```
In [1]: import numpy as np
        import pandas as pd
        import seaborn as sns
        import matplotlib.pyplot as plt
        import warnings
        warnings.filterwarnings('ignore')
In [2]: df = pd.read_csv('/content/heart.csv')
In [3]: df.head()
           Age Sex ChestPainType RestingBP Cholesterol FastingBS RestingECG
Out[3]:
        0
             40
                  Μ
                                ATA
                                           140
                                                        289
                                                                     0
                                                                             Normal
        1
             49
                                NAP
                                           160
                                                        180
                                                                     0
                                                                             Normal
        2
                                ATA
                                           130
                                                        283
                                                                     0
                                                                                 ST
            37
                  Μ
                                                                             Normal
        3
             48
                                ASY
                                           138
                                                        214
        4
                                                                     0
             54
                  Μ
                                NAP
                                           150
                                                        195
                                                                             Normal
        EDA
In [4]: df.columns
Out[4]: Index(['Age', 'Sex', 'ChestPainType', 'RestingBP', 'Cholesterol', 'FastingB
        S',
                'RestingECG', 'MaxHR', 'ExerciseAngina', 'Oldpeak', 'ST Slope',
                'HeartDisease'],
              dtype='object')
In [5]: df.shape
Out[5]: (918, 12)
```

In [6]: df.info()

<class 'pandas.core.frame.DataFrame'> RangeIndex: 918 entries, 0 to 917 Data columns (total 12 columns):

Ducu	cocamins (cocac	12 Cocamins / i					
#	Column	Non-Null Count	Dtype				
0	Age	918 non-null	int64				
1	Sex	918 non-null	object				
2	ChestPainType	918 non-null	object				
3	RestingBP	918 non-null	int64				
4	Cholesterol	918 non-null	int64				
5	FastingBS	918 non-null	int64				
6	RestingECG	918 non-null	object				
7	MaxHR	918 non-null	int64				
8	ExerciseAngina	918 non-null	object				
9	Oldpeak	918 non-null	float64				
10	ST_Slope	918 non-null	object				
11	HeartDisease	918 non-null	int64				
dtype	es: float64(1),	int64(6), object	(5)				

memory usage: 86.2+ KB

In [7]: df.describe()

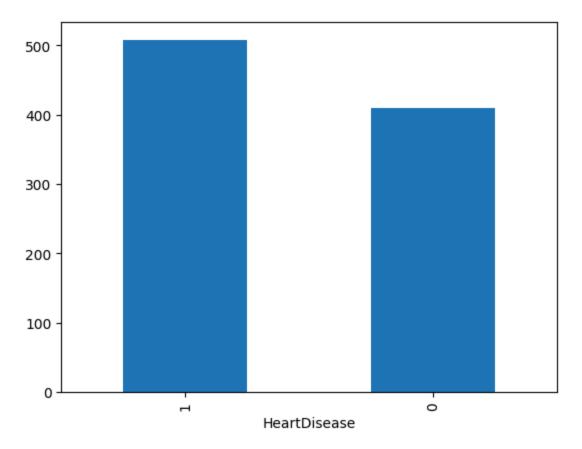
Out[7]:		Age	RestingBP	Cholesterol	FastingBS	MaxHR	Oldpeal
	count	918.000000	918.000000	918.000000	918.000000	918.000000	918.000000
	mean	53.510893	132.396514	198.799564	0.233115	136.809368	0.887364
	std	9.432617	18.514154	109.384145	0.423046	25.460334	1.066570
	min	28.000000	0.000000	0.000000	0.000000	60.000000	-2.600000
	25%	47.000000	120.000000	173.250000	0.000000	120.000000	0.000000
	<b>50</b> %	54.000000	130.000000	223.000000	0.000000	138.000000	0.600000
	<b>75</b> %	60.000000	140.000000	267.000000	0.000000	156.000000	1.500000
	max	77.000000	200.000000	603.000000	1.000000	202.000000	6.200000

```
In [8]: df.duplicated().sum()
```

Out[8]: np.int64(0)

```
In [9]: df['HeartDisease'].value_counts().plot(kind = 'bar')
```

