

```
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\sweet\Downloads\drug200.csv")
df
```

Out[2]:

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
0	23	F	HIGH	HIGH	25.355	drugY
1	47	M	LOW	HIGH	13.093	drugC
2	47	M	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	M	LOW	HIGH	12.006	drugC
197	52	M	NORMAL	HIGH	9.894	drugX
198	23	M	NORMAL	NORMAL	14.020	drugX
199	40	F	LOW	NORMAL	11.349	drugX

200 rows × 6 columns

In [3]: 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  
dtypes: float64(1), int64(1), object(4)
memory usage: 9.5+ KB
```

In [4]: df['Drug'].value_counts()

```
Out[4]: Drug
drugY    91
drugX    54
drugA    23
drugC    16
drugB    16
Name: count, dtype: int64
```

```
In [5]: df['Age'].value_counts()
```

Out[5]:

Age	
47	8
23	7
28	7
49	7
39	6
32	6
50	5
37	5
58	5
60	5
22	5
34	4
72	4
51	4
42	4
26	4
24	4
74	4
67	4
68	4
61	4
56	4
20	4
36	4
45	4
41	4
31	4
43	4
65	4
57	4
53	3
40	3
70	3
59	3
16	3
38	3
15	3
69	3
35	3
18	3

64	3
52	2
55	2
62	2
19	2
29	2
66	2
73	2
46	2
48	2
54	1
17	1
33	1
63	1
30	1
21	1
25	1

Name: count, dtype: int64

```
In [6]: convert={"Sex":{"F":1,"M":0}}
df=df.replace(convert)
df
```

Out[6]:

	Age	Sex	BP	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 [7]: convert={'BP':{'HIGH':1,"LOW":2,"NORMAL":3}}
df=df.replace(convert)
df
```

Out[7]:

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

200 rows × 6 columns

```
In [8]: x=["Age","Sex"]
y=["Yes","No"]
all_inputs=df[x]
all_classes=df["Cholesterol"]
```

```
In [9]: (x_train,x_test,y_train,y_test)=train_test_split(all_inputs,all_classes,test_size=0.5)
```

```
In [10]: clf=DecisionTreeClassifier(random_state=0)
```

```
In [11]: clf.fit(x_train,y_train)
```

```
Out[11]:
```

▼

DecisionTreeClassifier

DecisionTreeClassifier(random_state=0)

```
In [12]: score=clf.score(x_test,y_test)  
print(score)
```

0.4