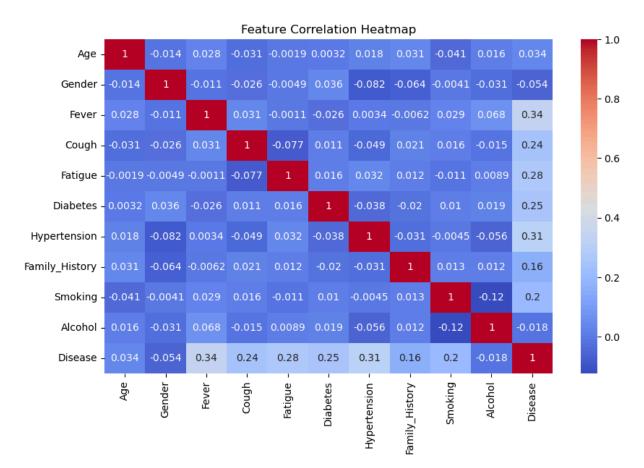
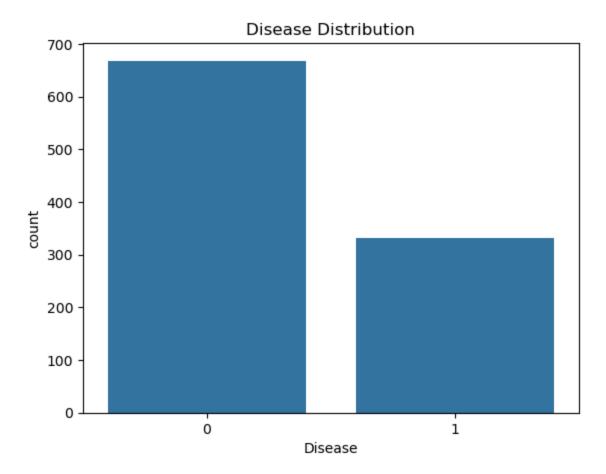
import pandas as pd from sklearn.preprocessing import LabelEncoder, StandardScaler

```
In [3]: df = pd.read_csv("Downloads/medical_dataset.csv")
In [5]: encoder = LabelEncoder()
         df["Gender"] = encoder.fit_transform(df["Gender"])
In [7]: scaler = StandardScaler()
         df[["Age"]] = scaler.fit_transform(df[["Age"]])
In [9]: print(df.head())
                                   Cough Fatigue Diabetes Hypertension \
                Age Gender
                             Fever
        0 0.449200
                          0
                                 0
                                        0
                                                 0
                                                           0
        1 1.197866
                         0
                                 1
                                        0
                                                 1
                                                           0
                                                                         0
        2 -0.126697
                         1
                                 0
                                        1
                                                 0
                                                           0
                                                                         0
        3 -0.932953
                          0
                                 0
                                        0
                                                 1
                                                                         0
        4 0.679558
                         1
                                 1
                                        1
                                                 0
                                                                         0
           Family_History Smoking Alcohol Disease
        0
                        0
                                 0
                                          1
                        1
                                 0
                                          1
                                                   0
        1
                        0
        2
                                 1
                                          0
                                                   0
        3
                        0
                                 1
                                          0
                                                   0
        4
                                          1
                                                   1
In [11]: import seaborn as sns
         import matplotlib.pyplot as plt
In [17]: plt.figure(figsize=(10, 6))
         sns.heatmap(df.corr(), annot=True, cmap="coolwarm")
         plt.title("Feature Correlation Heatmap")
         plt.show()
```



```
In [19]: sns.countplot(x="Disease", data=df)
  plt.title("Disease Distribution")
  plt.show()
```



```
Accuracy: 1.0
        Classification Report:
                       precision
                                    recall f1-score
                                                        support
                   0
                           1.00
                                     1.00
                                               1.00
                                                           130
                           1.00
                                     1.00
                                               1.00
                                                           70
                                                           200
            accuracy
                                               1.00
                           1.00
                                     1.00
                                               1.00
                                                           200
           macro avg
        weighted avg
                           1.00
                                     1.00
                                                           200
                                               1.00
In [57]: from sklearn.model selection import GridSearchCV
         param_grid = {
              "n_estimators": [50, 100, 150],
              "max_depth": [5, 10, 15],
         }
         grid_search = GridSearchCV(RandomForestClassifier(), param_grid, cv=5)
         grid_search.fit(X_train, y_train)
         print("Best Parameters:", grid_search.best_params_)
        Best Parameters: {'max_depth': 10, 'n_estimators': 50}
In [60]: new_patient = [[45, 1, 1, 0, 1, 0, 1, 0, 0, 1]]
         prediction = model.predict(new_patient)
         print("Disease Prediction:", "Yes" if prediction[0] == 1 else "No")
        Disease Prediction: Yes
```

C:\Users\640 G2\anaconda3\Lib\site-packages\sklearn\base.py:493: UserWarning: X does not have valid feature names, but RandomForestClassifier was fitted with feature nam

warnings.warn(

In []: