```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report, accuracy_score


df = pd.read_csv('/content/archive (3).zip')
```

```
df.head()
```

	Age	Sex	Chest pain type	BP	Cholesterol	FBS over 120	EKG results	Max HR	Exercise angina	ST depression	Slope of ST	Number of vessels fluro	Thallium	Heart Disease
0	70	1	4	130	322	0	2	109	0	2.4	2	3	3	Presence
1	67	0	3	115	564	0	2	160	0	1.6	2	0	7	Absence
2	57	1	2	124	261	0	0	141	0	0.3	1	0	7	Presence
3	64	1	4	128	263	0	0	105	1	0.2	2	1	7	Absence

Next steps: Generate code with df | View recommended plots | New interactive sheet

```
df.tail()
```

	Age	Sex	Chest pain type	BP	Cholesterol	FBS over 120	EKG results	Max HR	Exercise angina	ST depression	Slope of ST	Number of vessels fluro	Thallium	Heart Disease
265	52	1	3	172	199	1	0	162	0	0.5	1	0	7	Absence
266	44	1	2	120	263	0	0	173	0	0.0	1	0	7	Absence
267	56	0	2	140	294	0	2	153	0	1.3	2	0	3	Absence
268	57	1	4	140	192	0	0	148	0	0.4	2	0	6	Absence

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 270 entries, 0 to 269
Data columns (total 14 columns):
 #   Column                  Non-Null Count  Dtype  
---  ------                  --------------  -----  
 0   Age                     270 non-null    int64  
 1   Sex                     270 non-null    int64  
 2   Chest pain type         270 non-null    int64  
 3   BP                      270 non-null    int64  
 4   Cholesterol             270 non-null    int64  
 5   FBS over 120            270 non-null    int64  
 6   EKG results             270 non-null    int64  
 7   Max HR                  270 non-null    int64  
 8   Exercise angina         270 non-null    int64  
 9   ST depression           270 non-null    float64
 10  Slope of ST             270 non-null    int64  
 11  Number of vessels fluro 270 non-null    int64  
 12  Thallium                270 non-null    int64  
 13  Heart Disease           270 non-null    object 
dtypes: float64(1), int64(12), object(1)
memory usage: 29.7+ KB
```

```
df.isnull().sum()
```

	0
Age	0
Sex	0
Chest pain type	0
BP	0
Cholesterol	0
FBS over 120	0
EKG results	0
Max HR	0
Exercise angina	0
ST depression	0
Slope of ST	0
Number of vessels fluro	0
Thallium	0
Heart Disease	0

dtype: int64

```
label_encoder = LabelEncoder()
df['Heart Disease'] = label_encoder.fit_transform(df['Heart Disease'])


X = df.drop('Heart Disease', axis=1)
y = df['Heart Disease']


X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, random_state=42
)


model = RandomForestClassifier(random_state=42)


model.fit(X_train, y_train)
```

```
RandomForestClassifier
RandomForestClassifier(random_state=42)
```

```
y_pred = model.predict(X_test)


accuracy = accuracy_score(y_test, y_pred)
report = classification_report(y_test, y_pred, target_names=label_encoder.classes_)


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

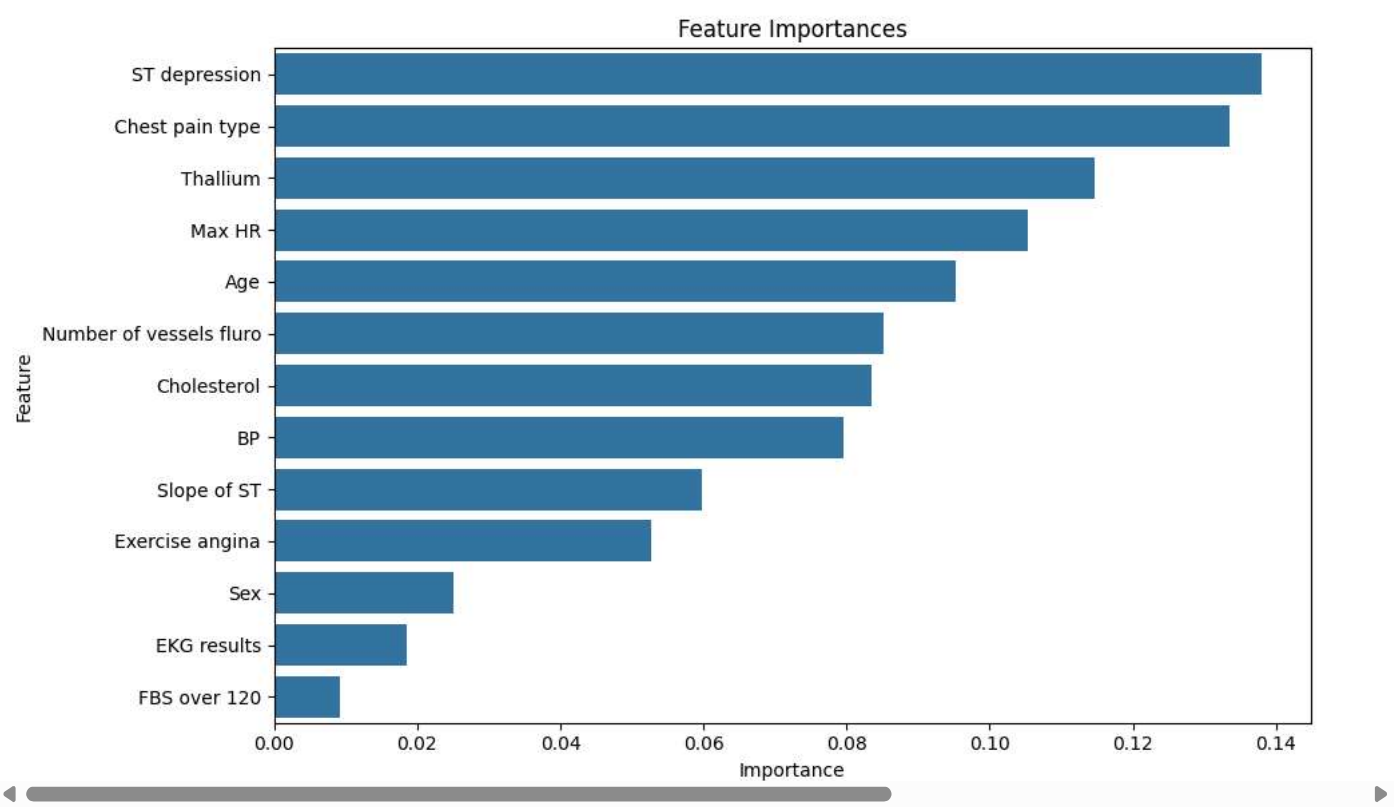
```
Accuracy: 0.80
```

```
print("Classification Report:")

print(report)
```

```
Classification Report:
                precision    recall  f1-score   support

       Absence       0.81      0.88      0.84        33
      Presence       0.78      0.67      0.72        21

      accuracy                           0.80        54
     macro avg       0.79      0.77      0.78        54
  weighted avg       0.79      0.80      0.79        54
```

```
importances = model.feature_importances_
```

```
features = X.columns
indices = importances.argsort()[::-1]


plt.figure(figsize=(10, 6))
sns.barplot(x=importances[indices], y=features[indices])
plt.title("Feature Importances")
plt.xlabel("Importance")
plt.ylabel("Feature")
plt.tight_layout()
plt.show()
```




```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LogisticRegression
from xgboost import XGBClassifier
from sklearn.metrics import classification_report, accuracy_score


df = pd.read_csv('/content/archive (3).zip')


df['Heart Disease'] = df['Heart Disease'].map({'Presence': 1, 'Absence': 0})


label_encoder = LabelEncoder()
df['Heart Disease'] = label_encoder.fit_transform(df['Heart Disease'])


X = df.drop('Heart Disease', axis=1)
y = df['Heart Disease']


scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)


X = df.drop('Heart Disease', axis=1)
y = df['Heart Disease']


X_train, X_test, y_train, y_test = train_test_split(
    X_scaled, y, test_size=0.2, random_state=42
)
```

```
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, random_state=42
)


# Drop rows where the target is NaN
df = df.dropna(subset=['Heart Disease'])

# Proceed with feature/target split
X = df.drop('Heart Disease', axis=1)
y = df['Heart Disease']

# (Continue with StandardScaler, train_test_split, etc.)



print(df['Heart Disease'].isna().sum())
```

```
0
```

```
print(df['Heart Disease'].isna().sum())
```

```
0
```

```
df = df.fillna(df.mean(numeric_only=True))


log_reg = LogisticRegression()



log_reg.fit(X_train, y_train)
```

```
/usr/local/lib/python3.11/dist-packages/sklearn/linear_model/_logistic.py:465: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. OF ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
  n_iter_i = _check_optimize_result(
▾ LogisticRegression
LogisticRegression()
```

```
log_preds = log_reg.predict(X_test)


print("=== Logistic Regression Results ===")
print("Accuracy:", accuracy_score(y_test, log_preds))
print(classification_report(y_test, log_preds))
```

```
=== Logistic Regression Results ===
Accuracy: 0.9074074074074074
              precision    recall  f1-score   support

           0       0.89      0.97      0.93        33
           1       0.94      0.81      0.87        21

    accuracy                           0.91        54
   macro avg       0.92      0.89      0.90        54
weighted avg       0.91      0.91      0.91        54
```

```
xgb = XGBClassifier(use_label_encoder=False, eval_metric='logloss')
xgb.fit(X_train, y_train)
```

```
/usr/local/lib/python3.11/dist-packages/xgboost/core.py:158: UserWarning: [14:29:40] WARNING: /workspace/src/learner.cc:740:
Parameters: { "use_label_encoder" } are not used.

  warnings.warn(smsg, UserWarning)
```

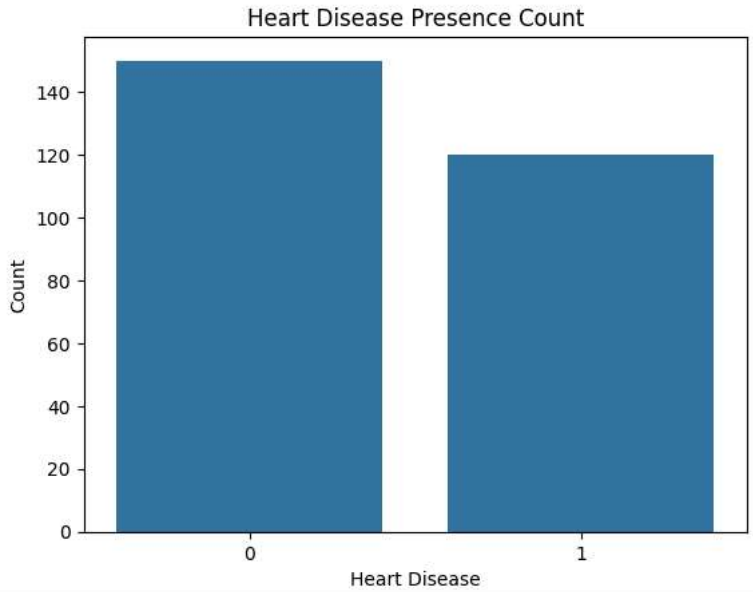
```
XGBClassifier(base_score=None, booster=None, callbacks=None,
              colsample_bylevel=None, colsample_bynode=None,
              colsample_bytree=None, device=None, early_stopping_rounds=None,
              enable_categorical=False, eval_metric='logloss',
              feature_types=None, gamma=None, grow_policy=None,
              importance_type=None, interaction_constraints=None,
              learning_rate=None, max_bin=None, max_cat_threshold=None,
              max_cat_to_onehot=None, max_delta_step=None, max_depth=None,
              max_leaves=None, min_child_weight=None, missing=nan,
              monotone_constraints=None, multi_strategy=None, n_estimators=None,
              n_jobs=None, num_parallel_tree=None, random_state=None, ...)
```

```
xgb_preds = xgb.predict(X_test)
```

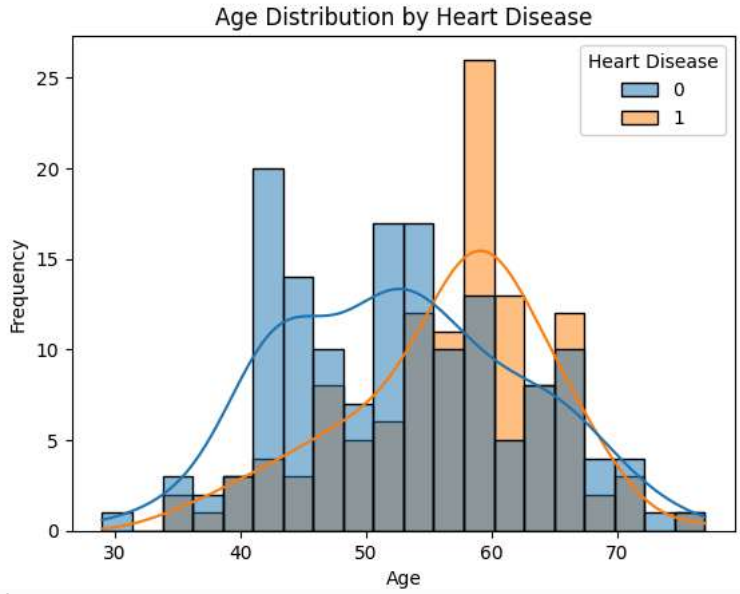
```
print("=== XGBoost Results ===")
print("Accuracy:", accuracy_score(y_test, xgb_preds))
print(classification_report(y_test, xgb_preds))
```

```
=== XGBoost Results ===
Accuracy: 0.8148148148148148
              precision    recall  f1-score   support

           0       0.83      0.88      0.85        33
           1       0.79      0.71      0.75        21

    accuracy                           0.81        54
   macro avg       0.81      0.80      0.80        54
weighted avg       0.81      0.81      0.81        54
```

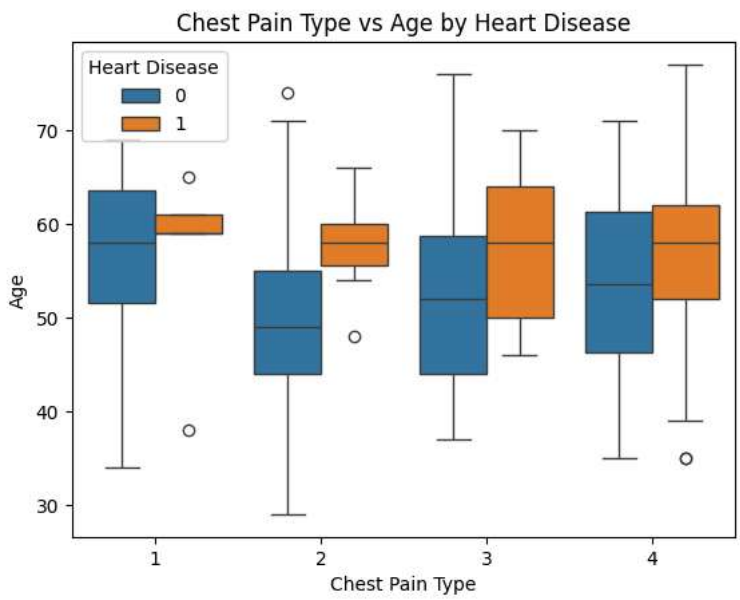
```
import seaborn as sns
import matplotlib.pyplot as plt

sns.countplot(data=df, x='Heart Disease')
plt.title('Heart Disease Presence Count')
plt.xlabel('Heart Disease')
plt.ylabel('Count')
plt.show()
```



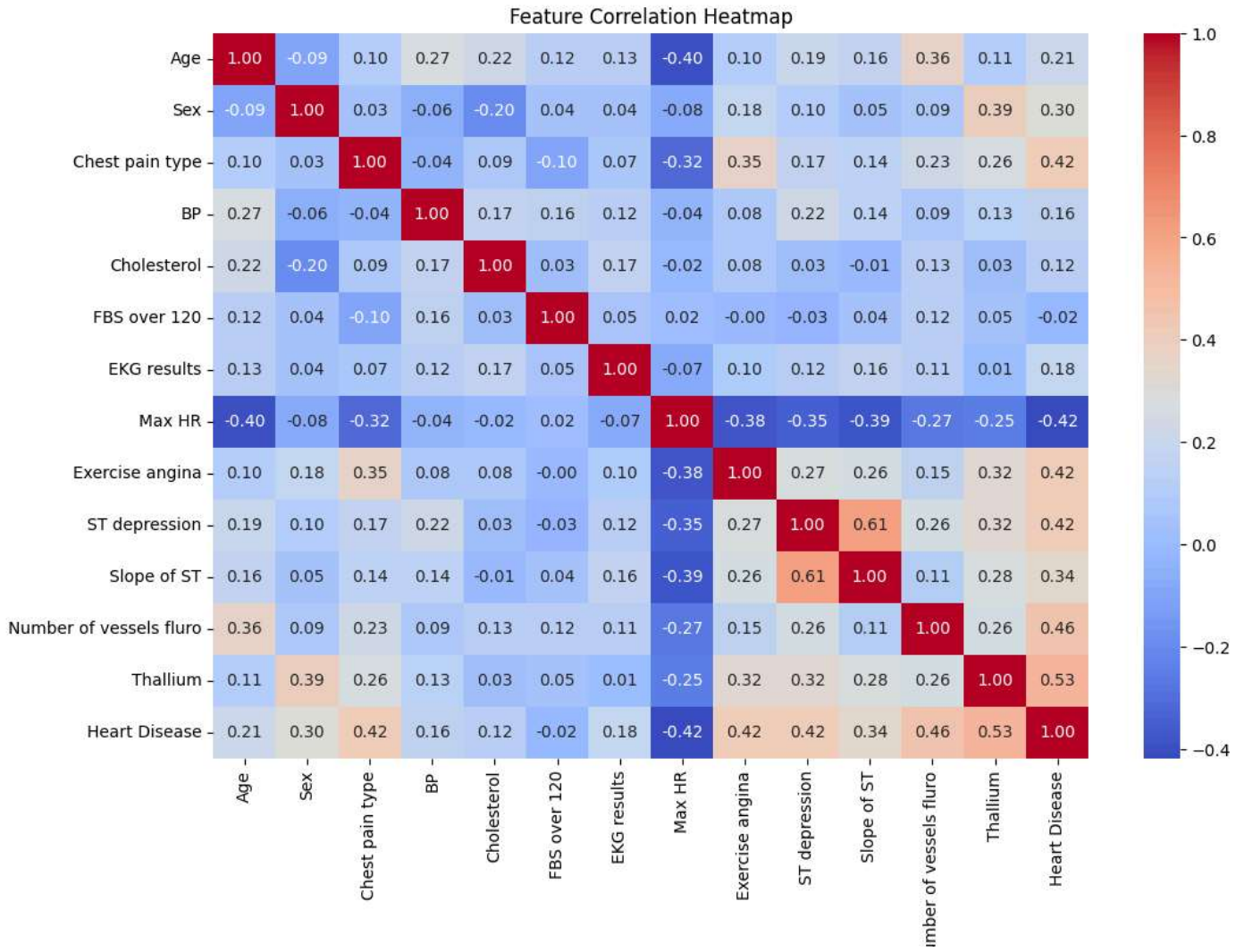

```
sns.histplot(data=df, x='Age', hue='Heart Disease', kde=True, bins=20)
plt.title('Age Distribution by Heart Disease')
plt.xlabel('Age')
plt.ylabel('Frequency')
plt.show()
```




```
sns.boxplot(data=df, x='Chest pain type', y='Age', hue='Heart Disease')
plt.title('Chest Pain Type vs Age by Heart Disease')
plt.xlabel('Chest Pain Type')
plt.ylabel('Age')
plt.show()
```




```
plt.figure(figsize=(12, 8))
corr = df.corr(numeric_only=True)
sns.heatmap(corr, annot=True, cmap='coolwarm', fmt=".2f")
plt.title('Feature Correlation Heatmap')
plt.show()
```


