

Heart Disease UCI – Exploratory Data Analysis (EDA)

MLOps Experimental Learning Assignment

This notebook covers:

- Dataset acquisition
- Data cleaning & preprocessing
- Exploratory Data Analysis with professional visualizations
- EDA checklist aligned with production ML requirements

```
In [55]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

sns.set_theme(style="whitegrid")
```

1. Data Acquisition

Dataset source: UCI Machine Learning Repository – Heart Disease Dataset

```
In [56]: df = pd.read_csv("../data/heart.csv")
df.head()
```

```
Out[56]:
```

| | age | sex | cp | trestbps | chol | fbs | restecg | thalach | exang | oldpeak | slope | ca | thal |
|---|------|-----|-----|----------|-------|-----|---------|---------|-------|---------|-------|-----|------|
| 0 | 63.0 | 1.0 | 1.0 | 145.0 | 233.0 | 1.0 | 2.0 | 150.0 | 0.0 | 2.3 | 3.0 | 0.0 | 6.0 |
| 1 | 67.0 | 1.0 | 4.0 | 160.0 | 286.0 | 0.0 | 2.0 | 108.0 | 1.0 | 1.5 | 2.0 | 3.0 | 3.0 |
| 2 | 67.0 | 1.0 | 4.0 | 120.0 | 229.0 | 0.0 | 2.0 | 129.0 | 1.0 | 2.6 | 2.0 | 2.0 | 7.0 |
| 3 | 37.0 | 1.0 | 3.0 | 130.0 | 250.0 | 0.0 | 0.0 | 187.0 | 0.0 | 3.5 | 3.0 | 0.0 | 3.0 |
| 4 | 41.0 | 0.0 | 2.0 | 130.0 | 204.0 | 0.0 | 2.0 | 172.0 | 0.0 | 1.4 | 1.0 | 0.0 | 3.0 |

2. Basic Dataset Inspection

```
In [57]: df.shape
```

```
Out[57]: (303, 14)
```

In [58]: `df.info()`

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):
#   Column      Non-Null Count  Dtype
---  -
0   age         303 non-null   float64
1   sex         303 non-null   float64
2   cp          303 non-null   float64
3   trestbps    303 non-null   float64
4   chol        303 non-null   float64
5   fbs         303 non-null   float64
6   restecg     303 non-null   float64
7   thalach     303 non-null   float64
8   exang       303 non-null   float64
9   oldpeak     303 non-null   float64
10  slope       303 non-null   float64
11  ca          303 non-null   object
12  thal        303 non-null   object
13  target      303 non-null   int64
dtypes: float64(11), int64(1), object(2)
memory usage: 33.3+ KB
```

In [59]: `df.describe()`

Out[59]:

| | age | sex | cp | trestbps | chol | fbs | restecg |
|-------|------------|------------|------------|------------|------------|------------|------------|
| count | 303.000000 | 303.000000 | 303.000000 | 303.000000 | 303.000000 | 303.000000 | 303.000000 |
| mean | 54.438944 | 0.679868 | 3.158416 | 131.689769 | 246.693069 | 0.148515 | 0.990099 |
| std | 9.038662 | 0.467299 | 0.960126 | 17.599748 | 51.776918 | 0.356198 | 0.994971 |
| min | 29.000000 | 0.000000 | 1.000000 | 94.000000 | 126.000000 | 0.000000 | 0.000000 |
| 25% | 48.000000 | 0.000000 | 3.000000 | 120.000000 | 211.000000 | 0.000000 | 0.000000 |
| 50% | 56.000000 | 1.000000 | 3.000000 | 130.000000 | 241.000000 | 0.000000 | 1.000000 |
| 75% | 61.000000 | 1.000000 | 4.000000 | 140.000000 | 275.000000 | 0.000000 | 2.000000 |
| max | 77.000000 | 1.000000 | 4.000000 | 200.000000 | 564.000000 | 1.000000 | 2.000000 |

3. Data Quality Checks

In [60]: `df.isnull().sum()`

