```
import pandas as pd

# Load the dataset
df = pd.read_csv('/content/heart_disease_dataset.csv')

# Display the first 5 rows
print("First 5 rows of the dataset:")
display(df.head())

# Display column information (data types and non-null counts)
print("\nColumn information:")
display(df.info())
```

	age	sex	ср	trtbps	chol	fbs	restecg	thalachh	exng	oldpeak	slp	caa	thal
0	63	1	3	145	233	1	0	150	0	2.3	0	0	
1	37	1	2	130	250	0	1	187	0	3.5	0	0	
2	41	0	1	130	204	0	0	172	0	1.4	2	0	
3	56	1	1	120	236	0	1	178	0	0.8	2	0	

1

163

1

0.6

2

0

0

Column information:

0 0

57

First 5 rows of the dataset:

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):

120 354

#	Column	Non-Null Count	Dtype
0	age	303 non-null	int64
1	sex	303 non-null	int64
2	ср	303 non-null	int64
3	trtbps	303 non-null	int64
4	chol	303 non-null	int64
5	fbs	303 non-null	int64
6	restecg	303 non-null	int64
7	thalachh	303 non-null	int64
8	exng	303 non-null	int64
9	oldpeak	303 non-null	float64
10	slp	303 non-null	int64
11	caa	303 non-null	int64
12	thall	303 non-null	int64
13	output	303 non-null	int64
dtyp	es: float6	4(1), int64(13)	

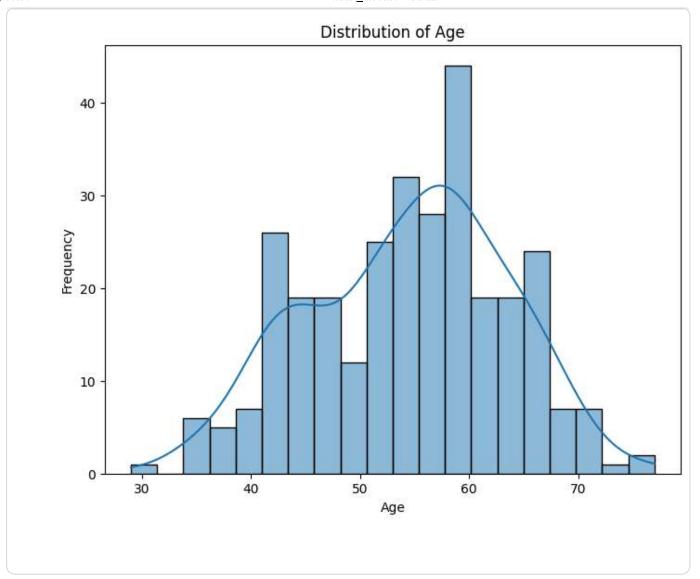
atypes: +10at64(1), 1nt64(13)

memory usage: 33.3 KB

None

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

# Plot a histogram of the 'Age' column
plt.figure(figsize=(8, 6))
sns.histplot(df['age'], bins=20, kde=True)
plt.title('Distribution of Age')
plt.xlabel('Age')
plt.ylabel('Frequency')
plt.show()
```