Feature Correlation Heatmap
Age 1.00 -0.09 0.10 0.27 0.22 0.12 0.13 -0.40 0.10 0.19 0.16 0.36 0.11 0.21
Sex -0.09 1.00 0.03 -0.06 -0.20 0.04 0.04 -0.08 0.18 0.10 0.05 0.09 0.39 0.30
Chest pain type 0.10 0.03 1.00 -0.04 0.09 -0.10 0.07 -0.32 0.35 0.17 0.14 0.23 0.26 0.42
BP 0.27 -0.06 -0.04 1.00 0.17 0.16 0.12 -0.04 0.08 0.22 0.14 0.09 0.13 0.16
Cholesterol 0.22 -0.20 0.09 0.17 1.00 0.03 0.17 -0.02 0.08 0.03 -0.01 0.13 0.03 0.12
FBS over 120 0.12 0.04 -0.10 0.16 0.03 1.00 0.05 0.02 -0.00 -0.03 0.04 0.12 0.05 -0.02
EKG results 0.13 0.04 0.07 0.12 0.17 0.05 1.00 -0.07 0.10 0.12 0.16 0.11 0.01 0.18
Max HR -0.40 -0.08 -0.32 -0.04 -0.02 0.02 -0.07 1.00 -0.38 -0.35 -0.39 -0.27 -0.25 -0.42
Exercise angina 0.10 0.18 0.35 0.08 0.08 -0.00 0.10 -0.38 1.00 0.27 0.26 0.15 0.32 0.42
ST depression 0.19 0.10 0.17 0.22 0.03 -0.03 0.12 -0.35 0.27 1.00 0.61 0.26 0.32 0.42
Slope of ST 0.16 0.05 0.14 0.14 -0.01 0.04 0.16 -0.39 0.26 0.61 1.00 0.11 0.28 0.34
Number of vessels fluro 0.36 0.09 0.23 0.09 0.13 0.12 0.11 -0.27 0.15 0.26 0.11 1.00 0.26 0.46
Thallium 0.11 0.39 0.26 0.13 0.03 0.05 0.01 -0.25 0.32 0.32 0.28 0.26 1.00 0.53
Heart Disease 0.21 0.30 0.42 0.16 0.12 -0.02 0.18 -0.42 0.42 0.42 0.34 0.46 0.53 1.00
Age
Sex
Chest pain type
BP
Cholesterol
FBS over 120
EKG results
Max HR
Exercise angina
ST depression
Slope of ST
imber of vessels fluro
Thallium
Heart Disease
1.0
0.8
0.6
0.4
0.2
0.0
−0.2
−0.4