

```

import pandas as pd

df = pd.read_csv('/content/HeartDiseaseTrain-Test.csv')

# 3 Data Exploration
print("\nFirst 5 rows of the dataset:")
print(df.head())

print("\nShape of the dataset:", df.shape)
print("\nColumns:", df.columns.tolist())
print("\nData Types & Missing Values:")
df.info()

print("\nSummary Statistics:")
print(df.describe())

print("\nMissing values per column:\n", df.isnull().sum())
print("\nDuplicate rows:", df.duplicated().sum())

# 4 Data Visualization (Optional, for better understanding)
import seaborn as sns
import matplotlib.pyplot as plt

# Example: Plot distribution of target variable (assuming 'target' is the disease indicator)
sns.countplot(x='target', data=df)
plt.title('Distribution of Disease Presence (0=No, 1=Yes)')
plt.show()

# 5 Prepare Features and Target
target = 'target' # Change to actual target column if named differently
features = df.columns.drop(target)
print("\nFeatures:", features.tolist())

# 6 Convert Categorical Columns
categorical_cols = df.select_dtypes(include=['object']).columns.tolist()
print("\nCategorical Columns:", categorical_cols)

df_encoded = pd.get_dummies(df, drop_first=True)

# 3 Check for Missing Values
print("\n🔴 Missing values per column:\n", df.isnull().sum())

# Fill missing numeric columns with median
numeric_cols = df.select_dtypes(include=['int64', 'float64']).columns
for col in numeric_cols:
    if df[col].isnull().sum() > 0:
        median_val = df[col].median()
        df[col].fillna(median_val, inplace=True)
        print(f"Filled missing values in '{col}' with median: {median_val}")

# Fill missing categorical columns with mode
categorical_cols = df.select_dtypes(include=['object']).columns
for col in categorical_cols:
    if df[col].isnull().sum() > 0:
        mode_val = df[col].mode()[0]
        df[col].fillna(mode_val, inplace=True)
        print(f"Filled missing values in '{col}' with mode: {mode_val}")

# 4 Check for Duplicates
duplicates = df.duplicated().sum()
print(f"\n🔴 Duplicate rows: {duplicates}")
if duplicates > 0:
    df.drop_duplicates(inplace=True)
    print("✅ Duplicates removed.")

# 5 Detect and Handle Outliers (Optional: here we cap them using IQR method)
def cap_outliers(column):
    Q1 = df[column].quantile(0.25)
    Q3 = df[column].quantile(0.75)
    IQR = Q3 - Q1
    lower = Q1 - 1.5 * IQR
    upper = Q3 + 1.5 * IQR
    df[column] = df[column].clip(lower, upper)
    print(f"✅ Outliers capped for '{column}'")

for col in numeric_cols:

```

```

cap_outliers(col)

# 4 Encode Categorical Features
print("\n✅ Categorical Columns:", categorical_cols.tolist())
df_encoded = pd.get_dummies(df, drop_first=True)

# 7 Feature Scaling
from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()
X = df_encoded.drop('target', axis=1) # Replace 'target' with your actual target column
y = df_encoded['target']

X_scaled = scaler.fit_transform(X)

# 8 Train-Test Split
from sklearn.model_selection import train_test_split

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

# 9 Model Training
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report

model = RandomForestClassifier(random_state=42)
model.fit(X_train, y_train)

# 10 Predictions
y_pred = model.predict(X_test)

# Evaluation
print("\nAccuracy:", accuracy_score(y_test, y_pred))
print("\nConfusion Matrix:\n", confusion_matrix(y_test, y_pred))
print("\nClassification Report:\n", classification_report(y_test, y_pred))

# Predicting a new patient
# Example: Replace with actual input values
new_patient = {
    'age': 55,
    'sex': 1,
    'cp': 3,
    'trestbps': 140,
    'chol': 250,
    'fbs': 0,
    'restecg': 1,
    'thalach': 150,
    'exang': 0,
    'oldpeak': 2.3,
    'slope': 0,
    'ca': 0,
    'thal': 2
}

# Convert input to DataFrame
new_df = pd.DataFrame([new_patient])

# Combine with original data to ensure same columns
df_temp = pd.concat([df.drop(target, axis=1), new_df], ignore_index=True)
df_temp_encoded = pd.get_dummies(df_temp, drop_first=True)

# Reindex to match training columns
df_temp_encoded = df_temp_encoded.reindex(columns=df_encoded.drop(target, axis=1).columns, fill_value=0)

# Scale new input
new_input_scaled = scaler.transform(df_temp_encoded.tail(1))

# Make prediction
predicted_disease = model.predict(new_input_scaled)

print("\nPredicted Disease Presence (1=Yes, 0=No):", predicted_disease[0])

