```
In [1]: import numpy as ny
import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
```

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.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
                         200 non-null
                                         object
         1
             Sex
                          200 non-null
                                       object
         2
             Cholesterol 200 non-null
                                       object
                                        float64
         4
             Na_to_K
                          200 non-null
         5
             Drug
                          200 non-null
                                          object
        dtypes: float64(1), int64(1), object(4)
        memory usage: 9.5+ KB
In [4]: df['BP'].value_counts()
Out[4]: BP
        HIGH
                  77
        LOW
                  64
        NORMAL
                  59
        Name: count, dtype: int64
In [5]: df['Cholesterol'].value counts()
Out[5]: Cholesterol
        HIGH
                  103
                   97
        NORMAL
        Name: count, dtype: int64
```

Out[6]:

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

200 rows × 6 columns

Out[7]:

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

200 rows × 6 columns

Out[8]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	F	2	1	25.355	2
1	47	М	0	1	13.093	5
2	47	М	0	1	10.114	5
3	28	F	1	1	7.798	1
4	61	F	0	1	18.043	2
195	56	F	0	1	11.567	5
196	16	М	0	1	12.006	5
197	52	М	1	1	9.894	1
198	23	М	1	0	14.020	1
199	40	F	0	0	11.349	1

200 rows × 6 columns

```
In [9]: convert={"Sex":{"M":1,"F":2}}
    df=df.replace(convert)
    df
```

Out[9]:

	Age	Sex	ВР	Cholesterol	Na_to_K	Drug
0	23	2	2	1	25.355	2
1	47	1	0	1	13.093	5
2	47	1	0	1	10.114	5
3	28	2	1	1	7.798	1
4	61	2	0	1	18.043	2
195	56	2	0	1	11.567	5
196	16	1	0	1	12.006	5
197	52	1	1	1	9.894	1
198	23	1	1	0	14.020	1
199	40	2	0	0	11.349	1

200 rows × 6 columns

```
In [10]: convert={"Sex":{1:"M",2:"F"}}
    df=df.replace(convert)
    df
```

Out[10]:

2
5
5
1
2
5
5
1
1
1

200 rows × 6 columns

```
In [11]: x=["Age","Drug","Cholesterol"]
y=["M","F"]
all_inputs=df[x]
all_classes=df["Sex"]
```

```
In [12]: (x_train,x_test,y_train,y_test)=train_test_split(all_inputs,all_classes,test_size=0.25)
```

```
In [13]: clf=DecisionTreeClassifier(random_state=0)
```