



**Machine Learning Lab/Even Sem 2023-23/Experiment 1b**

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**Title of Experiment :**

Introduction to scikit learn, matplotlib, seaborn library

**Objective of Experiment :**

To introduce platforms such as Anaconda, COLAB suitable to Machine learning.

**Outcome of Experiment :**

Implement various Machine learning models

**Problem Statement :**

Introduction to scikit learn, matplotlib, seaborn library

**Description / Theory :**

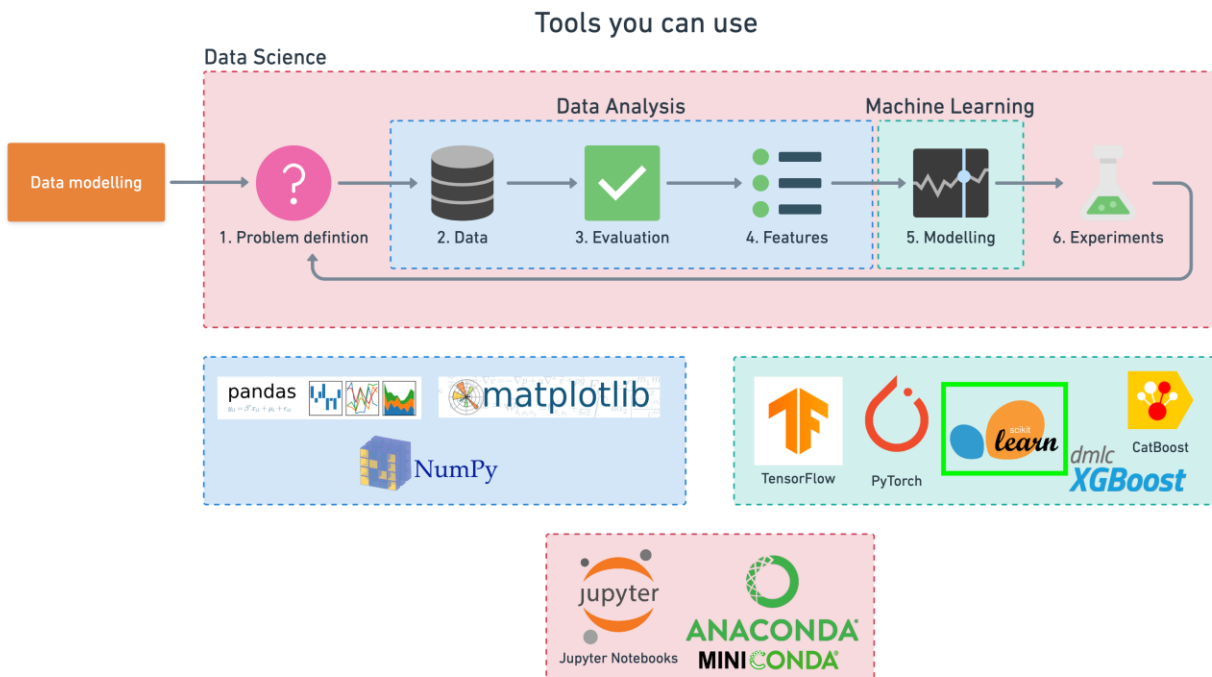


### Machine Learning Lab/Even Sem 2023-23/Experiment 1b

#### What is Scikit-Learn (sklearn)?

Scikit-Learn, also referred to as sklearn, is an open-source Python machine learning library.

It's built on top on NumPy (Python library for numerical computing) and Matplotlib (Python library for data visualization).



#### Why Scikit-Learn?

Although the fields of data science and machine learning are vast, the main goal is finding patterns within data and then using those patterns to make predictions.

And there are certain categories which a majority of problems fall into.

If you're trying to create a machine learning model to predict whether an email is spam and or not spam, you're working on a classification problem (whether something is one thing or another).

If you're trying to create a machine learning model to predict the price of houses given their characteristics, you're working on a regression problem (predicting a number).