```
Out[60]: age      0
sex      0
cp      0
trestbps 0
chol     0
fbs      0
restecg  0
thalach  0
exang    0
oldpeak  0
slope    0
ca       0
thal     0
target   0
dtype: int64
```

```
In [61]: df.duplicated().sum()
```

```
Out[61]: np.int64(0)
```

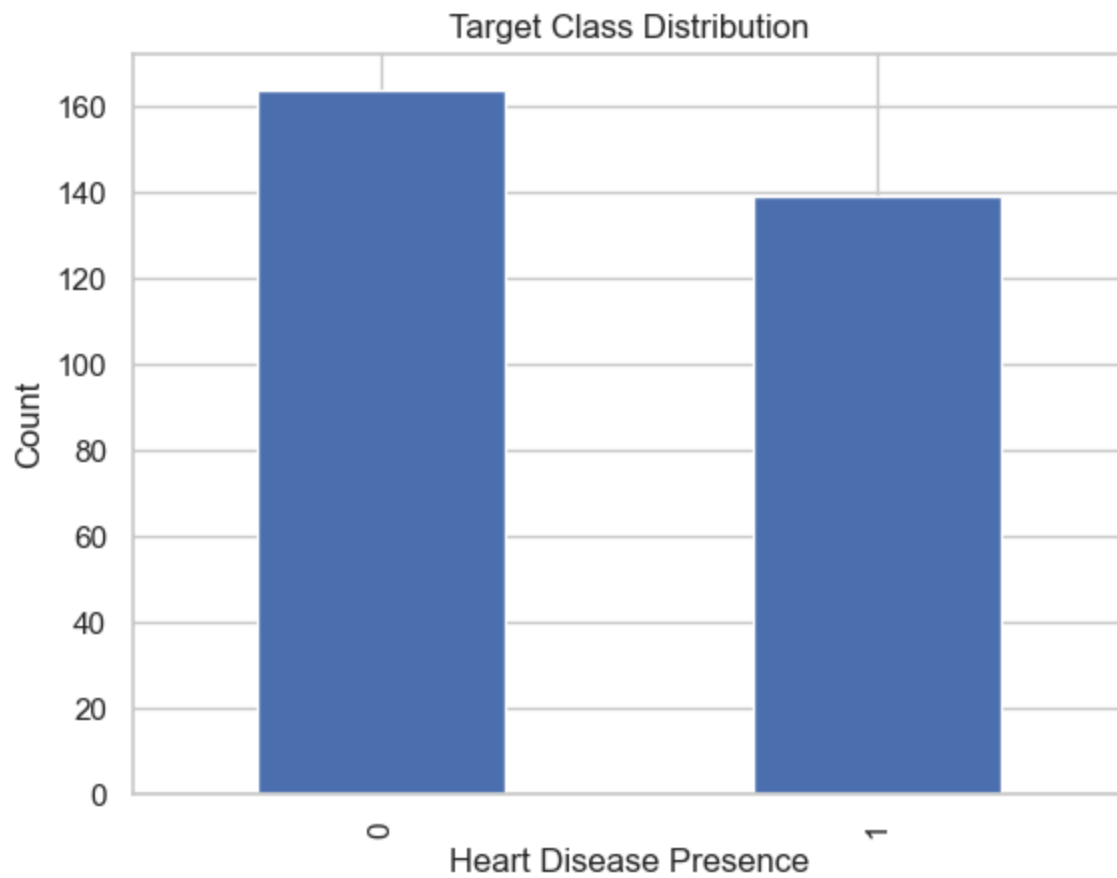
4. Target Variable Count

```
In [62]: df["target"] = (df["target"] > 0).astype(int)
df["target"].value_counts()
```

```
Out[62]: target
0      164
1      139
Name: count, dtype: int64
```

5. Target Variable Distribution