```



First 5 rows of the dataset:

	age	sex	chest_pain_type	resting_blood_pressure	cholesterol	\
0	52	Male	Typical angina	125	212	
1	53	Male	Typical angina	140	203	
2	70	Male	Typical angina	145	174	
3	61	Male	Typical angina	148	203	
4	62	Female	Typical angina	138	294	

	fasting_blood_sugar	rest_ecg	Max_heart_rate	\
0	Lower than 120 mg/ml	ST-T wave abnormality	168	
1	Greater than 120 mg/ml	Normal	155	
2	Lower than 120 mg/ml	ST-T wave abnormality	125	
3	Lower than 120 mg/ml	ST-T wave abnormality	161	
4	Greater than 120 mg/ml	ST-T wave abnormality	106	

	exercise_induced_angina	oldpeak	slope	vessels_colored_by_flourosopy	\
0	No	1.0	Downsloping	Two	
1	Yes	3.1	Upsloping	Zero	
2	Yes	2.6	Upsloping	Zero	
3	No	0.0	Downsloping	One	
4	No	1.9	Flat	Three	

	thalassemia	target
0	Reversable Defect	0
1	Reversable Defect	0
2	Reversable Defect	0
3	Reversable Defect	0
4	Fixed Defect	0

Shape of the dataset: (1025, 14)

Columns: ['age', 'sex', 'chest_pain_type', 'resting_blood_pressure', 'cholesterol', 'fasting_blood_sugar', 'rest_ecg', 'Max_heart_rate', 'exercise_induced_angina', 'oldpeak', 'slope', 'vessels_colored_by_flourosopy', 'thalassemia', 'target']

Data Types & Missing Values:

<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	object
2	chest_pain_type	1025 non-null	object
3	resting_blood_pressure	1025 non-null	int64
4	cholesterol	1025 non-null	int64
5	fasting_blood_sugar	1025 non-null	object
6	rest_ecg	1025 non-null	object
7	Max_heart_rate	1025 non-null	int64
8	exercise_induced_angina	1025 non-null	object
9	oldpeak	1025 non-null	float64
10	slope	1025 non-null	object
11	vessels_colored_by_flourosopy	1025 non-null	object
12	thalassemia	1025 non-null	object
13	target	1025 non-null	int64

dtypes: float64(1), int64(5), object(8)

memory usage: 112.2+ KB

Summary Statistics:

	age	resting_blood_pressure	cholesterol	Max_heart_rate	\
count	1025.000000	1025.000000	1025.000000	1025.000000	
mean	54.434146	131.611707	246.000000	149.114146	
std	9.072290	17.516718	51.59251	23.005724	
min	29.000000	94.000000	126.000000	71.000000	
25%	48.000000	120.000000	211.000000	132.000000	
50%	56.000000	130.000000	240.000000	152.000000	
75%	61.000000	140.000000	275.000000	166.000000	
max	77.000000	200.000000	564.000000	202.000000	