```
import pandas as pd

# Load the dataset
df = pd.read_csv('/content/heart_disease_dataset.csv')

# Display the first 5 rows
print("First 5 rows of the dataset:")
display(df.head())

# Display column information (data types and non-null counts)
print("\nColumn information:")
display(df.info())
```

First	5	rows	of	the	dat	aset:
-------	---	------	----	-----	-----	-------

	age	sex	ср	trtbps	chol	fbs	restecg	thalachh	exng	oldpeak	slp	caa	thal
(	63	1	3	145	233	1	0	150	0	2.3	0	0	
1	I 37	1	2	130	250	0	1	187	0	3.5	0	0	
2	2 41	0	1	130	204	0	0	172	0	1.4	2	0	
3	<b>3</b> 56	1	1	120	236	0	1	178	0	0.8	2	0	
4	<b>.</b> 57	0	0	120	354	0	1	163	1	0.6	2	0	

Column information:

<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	int64
1	sex	303 non-null	int64
2	ср	303 non-null	int64
3	trtbps	303 non-null	int64
4	chol	303 non-null	int64
5	fbs	303 non-null	int64
6	restecg	303 non-null	int64
7	thalachh	303 non-null	int64
8	exng	303 non-null	int64
9	oldpeak	303 non-null	float64
10	slp	303 non-null	int64
11	caa	303 non-null	int64
12	thall	303 non-null	int64
13	output	303 non-null	int64
dtyp	es: float6	4(1), int64(13)	

atypes:  $\pm 10at64(1)$ , 1nt64(13)

memory usage: 33.3 KB

None

```
# Check for missing values
print("Missing values before handling:")
display(df.isnull().sum())

# Verify that missing values have been handled
print("\nMissing values after handling:")
display(df.isnull().sum())
```

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## sns.scatterplot

## seaborn.relational.scatterplot

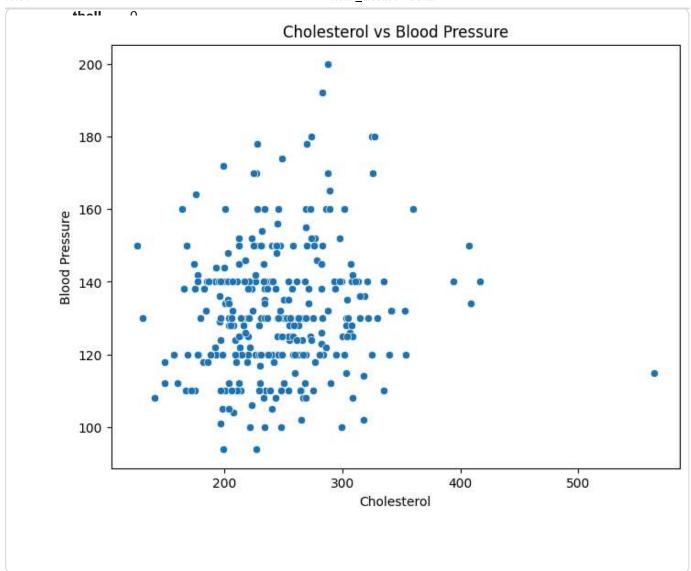
def scatterplot(data=None, \*, x=None, y=None, hue=None, size=None, style=None,
palette=None, hue\_order=None, hue\_norm=None, sizes=None, size\_order=None,
size\_norm=None, markers=True, style\_order=None, legend='auto', ax=None,
\*\*kwargs)

 ${f restecg} = 0$  Draw a scatter plot with possibility of several semantic groupings.

## thalachh 0

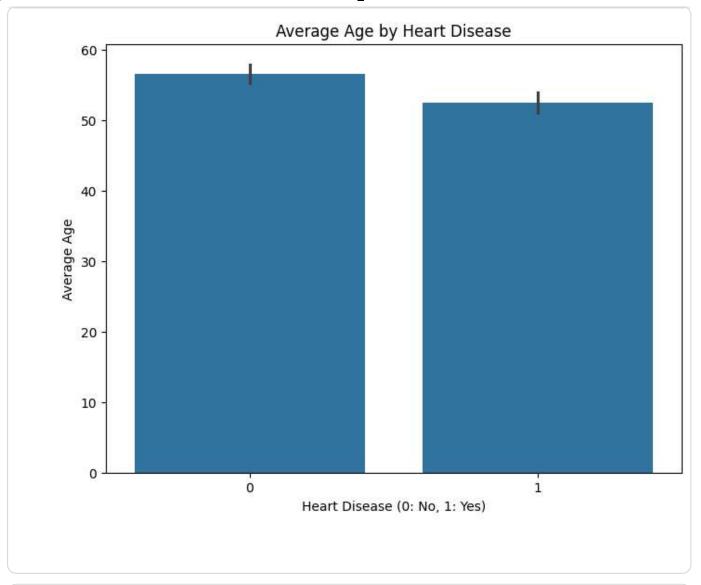
The relationship between `x` and `y` can be shown for different subsets of the control what visual semantics are used to identify the different oldweak. Of it is possible to show up to three dimensions independently by

