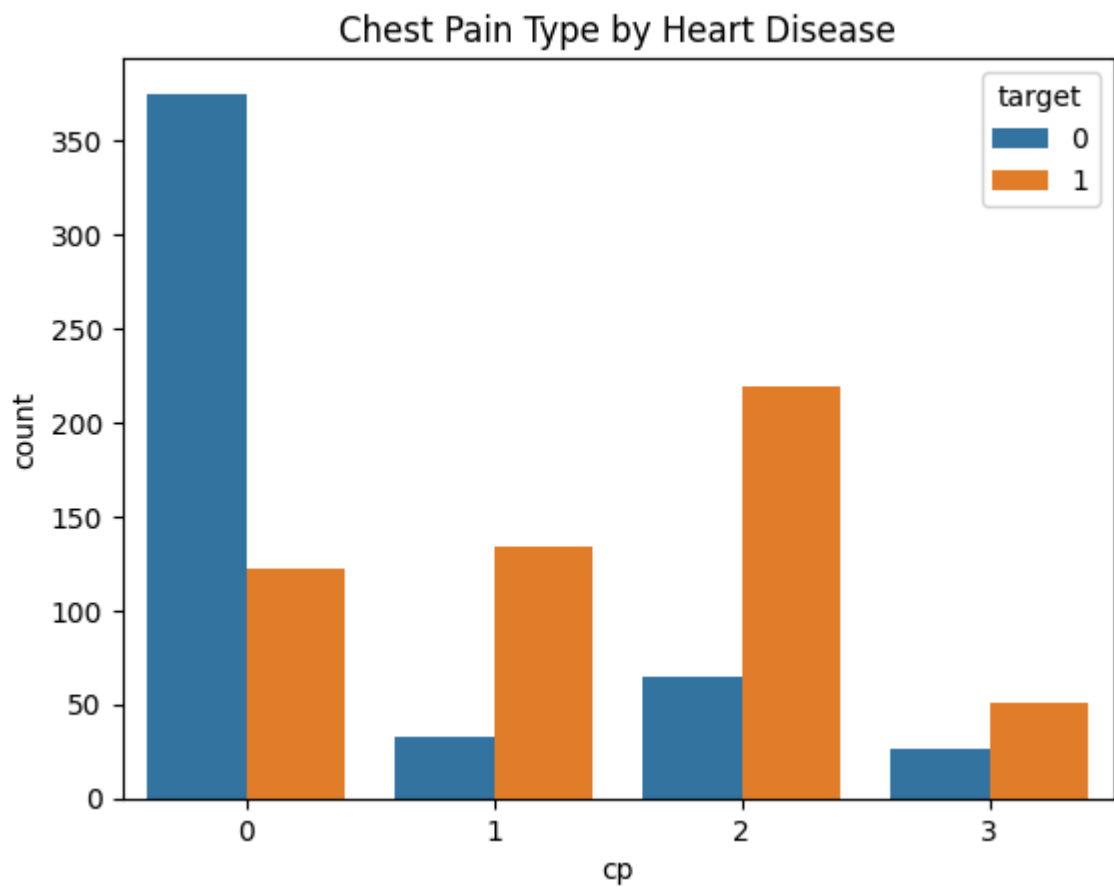


```
In [14]: # 4. Which chest pain type is most prone to heart disease?
print(questions[3])
print(pd.crosstab(df.cp, df.target))

sns.countplot(x='cp', data=df, hue='target')
plt.title("Chest Pain Type by Heart Disease")
plt.show()
```

4. Which chest pain type is most prone to heart disease?

target	0	1
cp		
0	375	122
1	33	134
2	65	219
3	26	51




```
In [15]: # 5. Distribution of age and its relation to heart disease
sns.displot(x='age', data=df, kde=True, bins=30, hue='target')
plt.title("Age Distribution with Heart Disease")
plt.show()
```

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: Future Warning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.

with pd.option_context('mode.use_inf_as_na', True):

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)

