Install and Import Required Libraries

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from google.colab import files
```

Load the Dataset

```
df = pd.read_csv('heart_disease_uci.csv')
```

normal

Display first few rows

```
print(df.head(5))
```

```
\rightarrow
                                                                      fbs \
       id
                                               cp trestbps
                                                              chol
           age
                   sex
                          dataset
        1
                 Male Cleveland
                                   typical angina
                                                      145.0
                                                             233.0
                                                                     True
                                     asymptomatic
                                                             286.0
            67
                 Male Cleveland
                                                      160.0
                                                                    False
        3
            67
                 Male Cleveland
                                     asymptomatic
                                                      120.0
                                                             229.0 False
            37
                 Male Cleveland
                                      non-anginal
                                                             250.0 False
                                                      130.0
            41 Female Cleveland atypical angina
                                                      130.0 204.0 False
              restecg thalch exang oldpeak
                                                    slope
                                                            ca \
      lv hypertrophy
                              False
                                              downsloping
                                                           0.0
                       150.0
                                         2.3
       lv hypertrophy
                       108.0
                                         1.5
                                                     flat 3.0
                               True
       lv hypertrophy
                               True
                                         2.6
                                                     flat 2.0
                       129.0
    3
               normal
                       187.0 False
                                         3.5
                                              downsloping
                                                           0.0
    4 lv hypertrophy
                       172.0 False
                                                upsloping 0.0
                                         1.4
                    thal num
    0
            fixed defect
                            0
```

```
reversable defect
                             1
     3
                   normal
                   normal
                             0
     4
Inspect Data
print("\nDataset Info:")
print(df.info())
print("\nMissing Values:")
print(df.isnull().sum())
     Dataset Info:
     <class 'pandas.core.frame.DataFrame'>
     RangeIndex: 920 entries, 0 to 919
     Data columns (total 16 columns):
         Column
                    Non-Null Count Dtype
                    _____
                                    ----
         id
                                    int64
      0
                    920 non-null
                                   int64
                    920 non-null
      1
         age
                    920 non-null
      2
                                   obiect
         sex
                    920 non-null
                                   object
         dataset
                    920 non-null
                                   object
      4
         ср
      5
         trestbps
                    861 non-null
                                   float64
      6
         chol
                    890 non-null
                                   float64
      7
         fbs
                    830 non-null
                                   object
         restecg
                   918 non-null
                                   object
      9
         thalch
                    865 non-null
                                   float64
                    865 non-null
                                   object
      10
         exang
     11 oldpeak
                   858 non-null
                                   float64
     12 slope
                    611 non-null
                                   obiect
                    309 non-null
      13 ca
                                   float64
     14 thal
                                   object
                    434 non-null
                    920 non-null
                                   int64
      15 num
     dtypes: float64(5), int64(3), object(8)
     memory usage: 115.1+ KB
```

None

 $\overline{\mathbf{x}}$

```
Missing Values:
id
              0
age
sex
dataset
ср
trestbps
             59
chol
             30
fbs
             90
restecg
              2
thalch
             55
             55
exang
oldpeak
             62
slope
            309
            611
ca
thal
            486
num
dtype: int64
```

Rename target column and clean

```
df.rename(columns={'num': 'target'}, inplace=True)
df['target'] = (df['target'] > 0).astype(int)

Drop'id' column (not useful)

df.drop(columns=['id'], inplace=True)

Handle Missing Values

df.replace({'': np.nan, '?': np.nan}, inplace=True)
```

Convert numeric columns to float where necessary

```
numeric_cols = ['age', 'trestbps', 'chol', 'thalch', 'oldpeak', 'ca']
for col in numeric cols:
    df[col] = pd.to numeric(df[col], errors='coerce')
Fill missing numeric values with median
for col in numeric cols:
    df[col].fillna(df[col].median(), inplace=True)
/tmp/ipython-input-3154000584.py:2: FutureWarning: A value is trying to be set on a copy of a DataFrame or Series through of
     The behavior will change in pandas 3.0. This inplace method will never work because the intermediate object on which we are
     For example, when doing 'df[col].method(value, inplace=True)', try using 'df.method({col: value}, inplace=True)' or df[col]
       df[col].fillna(df[col].median(), inplace=True)
Categorical columns: fill with mode
categorical cols = ['cp', 'fbs', 'restecg', 'exang', 'slope', 'thal', 'sex', 'dataset']
for col in categorical cols:
    df[col].fillna(df[col].mode()[0], inplace=True)
/tmp/ipython-input-3512451438.py:3: FutureWarning: A value is trying to be set on a copy of a DataFrame or Series through
     The behavior will change in pandas 3.0. This inplace method will never work because the intermediate object on which we are
     For example, when doing 'df[col].method(value, inplace=True)', try using 'df.method({col: value}, inplace=True)' or df[col]
       df[col].fillna(df[col].mode()[0], inplace=True)
     /tmp/ipython-input-3512451438.py:3: FutureWarning: Downcasting object dtype arrays on .fillna, .ffill, .bfill is deprecated
```