```
import matplotlib.pyplot as plt
import seaborn as sns
# Create a scatterplot of 'Cholesterol' vs 'Blood Pressure'
plt.figure(figsize=(8, 6))
sns.scatterplot(x='chol', y='trtbps', data=df)
plt.title('Cholesterol vs Blood Pressure')
plt.xlabel('Cholesterol')
plt.ylabel('Blood Pressure')
plt.show()
          0
   age
   sex
          0
          0
   Ср
  trtbps
          0
  chol
          0
          0
   fbs
 restecq
          0
thalachh 0
          0
  exng
 oldpeak
          0
   slp
          0
          0
   caa
```



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

# Create a barplot of average age by heart disease
plt.figure(figsize=(8, 6))
sns.barplot(x='output', y='age', data=df)
plt.title('Average Age by Heart Disease')
plt.xlabel('Heart Disease (0: No, 1: Yes)')
plt.ylabel('Average Age')
plt.show()
```



```
# Identify object type columns
print("Object type columns:")
for col in df.columns:
   if df[col].dtype == 'object':
      print(col)

Object type columns:
```

```
# Identify object type columns
# In this dataset, all columns are numerical as seen from df.info()
# There are no categorical columns to encode based on the current dataframe struct
# If there were object type columns, we would identify them here and decide on encode
```

```
# For this dataset, we can skip the encoding step as there are no object columns.

print("No object type columns to encode in this dataset based on df.info().")

# If you had object columns and wanted to define encoding methods, you would do it # encoding_methods = {col: 'One-Hot Encoding' for col in categorical_cols} # print("Chosen encoding methods for each categorical column:")

# for col, method in encoding_methods.items():

# print(f"{col}: {method}")

No object type columns to encode in this dataset based on df.info().
```

```
# import from sklearn.preprocessing import OneHotEncoder
# Based on the analysis, there are no object type columns to encode in this datase
# Therefore, the one-hot encoding step is not necessary for this dataset.
# The dataframe 'df' is already in a suitable format for modeling.
# If you had categorical columns to encode, the code would be as follows:
# categorical cols = list(encoding methods.keys())
# encoder = OneHotEncoder(sparse_output=False, handle_unknown='ignore')
# encoded data = encoder.fit transform(df[categorical cols])
# encoded_df = pd.DataFrame(encoded_data, columns=encoder.get_feature_names_out(ca
# df = pd.concat([df.drop(columns=categorical cols), encoded df], axis=1)
# print("DataFrame after checking for one-hot encoding:")
# display(df.head())
# print("\nColumn information after checking for encoding:")
# display(df.info())
print("Skipping one-hot encoding as there are no object columns in the dataset.")
Skipping one-hot encoding as there are no object columns in the dataset.
```

```
#
      encoder = OneHotEncoder(sparse output=False, handle unknown='ignore')
      encoded_data = encoder.fit_transform(df[categorical_cols])