Out[9]: <Axes: xlabel='HeartDisease'>



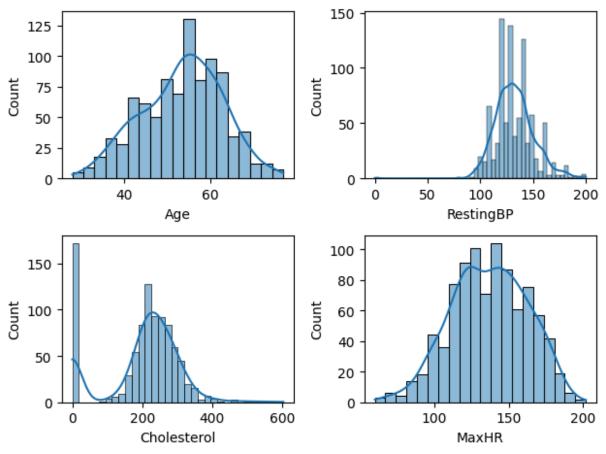
dtype: int64

```
In [11]: def plotting(var, num):
    plt.subplot(2,2,num)
```

```
sns.histplot(df[var], kde = True)

plotting('Age', 1)
plotting('RestingBP', 2)
plotting('Cholesterol', 3)
plotting('MaxHR', 4)

plt.tight_layout()
```



In [12]: df['Cholesterol'].value\_counts()

Out[12]: count

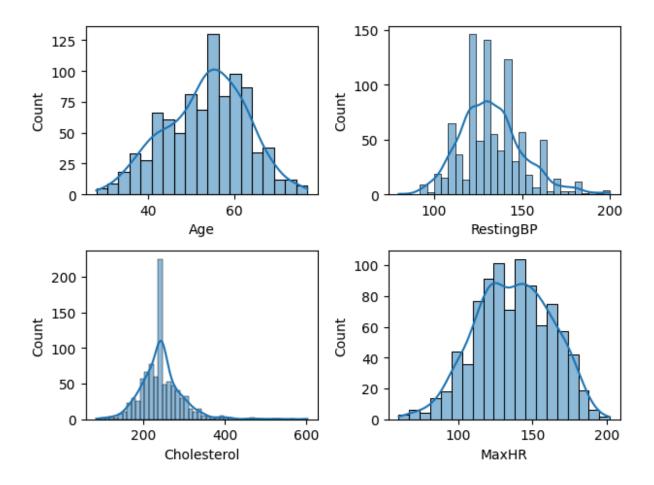
Cholesterol
-------------

0	172
254	11
220	10
223	10
204	9
353	1
278	1
157	1
176	1
131	1

222 rows × 1 columns

## dtype: int64

```
In [13]: ch_mean = df.loc[df['Cholesterol'] != 0, 'Cholesterol'].mean()
In [14]: ch mean
Out[14]: np.float64(244.6353887399464)
In [15]: df['Cholesterol'] = df['Cholesterol'].replace(0,ch_mean)
         df['Cholesterol'] = df['Cholesterol'].round(2)
In [16]: resting bp mean = df.loc[df['RestingBP'] != 0, 'RestingBP'].mean()
         df['RestingBP'] = df['RestingBP'].replace(0, resting bp mean)
         df['RestingBP'] = df['RestingBP'].round(2)
In [17]: def plotting(var, num):
           plt.subplot(2,2,num)
           sns.histplot(df[var], kde = True)
         plotting('Age', 1)
         plotting('RestingBP', 2)
         plotting('Cholesterol', 3)
         plotting('MaxHR', 4)
         plt.tight layout()
```