```
df[col].fillna(df[col].mode()[0], inplace=True)
```

Encode Categorical Variables

```
categorical cols.remove('sex') # We'll handle 'sex' separately as binary
df = pd.get dummies(df, columns=categorical cols, drop first=True)
Encode 'sex': Male = 1, Female = 0
df['sex'] = df['sex'].map({'Male': 1, 'Female': 0})
print("Shape after encoding:", df.shape)
Shape after encoding: (920, 22)
Define Features and Target
X = df.drop('target', axis=1)
y = df['target']
Train-Test Split
from sklearn.model selection import train test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)
```

Scale the Features

```
from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)
```

Train Models

```
from sklearn.linear_model import LogisticRegression
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_score, roc_auc_score, confusion_matrix, roc_curve
```

Dictionary to store results

```
models = {}
results = []
```

Logistic Regression

```
lr = LogisticRegression(max_iter=1000, random_state=42)
lr.fit(X_train_scaled, y_train)
y_pred_lr = lr.predict(X_test_scaled)
y_prob_lr = lr.predict_proba(X_test_scaled)[:, 1]
models['Logistic Regression'] = lr
results.append({
    'Model': 'Logistic Regression',
    'Accuracy': accuracy_score(y_test, y_pred_lr),
    'Precision': precision_score(y_test, y_pred_lr),
    'Recall': recall_score(y_test, y_pred_lr),
    'F1': f1_score(y_test, y_pred_lr),
```

```
'ROC-AUC': roc auc score(y test, y prob lr)
})
Decision Tree
dt = DecisionTreeClassifier(random state=42, max depth=8, min samples split=5)
dt.fit(X train, y train) # No scaling needed
y pred dt = dt.predict(X test)
y prob dt = dt.predict_proba(X_test)[:, 1]
models['Decision Tree'] = dt
results.append({
    'Model': 'Decision Tree',
    'Accuracy': accuracy score(y test, y pred dt),
    'Precision': precision score(y test, y pred dt),
    'Recall': recall score(y test, y pred dt),
    'F1': f1 score(y test, y pred dt),
    'ROC-AUC': roc auc_score(y_test, y_prob_dt)
})
Random Forest
rf = RandomForestClassifier(n estimators=100, max depth=7, min samples split=5,
                           random state=42, class weight='balanced')
rf.fit(X train, y train)
y pred rf = rf.predict(X test)
y prob rf = rf.predict proba(X test)[:, 1]
models['Random Forest'] = rf
results.append({
    'Model': 'Random Forest',
    'Accuracy': accuracy_score(y_test, y_pred_rf),
    'Precision': precision_score(y_test, y_pred_rf),
    'Recall': recall score(y test, y pred rf),
    'F1': f1 score(y test, y pred rf),
    'ROC-AUC': roc_auc_score(y_test, y_prob_rf)
```

```
})
```

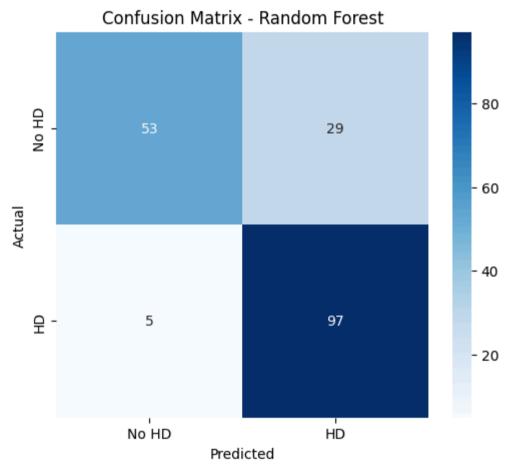
Display Results

```
results df = pd.DataFrame(results)
print("\n Model Performance Comparison:")
print(results df.round(4))
\overline{2}
     Model Performance Comparison:
                      Model Accuracy Precision Recall
                                                                F1 ROC-AUC
     0 Logistic Regression
                                0.8370
                                           0.8273 0.8922 0.8585
                                                                     0.9213
                               0.7989
                                           0.7876 0.8725 0.8279
              Decision Tree
                                                                     0.8488
     2
              Random Forest
                                           0.8505 0.8922 0.8708
                               0.8533
                                                                     0.9219
Highlight best model
best model = results df.loc[results df['F1'].idxmax()]['Model']
print(f"\n Best Model (by F1-score): {best model}")
\overline{\mathbf{x}}
      Best Model (by F1-score): Random Forest
Confusion Matrix for Best Model
def plot confusion matrix(model name):
    y_pred = models[model_name].predict(X_test if 'Tree' in model_name else X_test_scaled)
    cm = confusion matrix(y test, y pred)
    plt.figure(figsize=(6,5))
    sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['No HD', 'HD'], yticklabels=['No HD', 'HD'])
    plt.title(f'Confusion Matrix - {model name}')
    plt.ylabel('Actual')
```

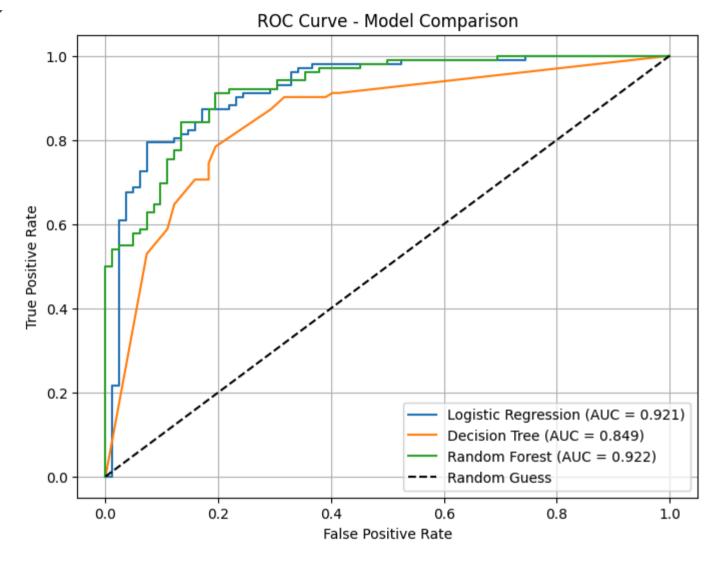
```
plt.xlabel('Predicted')
plt.show()
```

plot_confusion_matrix(best_model)

/usr/local/lib/python3.12/dist-packages/sklearn/utils/validation.py:2739: UserWarning: X does not have valid feature names, warnings.warn(