#
      encoded df = pd.DataFrame(encoded data, columns=encoder.get feature names ou
#
      df = pd.concat([df.drop(columns=categorical_cols), encoded_df], axis=1)
#
      print("DataFrame after one-hot encoding:")
#
      display(df.head())
#
#
      print("\nColumn information after encoding:")
      display(df.info())
#
# else:
      print("DataFrame not loaded, cannot proceed with encoding.")
print("Skipping one-hot encoding as there are no object columns in the dataset.")
Skipping one-hot encoding as there are no object columns in the dataset.
```

```
import pandas as pd
# from sklearn.preprocessing import OneHotEncoder
# Load the dataset again (This cell seems redundant as df is already loaded)
# try:
      df = pd.read_csv('/content/heart_disease_dataset.csv')
# except FileNotFoundError:
      print("Error: heart disease dataset.csv not found. Please make sure the file
      # Indicate failure if the file is not found
#
      df = None
# Based on the analysis, there are no object type columns to encode in this datase
# Therefore, the one-hot encoding step is not necessary for this dataset.
# The dataframe 'df' is already in a suitable format for modeling.
# if df is not None:
      categorical_cols = list(encoding_methods.keys()) # Need to ensure encoding_n
      # Create a OneHotEncoder instance
#
      encoder = OneHotEncoder(sparse output=False, handle unknown='ignore')
#
#
      # Fit and transform the selected categorical columns
#
      encoded_data = encoder.fit_transform(df[categorical_cols])
#
      # Create a new DataFrame from the encoded data
#
      encoded_df = pd.DataFrame(encoded_data, columns=encoder.get_feature_names_ou
#
      # Concatenate the new encoded DataFrame with the original DataFrame, droppir
      df = pd.concat([df.drop(columns=categorical cols), encoded df], axis=1)
#
      # Display the first few rows of the updated DataFrame
#
      print("DataFrame after one-hot encoding:")
#
#
      display(df.head())
```

```
# # Display column information to verify the new columns and data types
# print("\nColumn information after encoding:")
# display(df.info())
# else:
# print("DataFrame not loaded, cannot proceed with encoding.")

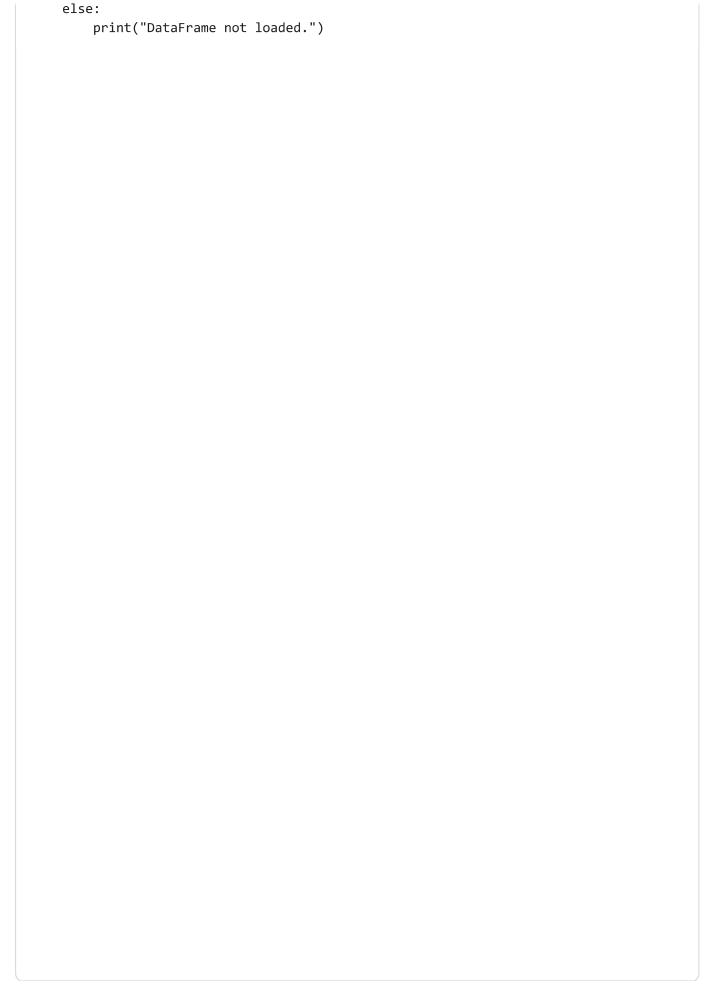
print("Skipping one-hot encoding as there are no object columns in the dataset.")

Skipping one-hot encoding as there are no object columns in the dataset.
```

```
# from sklearn.preprocessing import OneHotEncoder
# Based on the analysis, there are no object type columns to encode in this datase
# Therefore, the one-hot encoding step is not necessary for this dataset.
# The dataframe 'df' is already in a suitable format for modeling.
# if 'encoding methods' in locals():
      categorical_cols = list(encoding_methods.keys())
# else:
      categorical cols = [] # Or identify based on df.columns if needed
# if categorical cols:
#
      # Create a OneHotEncoder instance
      encoder = OneHotEncoder(sparse_output=False, handle_unknown='ignore')