In [18]: pip install sheryanalysis==0.1.0

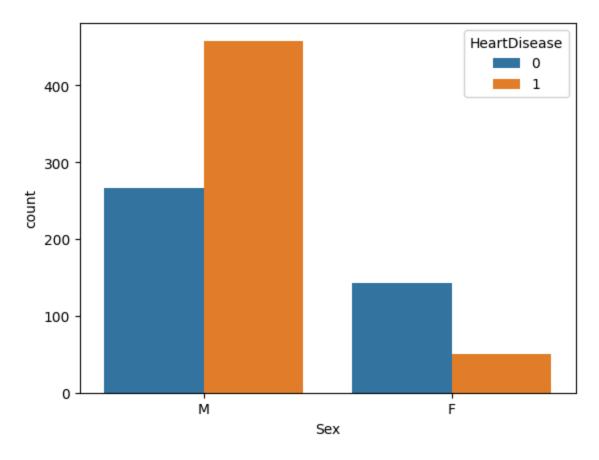
```
Collecting sheryanalysis==0.1.0
          Downloading sheryanalysis-0.1.0-py3-none-any.whl.metadata (574 bytes)
        Requirement already satisfied: pandas>=1.0.0 in /usr/local/lib/python3.11/di
        st-packages (from sheryanalysis==0.1.0) (2.2.2)
        Requirement already satisfied: numpy>=1.18.0 in /usr/local/lib/python3.11/di
        st-packages (from sheryanalysis==0.1.0) (2.0.2)
        Requirement already satisfied: scikit-learn>=0.22.0 in /usr/local/lib/python
        3.11/dist-packages (from sheryanalysis==0.1.0) (1.6.1)
        Requirement already satisfied: python-dateutil>=2.8.2 in /usr/local/lib/pyth
        on3.11/dist-packages (from pandas>=1.0.0->sheryanalysis==0.1.0) (2.9.0.post
        0)
        Requirement already satisfied: pytz>=2020.1 in /usr/local/lib/python3.11/dis
        t-packages (from pandas>=1.0.0->sheryanalysis==0.1.0) (2025.2)
        Requirement already satisfied: tzdata>=2022.7 in /usr/local/lib/python3.11/d
        ist-packages (from pandas>=1.0.0->sheryanalysis==0.1.0) (2025.2)
        Requirement already satisfied: scipy>=1.6.0 in /usr/local/lib/python3.11/dis
        t-packages (from scikit-learn>=0.22.0->sheryanalysis==0.1.0) (1.16.1)
        Requirement already satisfied: joblib>=1.2.0 in /usr/local/lib/python3.11/di
        st-packages (from scikit-learn>=0.22.0->sheryanalysis==0.1.0) (1.5.1)
        Requirement already satisfied: threadpoolctl>=3.1.0 in /usr/local/lib/python
        3.11/dist-packages (from scikit-learn>=0.22.0->sheryanalysis==0.1.0) (3.6.0)
        Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.11/dist-pa
        ckages (from python-dateutil>=2.8.2->pandas>=1.0.0->sheryanalysis==0.1.0)
        (1.17.0)
        Downloading sheryanalysis-0.1.0-py3-none-any.whl (10 kB)
        Installing collected packages: sheryanalysis
        Successfully installed sheryanalysis-0.1.0
In [19]: import sheryanalysis as sh
         sh.analyze(df)
        Basic Analysis Report
        INFO:sheryanalysis:
        Basic Analysis Report
        INFO:sheryanalysis:-----
        \ Shape: (918, 12)
        INFO:sheryanalysis: \Shape: (918, 12)
        Columns: ['Age', 'Sex', 'ChestPainType', 'RestingBP', 'Cholesterol', 'Fas
        tingBS', 'RestingECG', 'MaxHR', 'ExerciseAngina', 'Oldpeak', 'ST Slope', 'He
        artDisease']
        INFO:sheryanalysis: Columns: ['Age', 'Sex', 'ChestPainType', 'RestingBP',
        'Cholesterol', 'FastingBS', 'RestingECG', 'MaxHR', 'ExerciseAngina', 'Oldpea
        k', 'ST Slope', 'HeartDisease']

✓ No null values found

        INFO:sheryanalysis:
        ✓ No null values found
        Ell Categorical Columns: ['Sex', 'ChestPainType', 'FastingBS', 'RestingECG',
        'ExerciseAngina', 'ST Slope', 'HeartDisease']
        INFO:sheryanalysis:
        🔠 Categorical Columns: ['Sex', 'ChestPainType', 'FastingBS', 'RestingECG',
        'ExerciseAngina', 'ST_Slope', 'HeartDisease']
        🔢 Numerical Columns: ['Age', 'RestingBP', 'Cholesterol', 'MaxHR', 'Oldpea
        k']
```