```
plt.figure(figsize=(8,6))
for model name in models.keys():
    if model name == 'Logistic Regression':
       y prob = lr.predict proba(X test scaled)[:, 1]
       fpr, tpr, = roc curve(y test, y prob)
        plt.plot(fpr, tpr, label=f"{model name} (AUC = {roc auc score(y test, y prob):.3f})")
    elif model name == 'Decision Tree':
       v prob = dt.predict proba(X test)[:, 1]
       fpr, tpr, _ = roc_curve(y_test, y_prob)
        plt.plot(fpr, tpr, label=f"{model name} (AUC = {roc auc score(y test, y prob):.3f})")
    elif model name == 'Random Forest':
       y prob = rf.predict proba(X test)[:, 1]
       fpr, tpr, = roc curve(y test, y prob)
        plt.plot(fpr, tpr, label=f"{model name} (AUC = {roc auc score(y test, y prob):.3f})")
plt.plot([0,1], [0,1], 'k--', label='Random Guess')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('ROC Curve - Model Comparison')
plt.legend()
plt.grid(True)
plt.show()
```



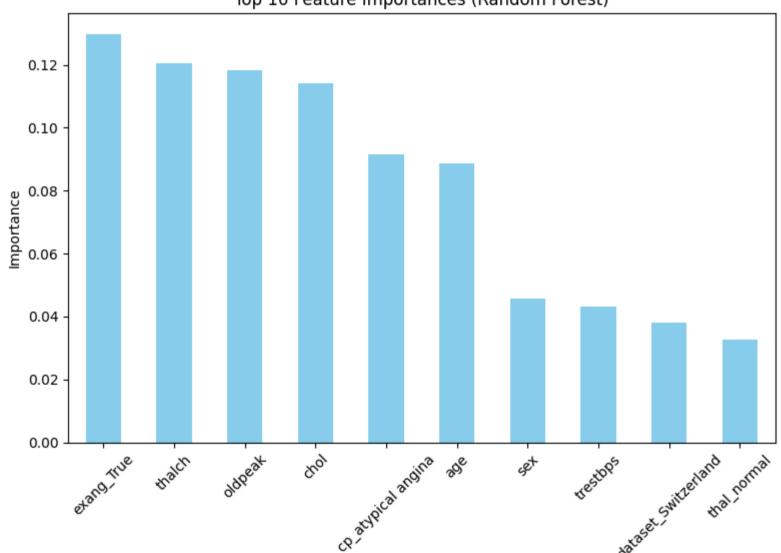
Feature Importance (Random Forest)

```
if best_model == 'Random Forest':
    feat_importance = pd.Series(rf.feature_importances_, index=X.columns).sort_values(ascending=False)
    plt.figure(figsize=(8,6))
    feat_importance.head(10).plot(kind='bar', color='skyblue')
```

```
plt.title('Top 10 Feature Importances (Random Forest)')
plt.ylabel('Importance')
plt.xticks(rotation=45)
plt.tight_layout()
plt.show()
```

→

Top 10 Feature Importances (Random Forest)



save model

```
import joblib

# Select best model
best_model_instance = models[best_model]

# Save model and scaler
joblib.dump(best_model_instance, 'heart_disease_model.pkl')
joblib.dump(scaler, 'scaler.pkl')
joblib.dump(X.columns.tolist(), 'feature_columns.pkl') # Save feature names for inference

print("\n Best model and preprocessing objects saved!")
print("Files: 'heart_disease_model.pkl', 'scaler.pkl', 'feature_columns.pkl'")

Best model and preprocessing objects saved!
Files: 'heart_disease_model.pkl', 'scaler.pkl', 'feature_columns.pkl'
```