#
      # Fit and transform the selected categorical columns
#
      encoded data = encoder.fit transform(df[categorical cols])
#
      # Create a new DataFrame from the encoded data
#
#
      encoded_df = pd.DataFrame(encoded_data, columns=encoder.get_feature_names_ou
      # Concatenate the new encoded DataFrame with the original DataFrame, droppir
#
      df = pd.concat([df.drop(columns=categorical_cols), encoded_df], axis=1)
#
#
      # Display the first few rows of the updated DataFrame
      print("DataFrame after one-hot encoding:")
#
      display(df.head())
#
      # Display column information to verify the new columns and data types
#
#
      print("\nColumn information after encoding:")
      display(df.info())
#
# else:
      print("No categorical columns to encode.")
print("Skipping one-hot encoding as there are no object columns in the dataset.")
Skipping one-hot encoding as there are no object columns in the dataset.
```

```
import pandas as pd
# from sklearn.preprocessing import OneHotEncoder
```

```
# Load the dataset
try:
    df = pd.read csv('/content/heart disease dataset.csv')
except FileNotFoundError:
    print("Error: heart disease dataset.csv not found. Please make sure the file j
    df = None
if df is not None:
    # Check for missing values
    print("Missing values before handling:")
    display(df.isnull().sum())
   # Based on the dataset info, there are no missing values and no object columns
    # Removing the missing value handling for 'Alcohol Intake' as it's not in the
    # Removing the encoding part as there are no object columns.
   # Identify object type columns - this will be empty based on df.info()
    categorical cols = [col for col in df.columns if df[col].dtype == 'object']
    if categorical cols:
        # Ensure encoding_methods dictionary is defined, or define it based on ide
        if 'encoding_methods' not in locals():
            encoding methods = {col: 'One-Hot Encoding' for col in categorical col
        # Create a OneHotEncoder instance
        encoder = OneHotEncoder(sparse_output=False, handle_unknown='ignore')
        # Fit and transform the selected categorical columns
        encoded_data = encoder.fit_transform(df[categorical_cols])
        # Create a new DataFrame from the encoded data
        encoded_df = pd.DataFrame(encoded_data, columns=encoder.get_feature_names_
        # Concatenate the new encoded DataFrame with the original DataFrame, dropp
        df = pd.concat([df.drop(columns=categorical cols), encoded df], axis=1)
        print("DataFrame after one-hot encoding:")
        display(df.head())
        print("\nColumn information after encoding:")
        display(df.info())
    else:
        print("No object type columns to encode.")
        print("DataFrame is ready for further analysis/modeling.")
        # Display the first few rows of the DataFrame
        print("DataFrame head:")
        display(df.head())
        print("\nColumn information:")
        display(df.info())
```