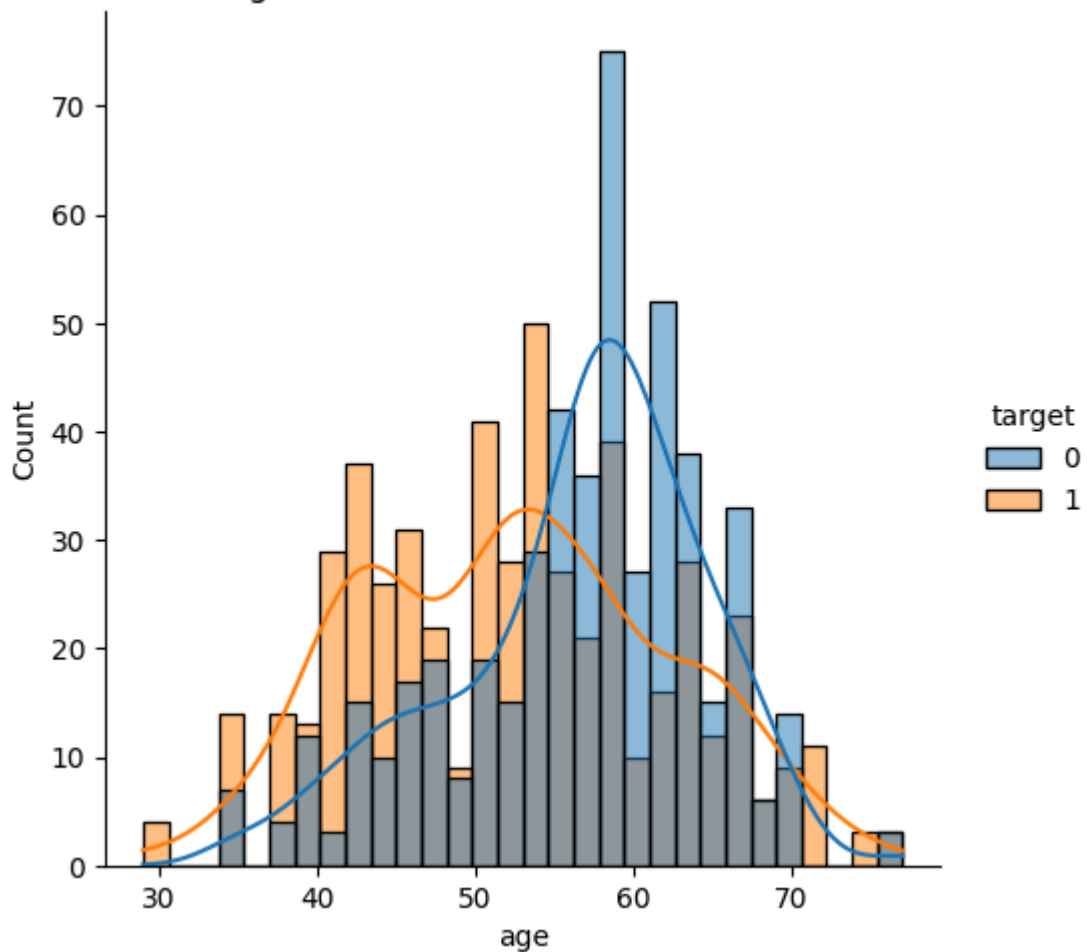
C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)

Age Distribution with Heart Disease



```
In [16]: # 6. Maximum heart rate and its impact on heart disease
sns.displot(x='thalach', data=df, kde=True, bins=30, color='chocolate', hue
plt.title("Maximum Heart Rate and Heart Disease")
plt.show()
```

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: Future Warning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.

with pd.option_context('mode.use_inf_as_na', True):