If you're trying to get a machine learning algorithm to group together similar samples (that you don't necessarily know which should go together), you're working on a clustering problem.



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Once you know what kind of problem you're working on, there are also similar steps you'll take for each. Steps like splitting the data into different sets, one for your machine learning algorithms to learn on (the training set) and another to test them on (the testing set).

Choosing a machine learning model and then evaluating whether or not your model has learned anything.

Scikit-Learn offers Python implementations for doing all of these kinds of tasks (from preparing data to modelling data). Saving you from having to build them from scratch.

What is matplotlib?

Matplotlib is a visualization library for Python.

As in, if you want to display something in a chart or graph, matplotlib can help you do that programmatically.

Many of the graphics you'll see in machine learning research papers or presentations are made with matplotlib.

Why matplotlib?

Matplotlib is part of the standard Python data stack (pandas, NumPy, matplotlib, Jupyter).

It has terrific integration with many other Python libraries.

pandas uses matplotlib as a backend to help visualize data in DataFrames.

What is seaborn?

Seaborn is a library for making statistical graphics in Python. It builds on top of matplotlib and integrates closely with pandas data structures.

Seaborn helps you explore and understand your data. Its plotting functions operate on dataframes and arrays containing whole datasets and internally perform the necessary semantic mapping and statistical aggregation to produce informative plots. Its dataset-oriented, declarative API lets you focus on what the different elements of your plots mean, rather than on the details of how to draw them.

## 0.1 Introduction to Scikit Learn

```
[1]: # Importing required libraries

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import sklearn
import seaborn as sns
```

```
[20]: # Reading data

heart_disease = pd.read_csv('../data/cleaned/heart.csv')
heart_disease.head()
```

```
[20]:
```

	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

```
[21]: # Create X (all the feature columns)
X = heart_disease.drop("target", axis=1)

# Create y (the target column)
y = heart_disease["target"]

# Check the head of the features DataFrame
X.head()
```

```
[21]:
```

	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
0	2	3
1	0	3
2	0	3
3	1	3
4	3	2

```
[22]: # Check the head and the value counts of the labels
y.head(), y.value_counts()
```

```
[22]: (0      0
      1      0
      2      0
      3      0
      4      0
      Name: target, dtype: int64,
      target
      1      526
      0      499
      Name: count, dtype: int64)
```

```
[23]: # Split the data into training and test sets
      from sklearn.model_selection import train_test_split

      X_train, X_test, y_train, y_test = train_test_split(X,
                                                         y,
                                                         test_size=0.25) # by default
      ↪train_test_split uses 25% of the data for the test set

      X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

```
[23]: ((768, 13), (257, 13), (768,), (257,))
```

```
[24]: from sklearn.ensemble import RandomForestClassifier

      clf = RandomForestClassifier()
```

```
[25]: clf.get_params()
```

```
[25]: {'bootstrap': True,
      'ccp_alpha': 0.0,
      'class_weight': None,
      'criterion': 'gini',
      'max_depth': None,
      'max_features': 'sqrt',
      'max_leaf_nodes': None,
      'max_samples': None,
      'min_impurity_decrease': 0.0,
      'min_samples_leaf': 1,
      'min_samples_split': 2,
      'min_weight_fraction_leaf': 0.0,
      'n_estimators': 100,
      'n_jobs': None,
      'oob_score': False,
      'random_state': None,
      'verbose': 0,
      'warm_start': False}
```

```
[26]: clf.fit(X=X_train, y=y_train)
```

```
[26]: RandomForestClassifier()
```

```
[27]: X_test.head()
```

```
[27]:
```

	age	sex	cp	trestbps	chol	fb	restecg	thalach	exang	oldpeak	\
81	49	1	2	118	149	0	0	126	0	0.8	
706	57	1	2	128	229	0	0	150	0	0.4	
792	68	1	0	144	193	1	1	141	0	3.4	
113	57	1	0	110	335	0	1	143	1	3.0	
643	65	1	0	120	177	0	1	140	0	0.4	

	slope	ca	thal
81	2	3	2
706	1	1	3
792	1	2	3
113	1	1	3
643	2	0	3