```
In [63]: plt.figure()
df['target'].value_counts().plot(kind='bar')
plt.title("Target Class Distribution")
plt.xlabel("Heart Disease Presence")
plt.ylabel("Count")
plt.show()
```



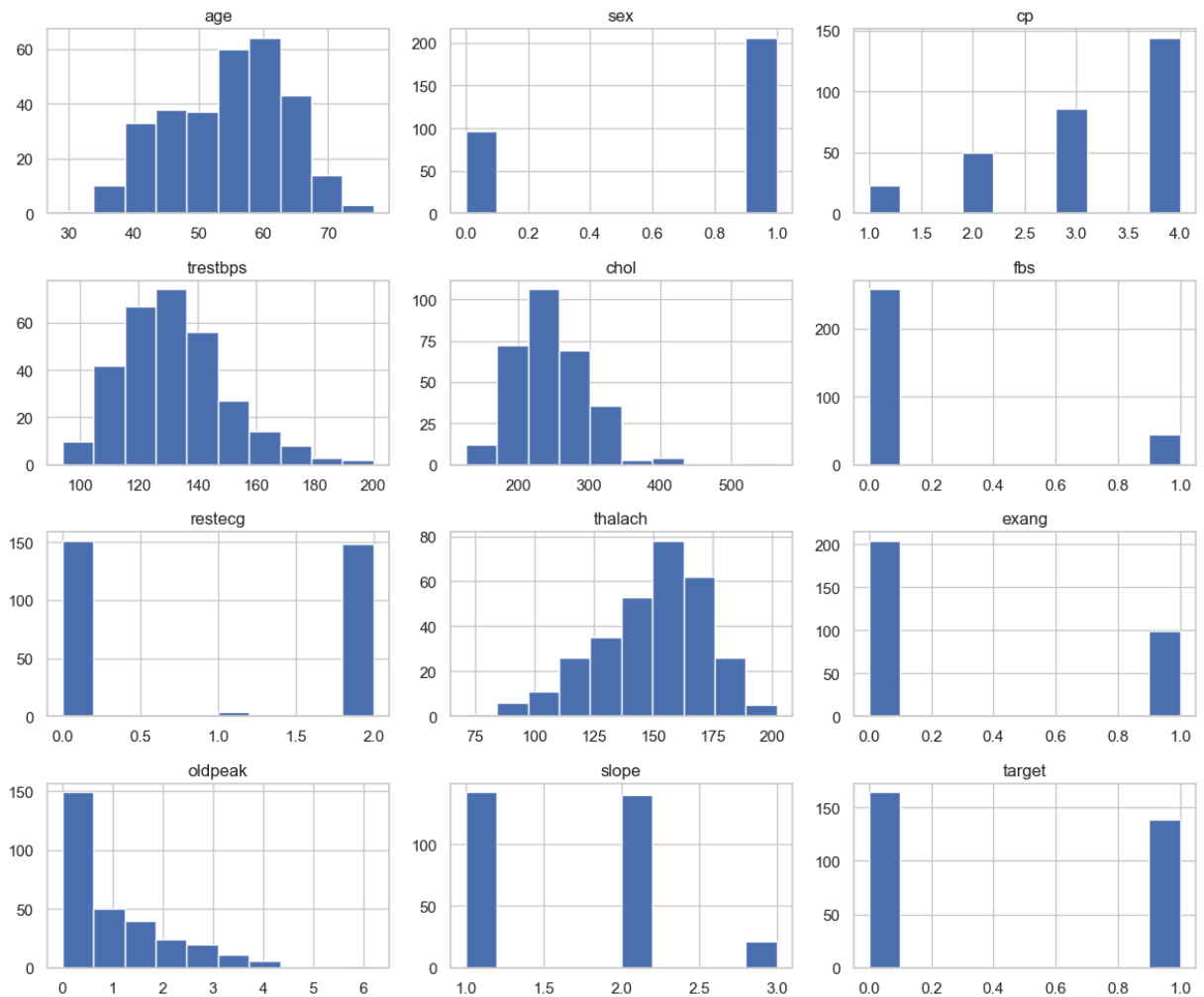
6. Class Balance Plot

```
In [64]: sns.countplot(x="target", data=df)
plt.title("Target Class Distribution")
plt.show()
```



7. Numerical Feature Distributions

```
In [65]: df.hist(figsize=(12,10))  
plt.tight_layout()  
plt.show()
```



8. Data Cleaning for Correlation Analysis

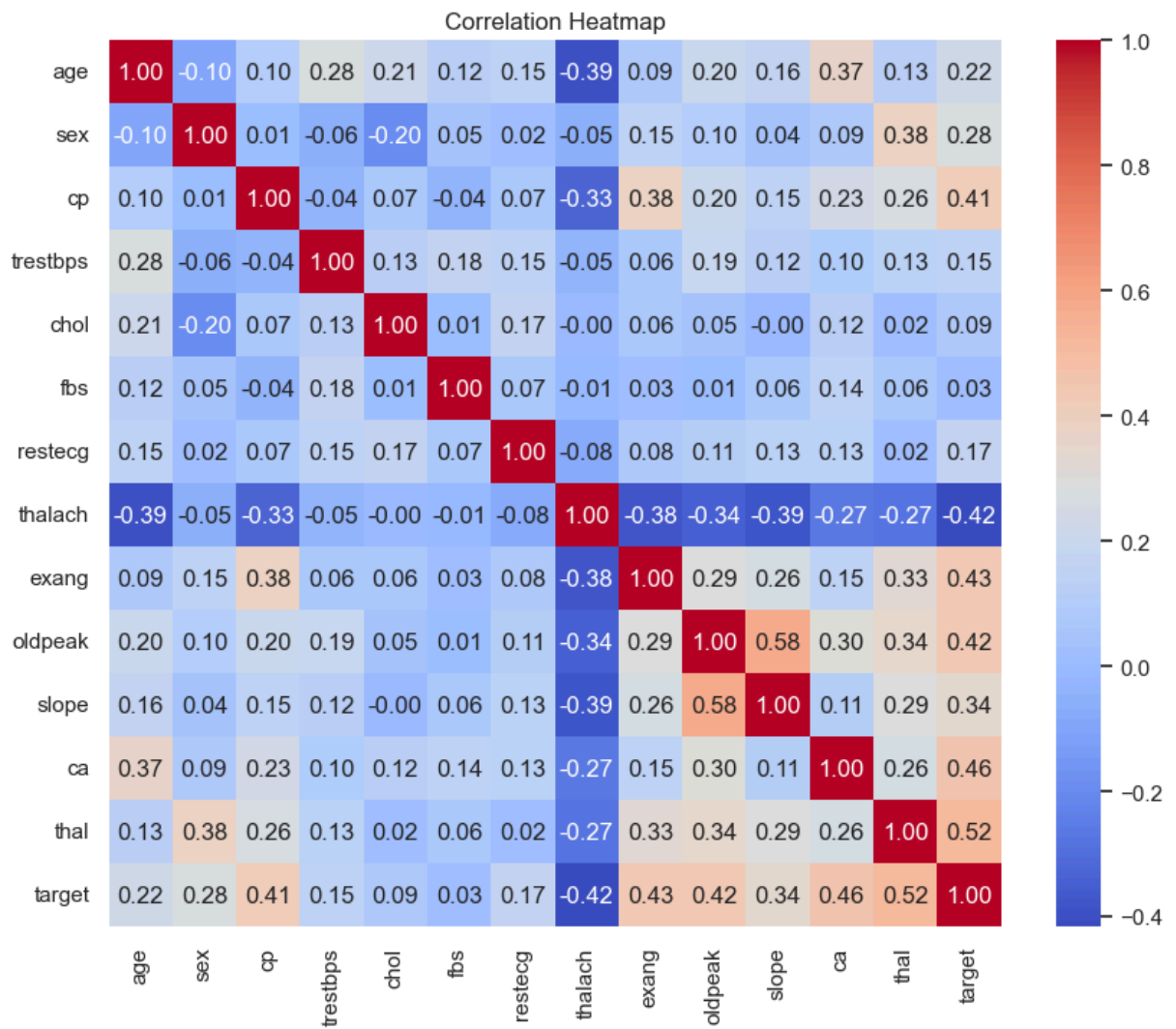
```
In [66]: # Replace invalid placeholders with NaN
df.replace('?', np.nan, inplace=True)

# Convert affected columns to numeric
df['ca'] = pd.to_numeric(df['ca'])
df['thal'] = pd.to_numeric(df['thal'])

# Impute missing values using median
df['ca'] = df['ca'].fillna(df['ca'].median())
df['thal'] = df['thal'].fillna(df['thal'].median())
```

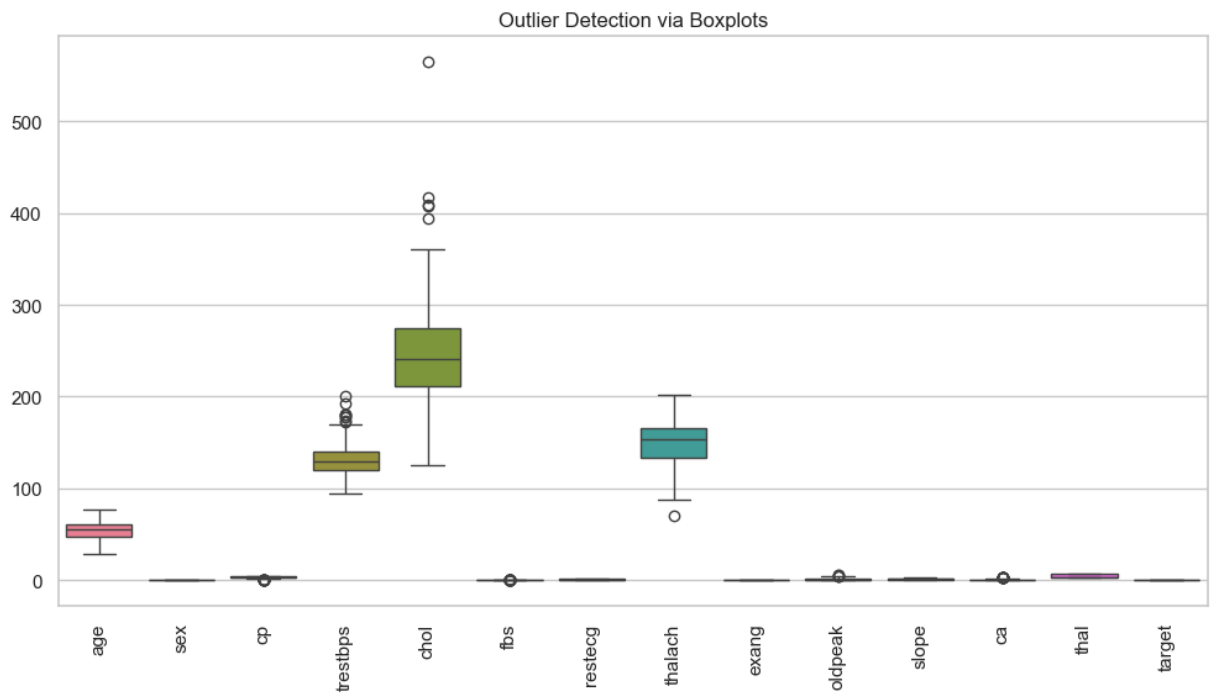
9. Correlation Analysis

```
In [67]: plt.figure(figsize=(10,8))
sns.heatmap(df.corr(), annot=True, cmap="coolwarm", fmt=".2f")
plt.title("Correlation Heatmap")
plt.show()
```



10. Outlier Detection

```
In [68]: plt.figure(figsize=(12,6))
sns.boxplot(data=df)
plt.xticks(rotation=90)
plt.title("Outlier Detection via Boxplots")
plt.show()
```



11. Key EDA Observations

- Dataset is clean with minimal missing values
- Target variable is fairly balanced
- Some features show moderate correlation with target
- Outliers are medically plausible