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

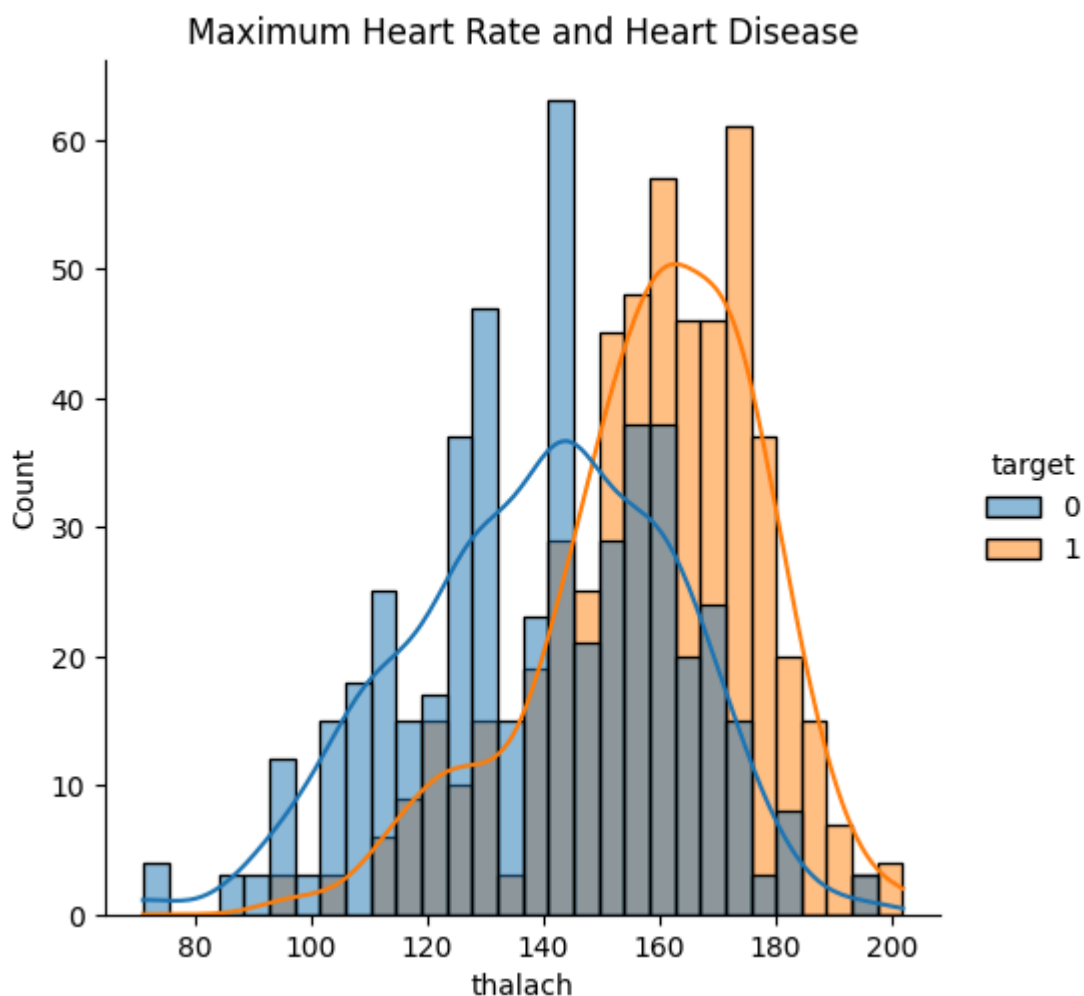
data_subset = grouped_data.get_group(pd_key)

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)



```
In [17]: # 7. Resting blood pressure and its relation to heart disease
sns.displot(x='trestbps', data=df, kde=True, bins=30, color='skyblue', hue=
plt.title("Resting Blood Pressure and Heart Disease")
plt.show()
```

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: Future Warning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.

with pd.option_context('mode.use_inf_as_na', True):