```
[28]: y_preds = clf.predict(X=X_test)
y_preds
```

```
[28]: array([0, 0, 0, 0, 1, 0, 1, 0, 1, 1, 1, 1, 0, 1, 1, 0, 0, 1, 1, 0, 1, 1,
        1, 0, 1, 0, 1, 0, 1, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 1, 0, 0, 1,
        0, 1, 1, 1, 1, 0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 1, 1, 0, 0, 0, 0, 0,
        1, 1, 1, 0, 1, 0, 0, 1, 0, 1, 0, 1, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1,
        0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 1, 0, 1, 1, 1, 0, 0, 1, 1, 1, 0,
        0, 1, 1, 1, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 1, 0, 1, 0, 0, 0, 1, 0,
        1, 0, 1, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 1, 0, 0, 1, 1, 1, 0,
        0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 1, 0, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1,
        0, 0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 1, 1, 0, 0, 0, 1, 0, 0,
        1, 1, 0, 1, 1, 1, 0, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0, 1, 1,
        0, 1, 1, 1, 1, 1, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 0, 1, 1, 0, 0, 1,
        1, 1, 0, 0, 0, 1, 1, 0, 1, 0, 1, 1, 0, 0, 1], dtype=int64)
```

```
[29]: from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

# Create a classification report
print(classification_report(y_test, y_preds))
```

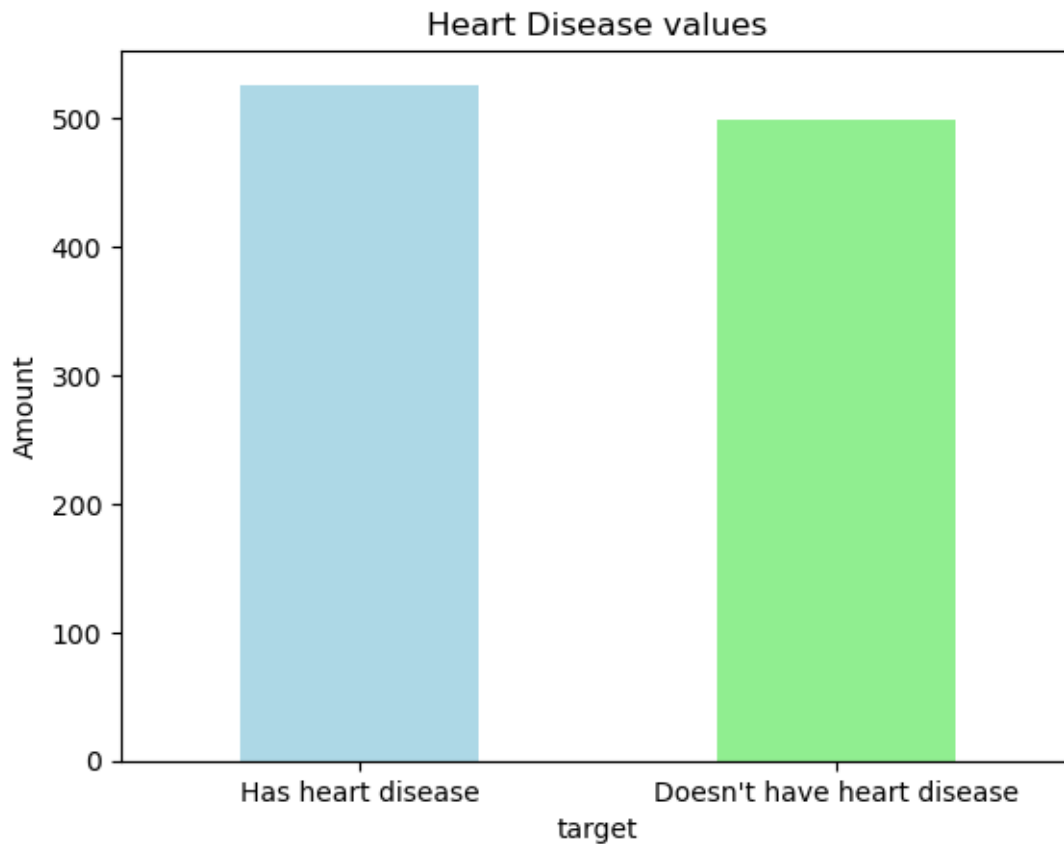
	precision	recall	f1-score	support
0	0.98	1.00	0.99	124
1	1.00	0.98	0.99	133
accuracy			0.99	257
macro avg	0.99	0.99	0.99	257
weighted avg	0.99	0.99	0.99	257