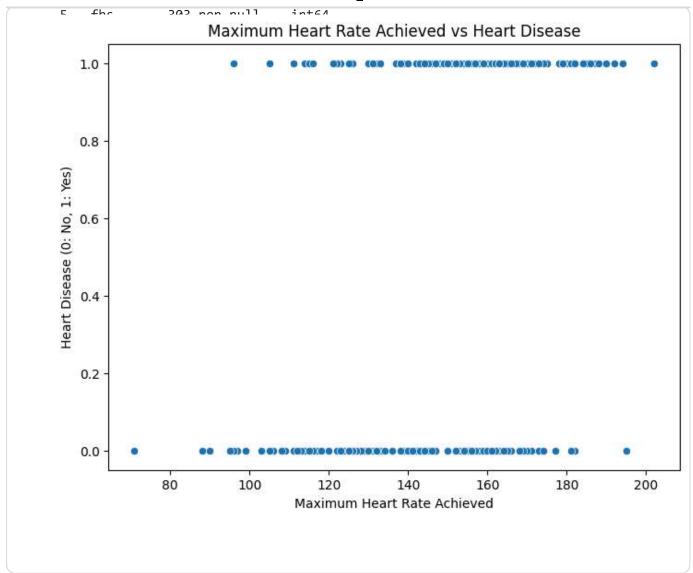


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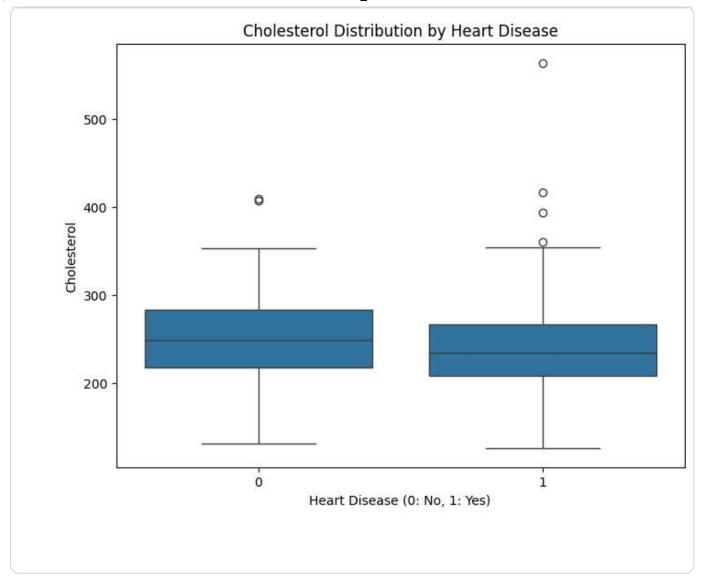
```
# Display column information to verify the new columns and data types
print("\nColumn information after encoding:")
display(df.info())
```

```
CoPXMP information after encoding:
RangeIndex: 303 entries, 0 to 302
Data Golumns (total 14 columns):
     Column
               Non-Null Count Dtype
---caa----მ
               303 non-null
                                int64
 0
     age
 1 thallex
         0
               303 non-null
                                int64
               303 non-null
                                int64
 3output pps
               303 non-null
                                int64
               303 non-null
     chol
                                int64
dtype:first64
               303 non-null
                                int64
N6 objectetgpe 201uman-tulencodent64
DataFthmeathreadd for-fulther anatysis/modeling.
DataFeams head:303 non-null
                                int64
             cp trtbps chol
303 non-null
                                float64
fbs restecg thalachh exng oldpeak slp
     oldpeak
                                                                             caa tha
   age
Slp
                                in‡64
               303 nqn5null
                                                                   2.3
                                           0
                                                    150
                                                           0
                                                                          0
                                                                               0
 12
               303 non-null
                                int64
    thall
13 37 protput
               2303 nqrgnu1250
                                int 64
                                           1
                                                    187
                                                           0
                                                                   3.5
                                                                          0
                                                                               0
dtypes: float64(1), int64(13)
m2mor4/1 usagle: 313.3 KE 30
                                  0
                                           0
                                                    172
                                                           0
                                                                   1.4
                                                                          2
                                                                               0
None
     56
                     120
                           236
                                           1
                                                   178
                                                           0
                                                                          2
                                                                               0
3
                                  0
                                                                   8.0
```