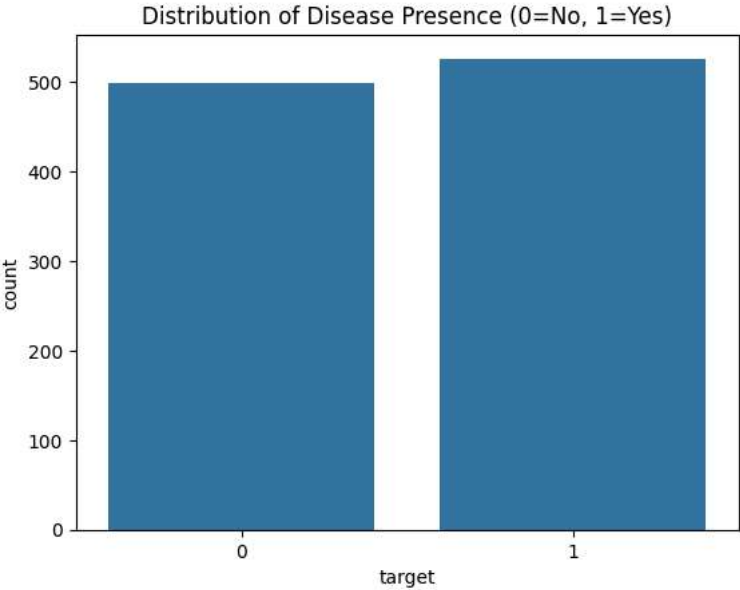
	oldpeak	target
count	1025.000000	1025.000000
mean	1.071512	0.513171
std	1.175053	0.500070
min	0.000000	0.000000
25%	0.000000	0.000000
50%	0.800000	1.000000
75%	1.800000	1.000000
max	6.200000	1.000000

Missing values per column:

Column	Missing Values
age	0
sex	0
chest_pain_type	0
resting_blood_pressure	0
cholesterol	0
fasting_blood_sugar	0
rest_ecg	0
Max_heart_rate	0
exercise_induced_angina	0
oldpeak	0
slope	0
vessels_colored_by_flourosopy	0
thalassemia	0
target	0

```
fasting_blood_sugar      0
rest_ecg                 0
Max_heart_rate           0
exercise_induced_angina  0
oldpeak                  0
slope                    0
vessels_colored_by_flourosopy  0
thalassemia              0
target                   0
dtype: int64
```

Duplicate rows: 723



Features: ['age', 'sex', 'chest_pain_type', 'resting_blood_pressure', 'cholestorl', 'fasting_blood_sugar', 'rest_ecg', 'Max_heart_ra
Categorical Columns: ['sex', 'chest_pain_type', 'fasting_blood_sugar', 'rest_ecg', 'exercise_induced_angina', 'slope', 'vessels_color

```
Missing values per column:
age      0
sex      0
chest_pain_type  0
resting_blood_pressure  0
cholestorl  0
fasting_blood_sugar  0
rest_ecg  0
Max_heart_rate  0
exercise_induced_angina  0
oldpeak  0
slope  0
vessels_colored_by_flourosopy  0
thalassemia  0
target  0
dtype: int64
```

- Duplicate rows: 723
- ✓ Duplicates removed.
- ✓ Outliers capped for 'age'
- ✓ Outliers capped for 'resting_blood_pressure'
- ✓ Outliers capped for 'cholestorl'
- ✓ Outliers capped for 'Max_heart_rate'
- ✓ Outliers capped for 'oldpeak'
- ✓ Outliers capped for 'target'
- ✓ Categorical Columns: ['sex', 'chest_pain_type', 'fasting_blood_sugar', 'rest_ecg', 'exercise_induced_angina', 'slope', 'vessels_cc

Accuracy: 0.8360655737704918

```
Confusion Matrix:
[[24  8]
 [ 2 27]]
```

Classification Report:

	precision	recall	f1-score	support
0	0.92	0.75	0.83	32
1	0.77	0.93	0.84	29
accuracy			0.84	61
macro avg	0.85	0.84	0.84	61

weighted avg 0.85 0.84 0.84 61

Predicted Disease Presence (1=Yes, 0=No): 1



```
!pip install gradio
```

```
Requirement already satisfied: gradio in /usr/local/lib/python3.11/dist-packages (5.29.0)
Requirement already satisfied: aiofiles<25.0,>=22.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (24.1.0)
Requirement already satisfied: anyio<5.0,>=3.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (4.9.0)
Requirement already satisfied: fastapi<1.0,>=0.115.2 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.115.12)
Requirement already satisfied: ffmpy in /usr/local/lib/python3.11/dist-packages (from gradio) (0.5.0)
Requirement already satisfied: gradio-client==1.10.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (1.10.0)
Requirement already satisfied: groovy~=0.1 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.1.2)
Requirement already satisfied: httpx>=0.24.1 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.28.1)
Requirement already satisfied: huggingface-hub>=0.28.1 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.31.1)
Requirement already satisfied: jinja2<4.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (3.1.6)
Requirement already satisfied: markupsafe<4.0,>=2.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (3.0.2)
Requirement already satisfied: numpy<3.0,>=1.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (2.0.2)
Requirement already satisfied: orjson==3.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (3.10.18)
Requirement already satisfied: packaging in /usr/local/lib/python3.11/dist-packages (from gradio) (24.2)
Requirement already satisfied: pandas<3.0,>=1.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (2.2.2)
Requirement already satisfied: pillow<12.0,>=8.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (11.2.1)
Requirement already satisfied: pydantic<2.12,>=2.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (2.11.4)
Requirement already satisfied: pydub in /usr/local/lib/python3.11/dist-packages (from gradio) (0.25.1)
Requirement already satisfied: python-multipart>=0.0.18 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.0.20)
Requirement already satisfied: pyyaml<7.0,>=5.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (6.0.2)
Requirement already satisfied: ruff>=0.9.3 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.11.9)
Requirement already satisfied: safehttpx<0.2.0,>=0.1.6 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.1.6)
Requirement already satisfied: semantic-version~=2.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (2.10.0)
Requirement already satisfied: starlette<1.0,>=0.40.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.46.2)
Requirement already satisfied: tomkit<0.14.0,>=0.12.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.13.2)
Requirement already satisfied: typer<1.0,>=0.12 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.15.3)
Requirement already satisfied: typing-extensions~=4.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (4.13.2)
Requirement already satisfied: uvicorn>=0.14.0 in /usr/local/lib/python3.11/dist-packages (from gradio) (0.34.2)
Requirement already satisfied: fsspec in /usr/local/lib/python3.11/dist-packages (from gradio-client==1.10.0->gradio) (2025.3.2)
Requirement already satisfied: websockets<16.0,>=10.0 in /usr/local/lib/python3.11/dist-packages (from gradio-client==1.10.0->gradio) (10.7)
Requirement already satisfied: idna>=2.8 in /usr/local/lib/python3.11/dist-packages (from anyio<5.0,>=3.0->gradio) (3.10)
Requirement already satisfied: sniffio>=1.1 in /usr/local/lib/python3.11/dist-packages (from anyio<5.0,>=3.0->gradio) (1.3.1)
Requirement already satisfied: certifi in /usr/local/lib/python3.11/dist-packages (from httpx>=0.24.1->gradio) (2025.4.26)
Requirement already satisfied: httpcore==1.* in /usr/local/lib/python3.11/dist-packages (from httpx>=0.24.1->gradio) (1.0.9)
Requirement already satisfied: h11>=0.16 in /usr/local/lib/python3.11/dist-packages (from httpcore==1.*->httpx>=0.24.1->gradio) (0.16.0)
Requirement already satisfied: filelock in /usr/local/lib/python3.11/dist-packages (from huggingface-hub>=0.28.1->gradio) (3.18.0)
Requirement already satisfied: requests in /usr/local/lib/python3.11/dist-packages (from huggingface-hub>=0.28.1->gradio) (2.32.3)
Requirement already satisfied: tqdm>=4.42.1 in /usr/local/lib/python3.11/dist-packages (from huggingface-hub>=0.28.1->gradio) (4.67.1)
Requirement already satisfied: hf-xet<2.0.0,>=1.1.0 in /usr/local/lib/python3.11/dist-packages (from huggingface-hub>=0.28.1->gradio) (1.1.1)
Requirement already satisfied: python-dateutil>=2.8.2 in /usr/local/lib/python3.11/dist-packages (from pandas<3.0,>=1.0->gradio) (2.9.0)
Requirement already satisfied: pytz>=2020.1 in /usr/local/lib/python3.11/dist-packages (from pandas<3.0,>=1.0->gradio) (2025.2)
Requirement already satisfied: tzdata>=2022.7 in /usr/local/lib/python3.11/dist-packages (from pandas<3.0,>=1.0->gradio) (2025.2)
Requirement already satisfied: annotated-types>=0.6.0 in /usr/local/lib/python3.11/dist-packages (from pydantic<2.12,>=2.0->gradio) (0.7.0)
Requirement already satisfied: pydantic-core==2.33.2 in /usr/local/lib/python3.11/dist-packages (from pydantic<2.12,>=2.0->gradio) (2.33.2)
Requirement already satisfied: typing-inspection>=0.4.0 in /usr/local/lib/python3.11/dist-packages (from pydantic<2.12,>=2.0->gradio) (0.12.1)
Requirement already satisfied: click>=8.0.0 in /usr/local/lib/python3.11/dist-packages (from typer<1.0,>=0.12->gradio) (8.1.8)
Requirement already satisfied: shellingham>=1.3.0 in /usr/local/lib/python3.11/dist-packages (from typer<1.0,>=0.12->gradio) (1.5.4)
Requirement already satisfied: rich>=10.11.0 in /usr/local/lib/python3.11/dist-packages (from typer<1.0,>=0.12->gradio) (13.9.4)
Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.11/dist-packages (from python-dateutil>=2.8.2->pandas<3.0,>=1.0->gradio) (1.17.0)
Requirement already satisfied: markdown-it-py>=2.2.0 in /usr/local/lib/python3.11/dist-packages (from rich>=10.11.0->typer<1.0,>=0.12->gradio) (3.0.0)
Requirement already satisfied: pygments<3.0.0,>=2.13.0 in /usr/local/lib/python3.11/dist-packages (from rich>=10.11.0->typer<1.0,>=0.12->gradio) (2.18.0)
Requirement already satisfied: charset-normalizer<4,>=2 in /usr/local/lib/python3.11/dist-packages (from requests->huggingface-hub>=0.28.1->gradio) (3.4.0)
Requirement already satisfied: urllib3<3,>=1.21.1 in /usr/local/lib/python3.11/dist-packages (from requests->huggingface-hub>=0.28.1->gradio) (2.2.3)
Requirement already satisfied: mdurl~0.1 in /usr/local/lib/python3.11/dist-packages (from markdown-it-py>=2.2.0->rich>=10.11.0->typer<1.0,>=0.12->gradio) (0.1.2)
```

```
import gradio as gr
import joblib
import pandas as pd

# Load the trained model
#model = joblib.load("heart_disease_model.pkl")

# Prediction function
def predict_heart_disease(age, sex, chest_pain_type, resting_blood_pressure, cholestoral,
                          fasting_blood_sugar, rest_ecg, max_heart_rate, exercise_induced_angina,
                          oldpeak, slope, vessels_colored_by_fluoroscopy, thalassemia):

    input_data = pd.DataFrame({
        "age": [age],
        "sex": [sex],
        "chest_pain_type": [chest_pain_type],
        "resting_blood_pressure": [resting_blood_pressure],
        "cholestoral": [cholestoral],
        "fasting_blood_sugar": [fasting_blood_sugar],
        "rest_ecg": [rest_ecg],
        "Max_heart_rate": [max_heart_rate],
        "exercise_induced_angina": [exercise_induced_angina],
```

```
"oldpeak": [oldpeak],
"slope": [slope],
"vessels_colored_by_flourosopy": [vessels_colored_by_flourosopy],
"thalassemia": [thalassemia]
})

prediction = model.predict(input_data)[0]
return "🔴 High Risk of Heart Disease" if prediction == 1 else "🟢 Low Risk of Heart Disease"

# Gradio interface
demo = gr.Interface(
    fn=predict_heart_disease,
    inputs=[
```