```
[30]: # Create a confusion matrix
conf_mat = confusion_matrix(y_test, y_preds)
conf_mat
```

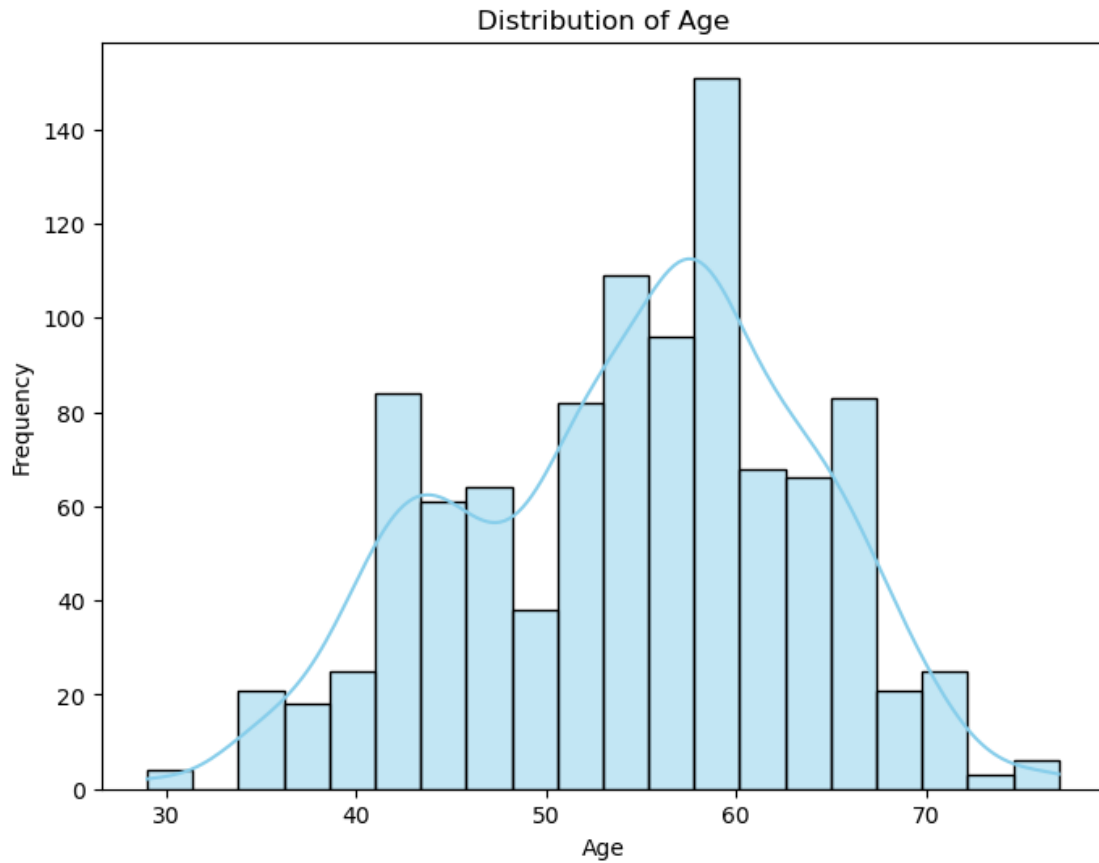
```
[30]: array([[124,  0],
        [ 3, 130]], dtype=int64)
```

## 0.2 Matplotlib and SnS