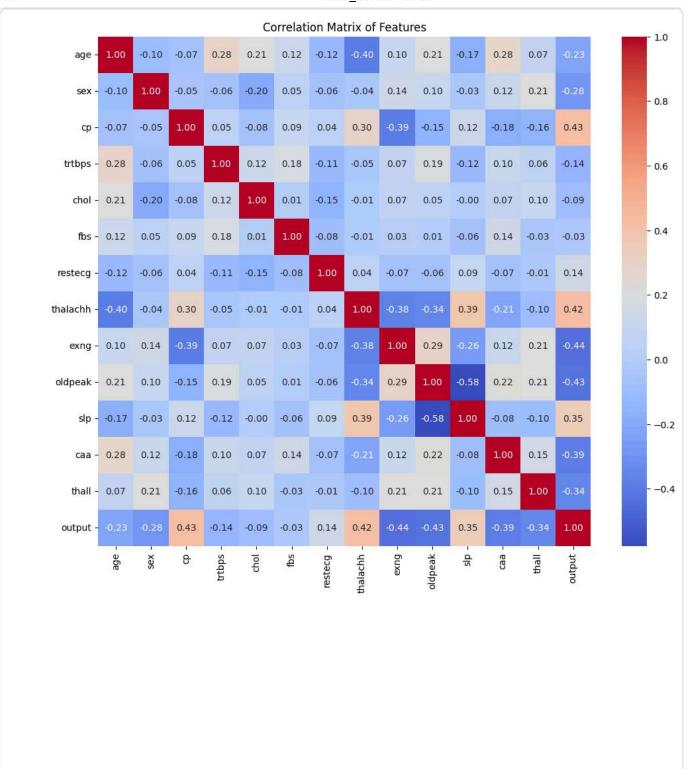
```
plt.figure(figsize=(8, 6))
sns.scatterplot(x='thalachh', y='output', data=df)
plt.title('Maximum Heart Rate Achieved vs Heart Disease')
plt.xlabel('Maximum Heart Rate Achieved')
plt.ylabel('Heart Disease (0: No, 1: Yes)')
plt.show()
0
               303 non-null
                                int64
     age
1
     sex
               303 non-null
                                int64
 2
               303 non-null
                                int64
     ср
 3
               303 non-null
                                int64
     trtbps
     chol
               303 non-null
                                int64
```



```
plt.figure(figsize=(8, 6))
sns.boxplot(x='output', y='chol', data=df)
plt.title('Cholesterol Distribution by Heart Disease')
plt.xlabel('Heart Disease (0: No, 1: Yes)')
plt.ylabel('Cholesterol')
plt.show()
```



```
plt.figure(figsize=(12, 10))
  correlation_matrix = df.corr()
  sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', fmt=".2f")
  plt.title('Correlation Matrix of Features')
  plt.show()
```