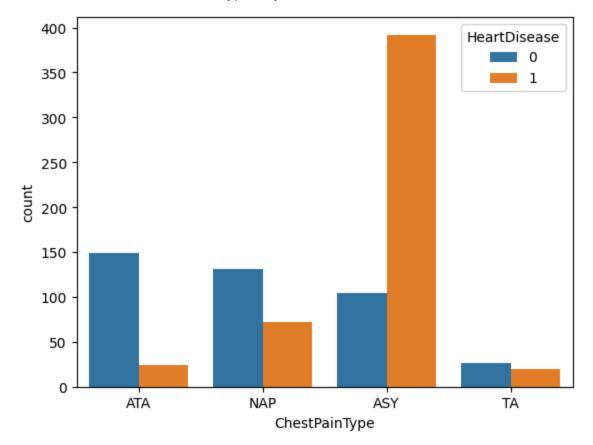
```
🔢 Numerical Columns: ['Age', 'RestingBP', 'Cholesterol', 'MaxHR', 'Oldpea
        k']
Out[19]: {'shape': (918, 12),
           'columns': ['Age',
            'Sex',
            'ChestPainType',
            'RestingBP',
            'Cholesterol',
            'FastingBS',
            'RestingECG',
            'MaxHR',
            'ExerciseAngina',
            'Oldpeak',
            'ST Slope',
            'HeartDisease'],
           'dtypes': {'Age': dtype('int64'),
            'Sex': dtype('0'),
            'ChestPainType': dtype('0'),
            'RestingBP': dtype('float64'),
            'Cholesterol': dtype('float64'),
            'FastingBS': dtype('int64'),
            'RestingECG': dtype('0'),
            'MaxHR': dtype('int64'),
            'ExerciseAngina': dtype('0'),
            'Oldpeak': dtype('float64'),
            'ST Slope': dtype('0'),
            'HeartDisease': dtype('int64')},
           'null_counts': {'Age': 0,
            'Sex': 0,
            'ChestPainType': 0,
            'RestingBP': 0,
            'Cholesterol': 0,
            'FastingBS': 0,
            'RestingECG': 0,
            'MaxHR': 0,
            'ExerciseAngina': 0,
            'Oldpeak': 0,
            'ST Slope': 0,
            'HeartDisease': 0},
           'total rows': 918,
           'column types': {'categorical': ['Sex',
             'ChestPainType',
             'FastingBS',
             'RestingECG',
             'ExerciseAngina',
             'ST Slope',
             'HeartDisease'],
            'numerical': ['Age', 'RestingBP', 'Cholesterol', 'MaxHR', 'Oldpeak'],
            'datetime': [],
            'text': []}}
In [20]: | sns.countplot(x = df['Sex'], hue = df['HeartDisease'])
Out[20]: <Axes: xlabel='Sex', ylabel='count'>
```

INFO:sheryanalysis:



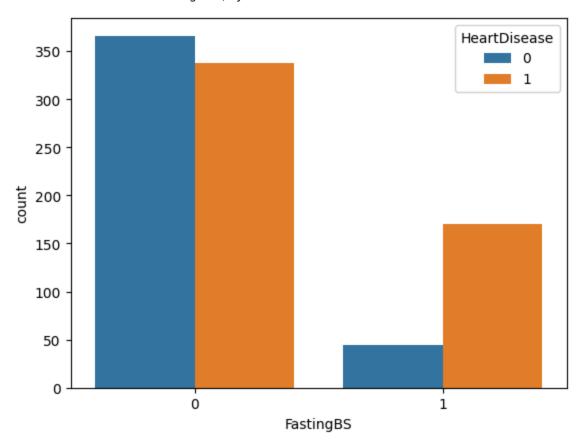
In [21]: sns.countplot(x = df['ChestPainType'], hue = df['HeartDisease'])