```
[31]: fig = heart_disease.target.value_counts().plot(kind = 'bar', color=["lightblue",  
    ↪ 'lightgreen'])  
fig.set_xticklabels(labels=['Has heart disease', "Doesn't have heart disease"],  
    ↪ rotation=0);  
plt.title("Heart Disease values")  
plt.ylabel("Amount");
```

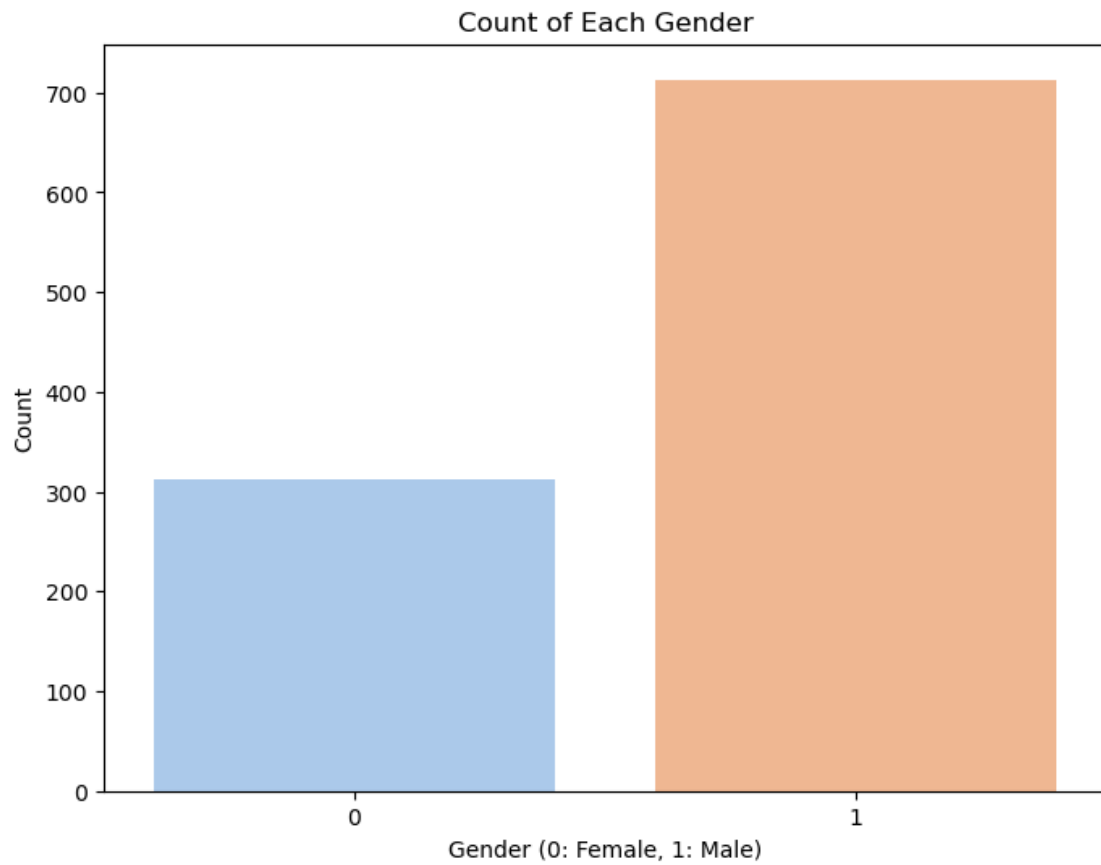