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

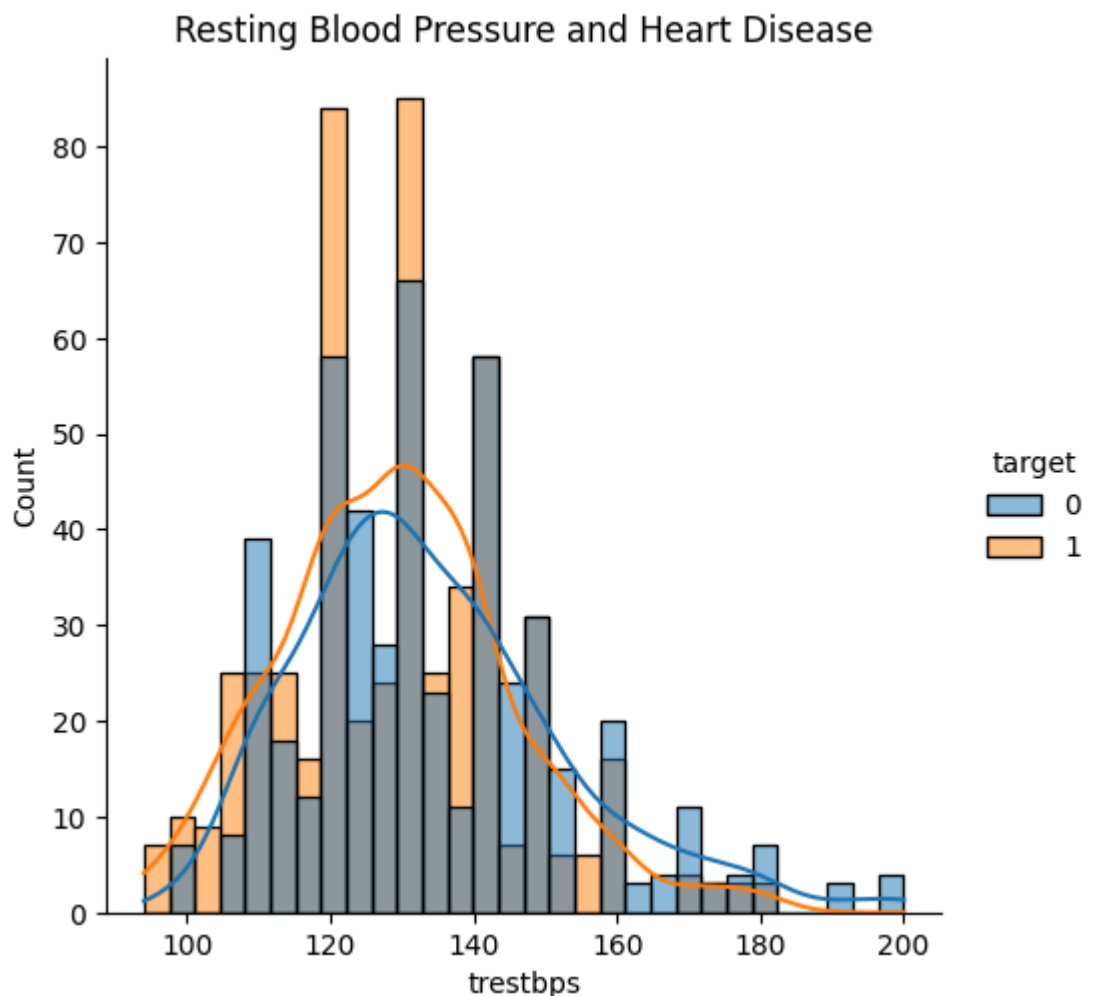
data_subset = grouped_data.get_group(pd_key)

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)

C:\Users\Ansh\anaconda3\Lib\site-packages\seaborn_oldcore.py:1075: Future Warning: When grouping with a length-1 list-like, you will need to pass a length-1 tuple to get_group in a future version of pandas. Pass `(name,)` instead of `name` to silence this warning.

data_subset = grouped_data.get_group(pd_key)



```
In [18]: # Feature Engineering: Creating new features (e.g., BMI, cholesterol-blood
df['chol_bp_ratio'] = df['chol'] / df['trestbps']
df['age_thalach_ratio'] = df['age'] / df['thalach']
```

```
In [19]: # Feature Selection: Using PCA and Feature Importance
# Splitting the dataset into features and target variable
X = df.drop('target', axis=1)
y = df['target']

# Standardizing the features
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
```

```
In [20]: # Applying PCA
pca = PCA(n_components=5) # Reducing to 5 principal components for simplicity
X_pca = pca.fit_transform(X_scaled)
```

```
In [21]: # Using SelectKBest for feature selection
selector = SelectKBest(score_func=f_classif, k=10)
X_new = selector.fit_transform(X, y)
```

```
In [22]: # Using RandomForestClassifier for feature importance
model = RandomForestClassifier(random_state=42)
model.fit(X_new, y)
importances = model.feature_importances_
```

```
In [23]: # Displaying feature importances
feature_names = X.columns[selector.get_support()]
importance_df = pd.DataFrame({
    'Feature': feature_names,
    'Importance': importances
}).sort_values(by='Importance', ascending=False)

print("\nTop Features based on Random Forest Importance:")
print(importance_df)
```

Top Features based on Random Forest Importance:

	Feature	Importance
2	cp	0.156111
7	ca	0.131463
5	oldpeak	0.123308
3	thalach	0.122212
8	thal	0.119441
9	age_thalach_ratio	0.114638
0	age	0.094604
4	exang	0.059173
6	slope	0.042675
1	sex	0.036375

```
In [24]: # Splitting data for model training and testing
X_train, X_test, y_train, y_test = train_test_split(X_new, y, test_size=0.3)

# Training the model
model.fit(X_train, y_train)
```

Out[24]:

▼ RandomForestClassifier ⓘ ?

RandomForestClassifier(random_state=42)

(<https://scikit-learn.org/1.5/modules/generated/sklearn.ensem>)

```
In [25]: # Making predictions and evaluating the model
y_pred = model.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)

print(f"\nModel Accuracy: {accuracy:.2f}")
```

Model Accuracy: 0.99

In []: