


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

```
# Upload the dataset
uploaded = files.upload()
```

```
# Load the dataset
df = pd.read_csv('heart.csv') # Adjust the filename if different
```

 heart.csv

- **heart.csv**(text/csv) - 38114 bytes, last modified: 22/08/2024 - 100% done


Saving heart.csv to heart (8).csv

```
import numpy as np # Make sure NumPy is imported
```

```
# Drop rows with missing values
df.dropna(inplace=True)
```

```
# Handle outliers (example: removing rows with cholesterol beyond 3 standard deviations)
if 'cholesterol' in df.columns:
    df = df[(np.abs(df['cholesterol'] - df['cholesterol'].mean()) <= (3 * df['cholesterol'].std()))]
else:
    print("The 'cholesterol' column does not exist in the dataframe.")
```

```
# Check the first few rows after cleaning
print(df.head())
```

 The 'cholesterol' column does not exist in the dataframe.

	age	sex	cp	trestbps	chol	fb	restecg	thalach	exang	oldpeak	slope	\
0	52	1	0	125	212	0	1	168	0	1.0	2	
1	53	1	0	140	203	1	0	155	1	3.1	0	
2	70	1	0	145	174	0	1	125	1	2.6	0	
3	61	1	0	148	203	0	1	161	0	0.0	2	
4	62	0	0	138	294	1	1	106	0	1.9	1	

	ca	thal	target
0	2	3	0
1	0	3	0
2	0	3	0
3	1	3	0
4	3	2	0

```
# Summary statistics
print(df.describe())
```

```
# Distribution of data (Histograms)
df.hist(figsize=(12, 10))
plt.show()
```

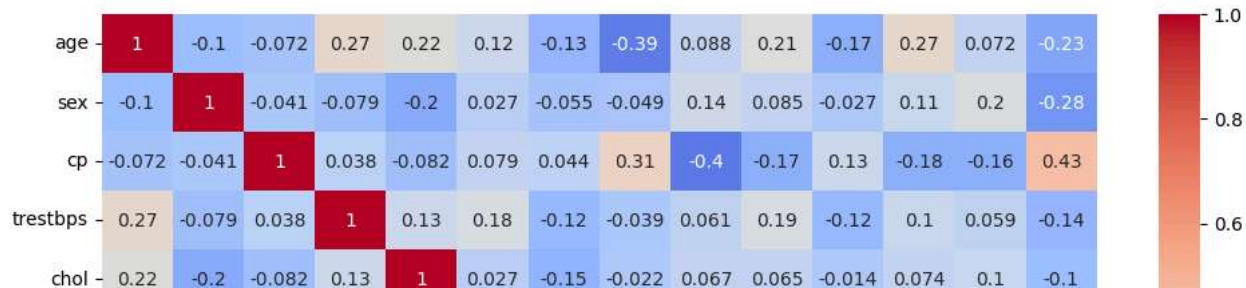
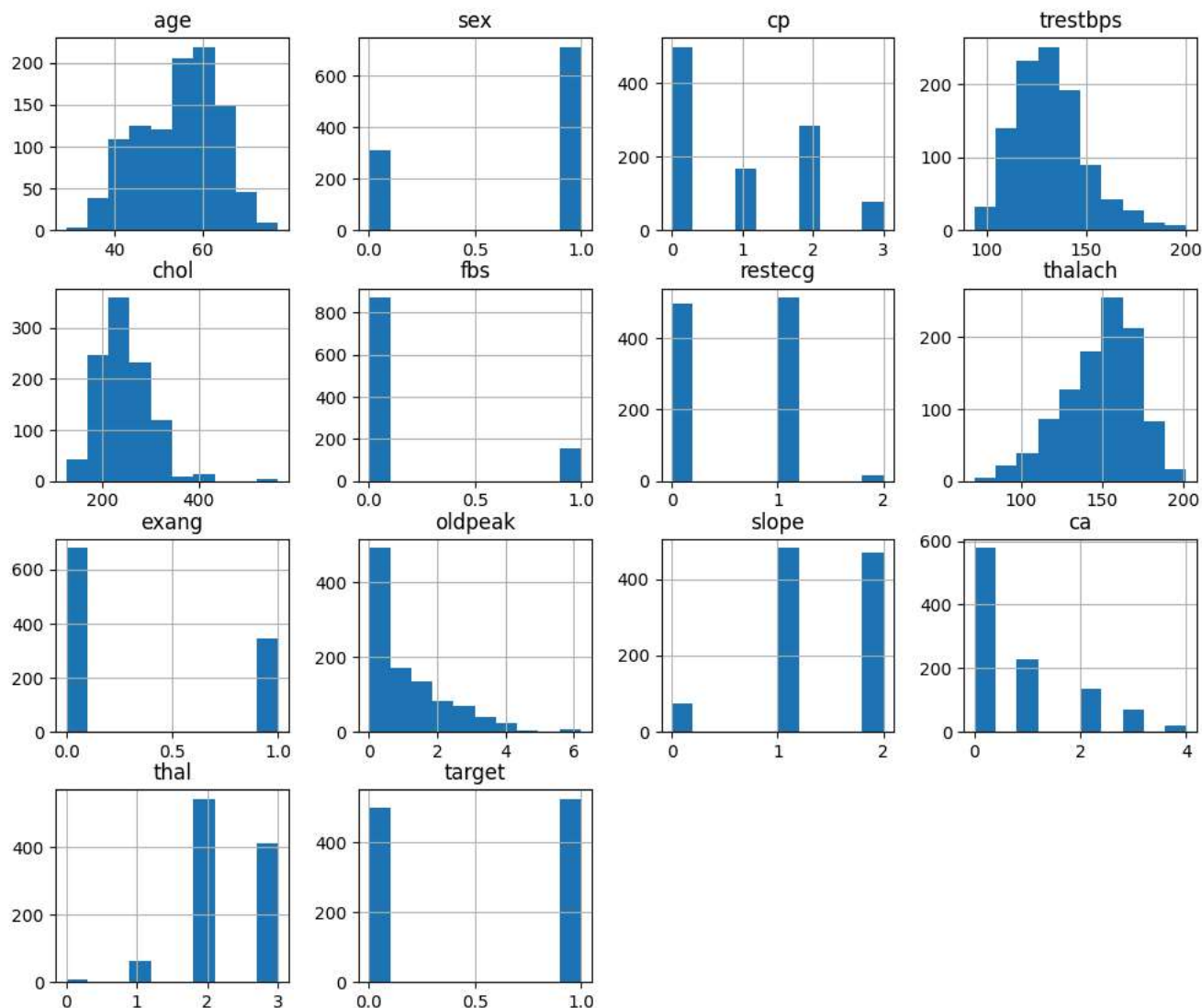
```
# Correlation matrix
plt.figure(figsize=(12, 8))
sns.heatmap(df.corr(), annot=True, cmap='coolwarm')
plt.show()
```

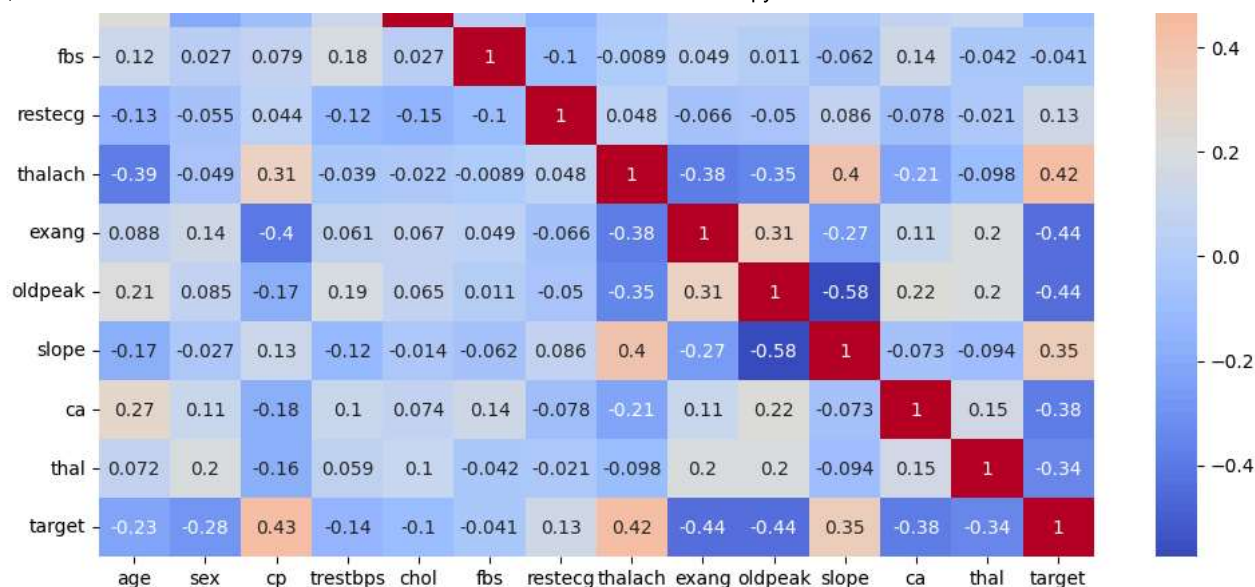


count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	54.434146	0.695610	0.942439	131.611707	246.000000
std	9.072290	0.460373	1.029641	17.516718	51.59251
min	29.000000	0.000000	0.000000	94.000000	126.00000
25%	48.000000	0.000000	0.000000	120.000000	211.00000
50%	56.000000	1.000000	1.000000	130.000000	240.00000
75%	61.000000	1.000000	2.000000	140.000000	275.00000
max	77.000000	1.000000	3.000000	200.000000	564.00000

	fbs	restecg	thalach	exang	oldpeak \
count	1025.000000	1025.000000	1025.000000	1025.000000	1025.000000
mean	0.149268	0.529756	149.114146	0.336585	1.071512
std	0.356527	0.527878	23.005724	0.472772	1.175053
min	0.000000	0.000000	71.000000	0.000000	0.000000
25%	0.000000	0.000000	132.000000	0.000000	0.000000
50%	0.000000	1.000000	152.000000	0.000000	0.800000
75%	0.000000	1.000000	166.000000	1.000000	1.800000
max	1.000000	2.000000	202.000000	1.000000	6.200000

	slope	ca	thal	target
count	1025.000000	1025.000000	1025.000000	1025.000000
mean	1.385366	0.754146	2.323902	0.513171
std	0.617755	1.030798	0.620660	0.500070
min	0.000000	0.000000	0.000000	0.000000
25%	1.000000	0.000000	2.000000	0.000000
50%	1.000000	0.000000	2.000000	1.000000
75%	2.000000	1.000000	3.000000	1.000000
max	2.000000	4.000000	3.000000	1.000000





```
# Check if the dataframe is loaded correctly and inspect the first few rows
print("First few rows of the dataframe:")
print(df.head())

# Check if the columns 'heart_disease' and 'age' exist in the dataframe
print("\nColumn Names in DataFrame:")
print(df.columns)

# If both 'heart_disease' and 'age' columns exist, check for missing values
if 'heart_disease' in df.columns and 'age' in df.columns:
    print("\nMissing Values in 'heart_disease' and 'age' Columns:")
    print(df[['heart_disease', 'age']].isnull().sum())
else:
    print("\nError: One or both of the columns 'heart_disease' and 'age' do not exist in the dataframe.")
```

```
First few rows of the dataframe:
   age  sex  cp  trestbps  chol  fbs  restecg  thalach  exang  oldpeak  slope  \
0   52   1   0    125    212   0         1    168      0      1.0      2
1   53   1   0    140    203   1         0    155      1      3.1      0
2   70   1   0    145    174   0         1    125      1      2.6      0
3   61   1   0    148    203   0         1    161      0      0.0      2
4   62   0   0    138    294   1         1    106      0      1.9      1
```

```
   ca  thal  target
0   2     3      0
1   0     3      0
2   0     3      0
3   1     3      0
4   3     2      0
```

```
Column Names in DataFrame:
Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
       'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
      dtype='object')
```

```
Error: One or both of the columns 'heart_disease' and 'age' do not exist in the dataframe.
```

```
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    print("\nMissing Values in 'heart_disease' and 'age' Columns:")
    print(df[['heart_disease', 'age']].isnull().sum())

# Pairplot to visualize relationships between variables
sns.pairplot(df, hue='heart_disease')
plt.show()

# Heatmap of correlation matrix with a focus on variables related to heart disease
plt.figure(figsize=(10, 6))
sns.heatmap(df.corr(), annot=True, cmap='coolwarm', center=0)
plt.title('Correlation Heatmap')
```