```
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\jangidi veena\OneDrive\Documents\jupyter\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]: convert={'Cholesterol':{"HIGH":1,"NORMAL":0}}
    df=df.replace(convert)
    df
```

Out[3]:

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

200 rows × 6 columns

```
In [4]: df.info()
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 200 entries, 0 to 199
         Data columns (total 6 columns):
                            Non-Null Count Dtype
              Column
          ---
          0
              Age
                            200 non-null
                                            int64
          1
                            200 non-null
                                            object
              Sex
          2
              BP
                            200 non-null
                                            object
          3
              Cholesterol 200 non-null
                                            int64
          4
              Na_to_K
                            200 non-null
                                            float64
          5
              Drug
                            200 non-null
                                            object
         dtypes: float64(1), int64(2), object(3)
         memory usage: 9.5+ KB
In [10]: df['Cholesterol'].value_counts()
Out[10]: Cholesterol
              103
         1
               97
         Name: count, dtype: int64
In [11]: |df['BP'].value_counts()
Out[11]: BP
                    77
         HIGH
         LOW
                    64
         NORMAL
                    59
         Name: count, dtype: int64
In [12]: |df['Drug'].value_counts()
Out[12]: Drug
         drugY
                  91
         drugX
                   54
         drugA
                  23
         drugC
                  16
         drugB
                   16
         Name: count, dtype: int64
```

```
In [13]: convert={'BP':{'HIGH':1,'LOW':2,'NORMAL':3}}
    df=df.replace(convert)
    df
```

Out[13]:

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

200 rows × 6 columns

```
In [14]: convert={'Sex':{'F':1,'M':2}}
    df=df.replace(convert)
    df
```

Out[14]:

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

200 rows × 6 columns