In [1]:

```
import numpy as np
import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
```

In [2]:

```
df=pd.read_csv(r"C:\Users\poorn\Downloads\drug200.csv")
df
```

Out[2]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	F	HIGH	HIGH	25.355	drugY
1	47	М	LOW	HIGH	13.093	drugC
2	47	М	LOW	HIGH	10.114	drugC
3	28	F	NORMAL	HIGH	7.798	drugX
4	61	F	LOW	HIGH	18.043	drugY
195	56	F	LOW	HIGH	11.567	drugC
196	16	М	LOW	HIGH	12.006	drugC
197	52	М	NORMAL	HIGH	9.894	drugX
198	23	М	NORMAL	NORMAL	14.020	drugX
199	40	F	LOW	NORMAL	11.349	drugX

200 rows × 6 columns

In [3]:

df.head()

Out[3]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	F	HIGH	HIGH	25.355	drugY
1	47	М	LOW	HIGH	13.093	drugC
2	47	М	LOW	HIGH	10.114	drugC
3	28	F	NORMAL	HIGH	7.798	drugX
4	61	F	LOW	HIGH	18.043	drugY

```
In [4]:
```

```
df.tail()
```

Out[4]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
195	56	F	LOW	HIGH	11.567	drugC
196	16	М	LOW	HIGH	12.006	drugC
197	52	М	NORMAL	HIGH	9.894	drugX
198	23	М	NORMAL	NORMAL	14.020	drugX
199	40	F	LOW	NORMAL	11.349	drugX

In [5]:

```
df.shape
```

Out[5]:

(200, 6)

In [6]:

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 200 entries, 0 to 199
Data columns (total 6 columns):
```

#	Column	Non-Null Count	Dtype				
0	Age	200 non-null	int64				
1	Sex	200 non-null	object				
2	BP	200 non-null	object				
3	Cholesterol	200 non-null	object				
4	Na_to_K	200 non-null	float64				
5	Drug	200 non-null	object				
d+vn	$\frac{1}{1}$						

dtypes: float64(1), int64(1), object(4)

memory usage: 9.5+ KB

In [7]:

```
df.isnull().sum()
```

Out[7]:

Age	0
Sex	0
BP	0
Cholesterol	0
Na_to_K	0
Drug	0
dtvpe: int64	

```
In [8]:
df['Sex'].value_counts()
Out[8]:
Sex
     104
Μ
F
      96
Name: count, dtype: int64
In [9]:
df['BP'].value_counts()
Out[9]:
BP
HIGH
          77
          64
LOW
NORMAL
          59
Name: count, dtype: int64
In [10]:
df['Cholesterol'].value_counts()
Out[10]:
Cholesterol
HIGH
          103
NORMAL
           97
Name: count, dtype: int64
In [11]:
df['Drug'].value_counts()
Out[11]:
Drug
drugY
         91
drugX
         54
drugA
         23
drugC
         16
drugB
         16
Name: count, dtype: int64
```

In [12]:

```
convert={'Sex':{"M":0,"F":1}}
df=df.replace(convert)
df
```

Out[12]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	1	HIGH	HIGH	25.355	drugY
1	47	0	LOW	HIGH	13.093	drugC
2	47	0	LOW	HIGH	10.114	drugC
3	28	1	NORMAL	HIGH	7.798	drugX
4	61	1	LOW	HIGH	18.043	drugY
195	56	1	LOW	HIGH	11.567	drugC
196	16	0	LOW	HIGH	12.006	drugC
197	52	0	NORMAL	HIGH	9.894	drugX
198	23	0	NORMAL	NORMAL	14.020	drugX
199	40	1	LOW	NORMAL	11.349	drugX

200 rows × 6 columns

In [13]:

```
convert={'BP':{"HIGH":1,"LOW":0,"NORMAL":2}}
df=df.replace(convert)
df
```

Out[13]:

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
0	23	1	1	HIGH	25.355	drugY
1	47	0	0	HIGH	13.093	drugC
2	47	0	0	HIGH	10.114	drugC
3	28	1	2	HIGH	7.798	drugX
4	61	1	0	HIGH	18.043	drugY
195	56	1	0	HIGH	11.567	drugC
196	16	0	0	HIGH	12.006	drugC
197	52	0	2	HIGH	9.894	drugX
198	23	0	2	NORMAL	14.020	drugX
199	40	1	0	NORMAL	11.349	drugX

200 rows × 6 columns

In [14]:

```
convert={'Cholesterol':{"HIGH":1,"NORMAL":0}}
df=df.replace(convert)
df
```

Out[14]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	1	1	1	25.355	drugY
1	47	0	0	1	13.093	drugC
2	47	0	0	1	10.114	drugC
3	28	1	2	1	7.798	drugX
4	61	1	0	1	18.043	drugY
195	56	1	0	1	11.567	drugC
196	16	0	0	1	12.006	drugC
197	52	0	2	1	9.894	drugX
198	23	0	2	0	14.020	drugX
199	40	1	0	0	11.349	drugX

200 rows × 6 columns

In [15]:

```
x=["Age","Sex","BP","Cholesterol"]
y=["drugY","drugC","drugX","drugA","drugB"]
all_inputs=df[x]
all_classes=df["Drug"]
```

In [16]:

```
(x_train,x_test,y_train,y_test)=train_test_split(all_inputs,all_classes,test_size=0.5)
```

In [17]:

```
clf=DecisionTreeClassifier(random_state=0)
```

In [18]:

```
clf.fit(x_train,y_train)
```

Out[18]:

```
DecisionTreeClassifier
DecisionTreeClassifier(random_state=0)
```

In [19]:	
<pre>score=clf.score(x_test,y_test) print(score)</pre>	
0.54	
In []:	