Out[21]: <Axes: xlabel='ChestPainType', ylabel='count'>



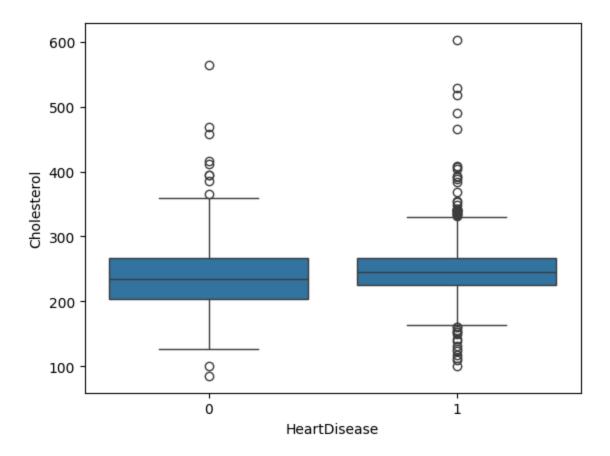
```
In [22]: sns.countplot(x = df['FastingBS'], hue = df['HeartDisease'])
```

Out[22]: <Axes: xlabel='FastingBS', ylabel='count'>

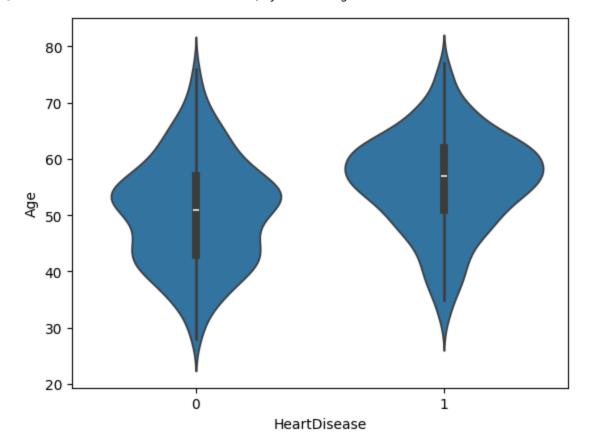


```
In [23]: sns.boxplot(x = 'HeartDisease', y = 'Cholesterol', data = df)
```

Out[23]: <Axes: xlabel='HeartDisease', ylabel='Cholesterol'>

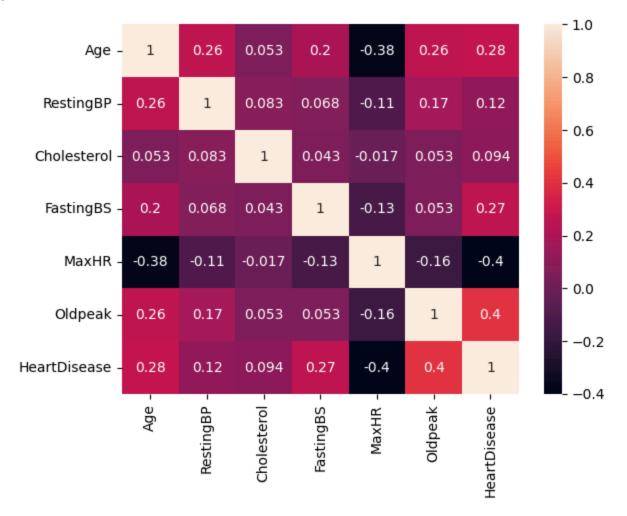


Out[24]: <Axes: xlabel='HeartDisease', ylabel='Age'>



In [25]: sns.heatmap(df.corr(numeric\_only = True), annot =True)

Out[25]: <Axes: >



## **Data Preprocessing and cleaning**

```
In [26]: df_encode = pd.get_dummies(df, drop_first = True)
    df_encode
```

Out[26]:		Age	RestingBP	Cholesterol	FastingBS	MaxHR	Oldpeak	HeartDisease
	0	40	140.0	289.0	0	172	0.0	0
	1	49	160.0	180.0	0	156	1.0	1
	2	37	130.0	283.0	0	98	0.0	0
	3	48	138.0	214.0	0	108	1.5	1
	4	54	150.0	195.0	0	122	0.0	0
	913	45	110.0	264.0	0	132	1.2	1
	914	68	144.0	193.0	1	141	3.4	1
	915	57	130.0	131.0	0	115	1.2	1
	916	57	130.0	236.0	0	174	0.0	1
	917	38	138.0	175.0	0	173	0.0	0

918 rows  $\times$  16 columns

In [27]:	<pre>df_encode = df_encode.astype(int)</pre>
----------	--

In [28]: df_encode	In	[28]:	df_	_encode
--------------------	----	-------	-----	---------