from sklearn.model\_selection import train\_test\_split
X = df.drop('output', axis=1)

```
y = df['output']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_s1
print("Shape of X_train:", X_train.shape)
print("Shape of X test:", X test.shape)
print("Shape of y_train:", y_train.shape)
print("Shape of y_test:", y_test.shape)
Shape of X train: (242, 13)
Shape of X_test: (61, 13)
Shape of y train: (242,)
Shape of y_test: (61,)
from sklearn.linear model import LogisticRegression
# Instantiate the model
model = LogisticRegression(max iter=1000)
# Train the model
model.fit(X_train, y_train)
print("Model training completed.")
Model training completed.
from sklearn.metrics import accuracy score, precision score, recall score, f1 scor
# Make predictions on the test set
y_pred = model.predict(X_test)
# Calculate evaluation metrics
accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred)
recall = recall_score(y_test, y_pred)
f1 = f1_score(y_test, y_pred)
# Print the evaluation metrics
print(f"Accuracy: {accuracy:.4f}")
print(f"Precision: {precision:.4f}")
print(f"Recall: {recall:.4f}")
print(f"F1-score: {f1:.4f}")
Accuracy: 0.8033
Precision: 0.7692
Recall: 0.9091
F1-score: 0.8333
```

```
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy score, precision score, recall score, f1 scor
# Instantiate the RandomForestClassifier model
rf_model = RandomForestClassifier(random_state=42)
# Train the new model
rf_model.fit(X_train, y_train)
# Make predictions with the new model
y pred rf = rf model.predict(X test)
# Evaluate the new model
accuracy rf = accuracy score(y test, y pred rf)
precision_rf = precision_score(y_test, y_pred_rf)
recall rf = recall score(y test, y pred rf)
f1_rf = f1_score(y_test, y_pred_rf)
# Print the evaluation metrics for the new model
print("Random Forest Classifier Performance:")
print(f"Accuracy: {accuracy rf:.4f}")
print(f"Precision: {precision_rf:.4f}")
print(f"Recall: {recall rf:.4f}")
print(f"F1-score: {f1 rf:.4f}")
# Compare with Logistic Regression
print("\nComparison with Logistic Regression:")
print(f"Logistic Regression Accuracy: {accuracy:.4f}")
print(f"Random Forest Accuracy: {accuracy rf:.4f}")
print(f"Logistic Regression Precision: {precision:.4f}")
print(f"Random Forest Precision: {precision rf:.4f}")
print(f"Logistic Regression Recall: {recall:.4f}")
print(f"Random Forest Recall: {recall rf:.4f}")
print(f"Logistic Regression F1-score: {f1:.4f}")
print(f"Random Forest F1-score: {f1_rf:.4f}")
Random Forest Classifier Performance:
Accuracy: 0.8361
Precision: 0.7805
Recall: 0.9697
F1-score: 0.8649
Comparison with Logistic Regression:
Logistic Regression Accuracy: 0.8033
Random Forest Accuracy: 0.8361
Logistic Regression Precision: 0.7692
Random Forest Precision: 0.7805
Logistic Regression Recall: 0.9091
Random Forest Recall: 0.9697
Logistic Regression F1-score: 0.8333
Random Forest F1-score: 0.8649
```

```
from sklearn.preprocessing import StandardScaler

# Initialize the StandardScaler
scaler = StandardScaler()

# Fit the scaler on the training data and transform both training and testing data
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

print("Data after standard scaling:")
print("Shape of X_train_scaled:", X_train_scaled.shape)
print("Shape of X_test_scaled:", X_test_scaled.shape)

Data after standard scaling:
Shape of X_train_scaled: (242, 13)
Shape of X_test_scaled: (61, 13)
```

```
# Instantiate the Logistic Regression model
scaled_lr_model = LogisticRegression(max_iter=1000)

# Train the model on the scaled training data
scaled_lr_model.fit(X_train_scaled, y_train)

print("Logistic Regression model trained on scaled data.")

Logistic Regression model trained on scaled data.
```

```
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_scor
# Make predictions on the scaled test set
y_pred_scaled_lr = scaled_lr_model.predict(X_test_scaled)
# Calculate evaluation metrics for the scaled Logistic Regression model
accuracy scaled lr = accuracy score(y test, y pred scaled lr)
precision_scaled_lr = precision_score(y_test, y_pred_scaled_lr)
recall_scaled_lr = recall_score(y_test, y_pred_scaled_lr)
f1 scaled lr = f1 score(y test, y pred scaled lr)
# Print the evaluation metrics
print("Logistic Regression Model Performance on Scaled Data:")
print(f"Accuracy: {accuracy_scaled_lr:.4f}")
print(f"Precision: {precision scaled lr:.4f}")
print(f"Recall: {recall_scaled_lr:.4f}")
print(f"F1-score: {f1 scaled lr:.4f}")
Logistic Regression Model Performance on Scaled Data:
Accuracy: 0.8033
Precision: 0.7692
Recall: 0.9091
F1-score: 0.8333
```