```
[32]: plt.figure(figsize=(8, 6))  
sns.histplot(heart_disease['age'], bins=20, kde=True, color='skyblue')  
plt.title('Distribution of Age')  
plt.xlabel('Age')  
plt.ylabel('Frequency')  
plt.show()
```

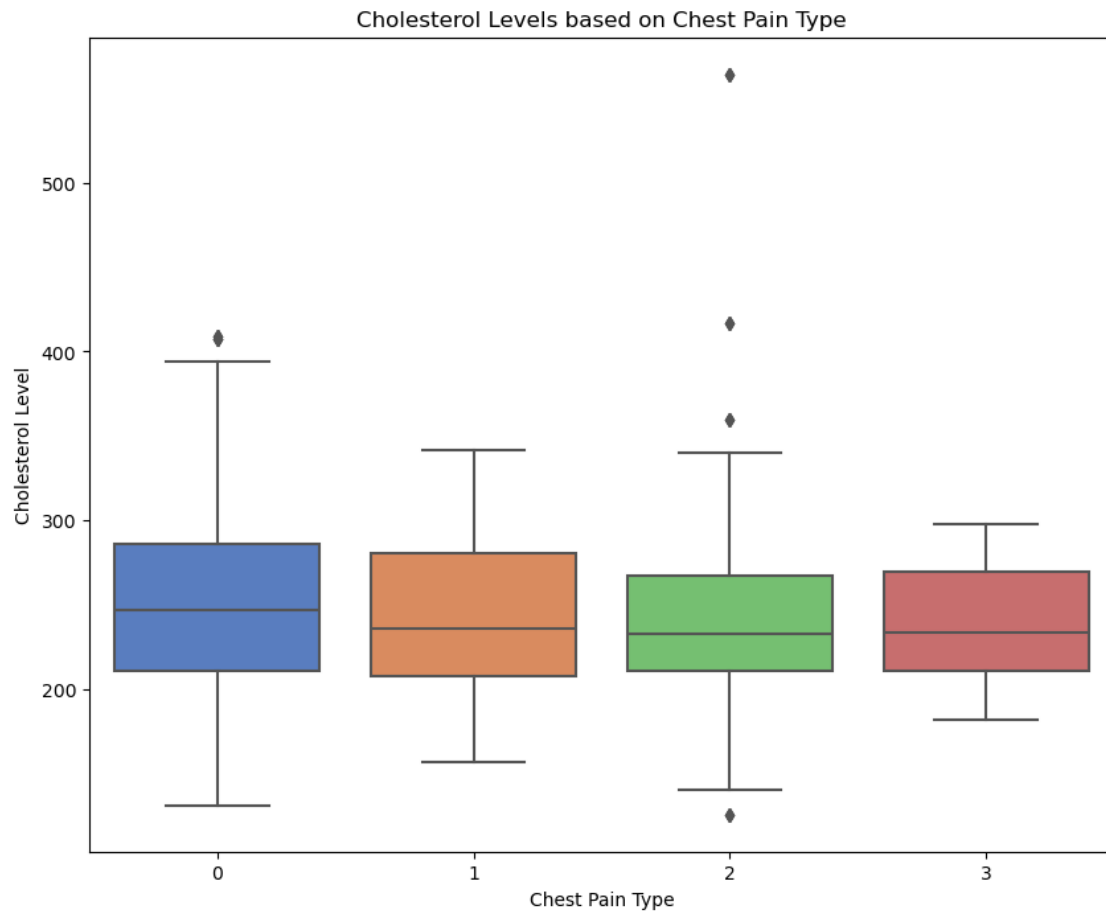