In [28]:	df_e	ncode						
Out[28]:		Age	RestingBP	Cholesterol	FastingBS	MaxHR	Oldpeak	HeartDisease
	0	40	140	289	0	172	0	0
	1	49	160	180	0	156	1	1
	2	37	130	283	0	98	0	0
	3	48	138	214	0	108	1	1
	4	54	150	195	0	122	0	0
	913	45	110	264	0	132	1	1
	914	68	144	193	1	141	3	1
	915	57	130	131	0	115	1	1
	916	57	130	236	0	174	0	1
	917	38	138	175	0	173	0	0

918 rows  $\times$  16 columns

## standard scaling

```
scaler = StandardScaler()
df_encode[numerical_cols] = scaler.fit_transform(df_encode[numerical_cols])
df_encode.head()
```

Out[29]:		Age	RestingBP	Cholesterol	FastingBS	MaxHR	Oldpeak	HeartDis
	0	-1.433140	0.414885	0.834754	0	1.382928	-0.727592	
	1	-0.478484	1.527224	-1.210675	0	0.754157	0.282891	
	2	-1.751359	-0.141284	0.722161	0	-1.525138	-0.727592	
	3	-0.584556	0.303651	-0.572651	0	-1.132156	0.282891	
	4	0.051881	0.971054	-0.929194	0	-0.581981	-0.727592	

In [30]: df_encode	In [30]:	df_encode				
--------------------	----------	-----------	--	--	--	--

Out[30]:		Age	RestingBP	Cholesterol	FastingBS	MaxHR	Oldpeak	Heart
	0	-1.433140	0.414885	0.834754	0	1.382928	-0.727592	
	1	-0.478484	1.527224	-1.210675	0	0.754157	0.282891	
	2	-1.751359	-0.141284	0.722161	0	-1.525138	-0.727592	
	3	-0.584556	0.303651	-0.572651	0	-1.132156	0.282891	
	4	0.051881	0.971054	-0.929194	0	-0.581981	-0.727592	
	913	-0.902775	-1.253622	0.365619	0	-0.188999	0.282891	
	914	1.536902	0.637353	-0.966725	1	0.164684	2.303858	
	915	0.370100	-0.141284	-2.130180	0	-0.857069	0.282891	
	916	0.370100	-0.141284	-0.159813	0	1.461525	-0.727592	
	917	-1.645286	0.303651	-1.304502	0	1.422226	-0.727592	

918 rows × 16 columns

```
from sklearn.svm import SVC
         from sklearn.neighbors import KNeighborsClassifier
In [35]: X = df encode.drop('HeartDisease', axis = 1)
         y = df encode['HeartDisease']
In [36]: X train, X test, y train, y test = train test split(X, y, test size=0.33, ra
In [45]: scaler = StandardScaler()
         X train scale = scaler.fit transform(X train)
         X test scale = scaler.fit transform(X test)
In [50]: models = {
             "Logistic Regression": LogisticRegression(),
             "Naive Bayes": GaussianNB(),
             "Decision Tree": DecisionTreeClassifier(),
             "SVM": SVC(probability = True),
             "KNN": KNeighborsClassifier()
In [51]: result =[]
In [52]: for name, model in models.items():
           model.fit(X train scale, y train)
           y pred = model.predict(X test scale)
           accuracy = accuracy score(y test, y pred)
           f1 = f1 score(y test, y pred)
           result.append({
               'model': name,
               'Accuracy': round(accuracy,4),
               'f1': round(f1,4)
               })
In [54]: result
Out[54]: [{'model': 'Logistic Regression', 'Accuracy': 0.8713, 'f1': 0.887},
          {'model': 'Naive Bayes', 'Accuracy': 0.8581, 'f1': 0.8746},
          {'model': 'Decision Tree', 'Accuracy': 0.7426, 'f1': 0.7651},
          {'model': 'SVM', 'Accuracy': 0.8614, 'f1': 0.88},
          {'model': 'KNN', 'Accuracy': 0.8449, 'f1': 0.863}]
In [55]: import joblib
         joblib.dump(models['Logistic Regression'], 'LR Heart.pkl')
         joblib.dump(scaler, 'scaler.pkl')
         joblib.dump(X.columns.tolist(), 'Columns.pkl')
Out[55]: ['Columns.pkl']
```