```
[33]: plt.figure(figsize=(8, 6))
sns.countplot(x='sex', data=heart_disease, palette='pastel')
plt.title('Count of Each Gender')
plt.xlabel('Gender (0: Female, 1: Male)')
plt.ylabel('Count')
plt.show()
```

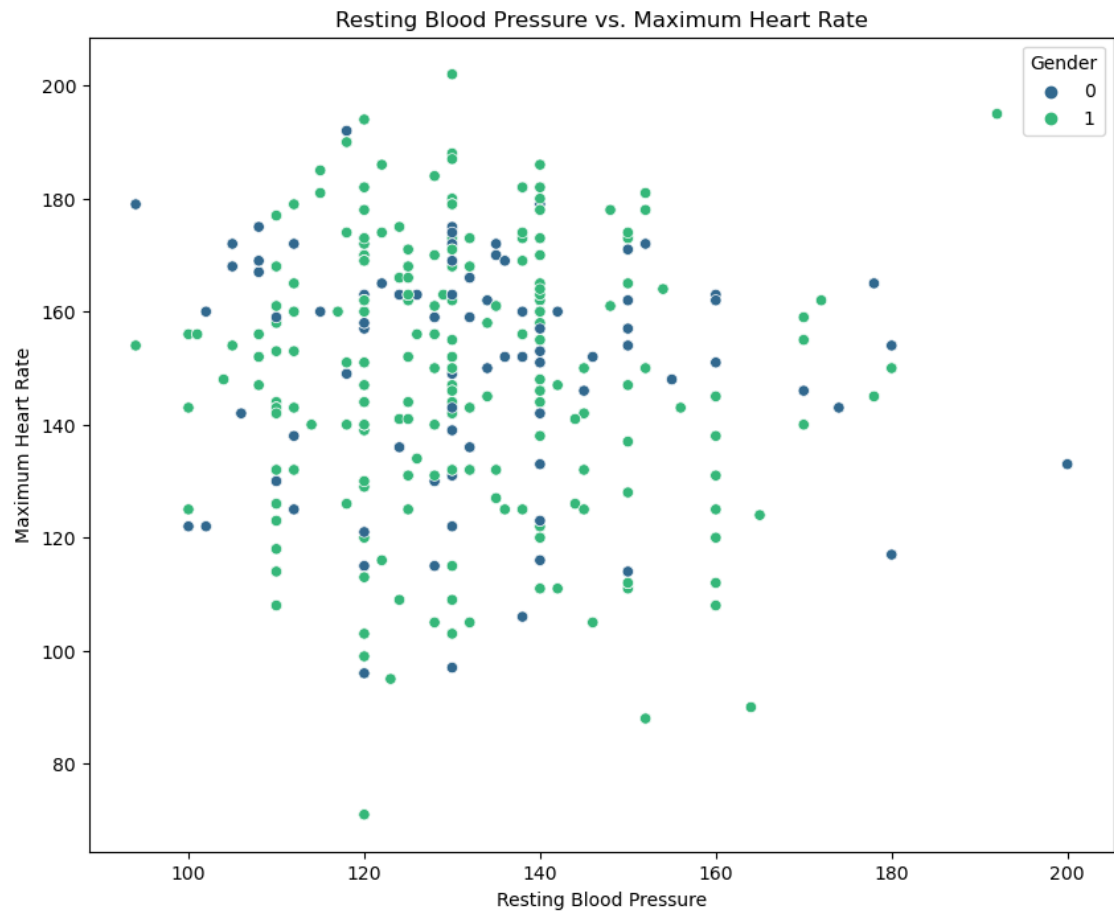




```
[34]: plt.figure(figsize=(10, 8))
sns.boxplot(x='cp', y='chol', data=heart_disease, palette='muted')
plt.title('Cholesterol Levels based on Chest Pain Type')
plt.xlabel('Chest Pain Type')
plt.ylabel('Cholesterol Level')
plt.show()
```



```
[35]: plt.figure(figsize=(10, 8))
sns.scatterplot(x='trestbps', y='thalach', data=heart_disease, hue='sex',
               palette='viridis')
plt.title('Resting Blood Pressure vs. Maximum Heart Rate')
plt.xlabel('Resting Blood Pressure')
plt.ylabel('Maximum Heart Rate')
plt.legend(title='Gender')
plt.show()
```





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## **Result and Discussion :**

Thus we have successfully used the python libraries.