

Assignment 6

Name: Khalate Shubham

Class: AIDS A

Rollno: 23107062

Batch: C

```
import pandas as pd
```

```
df=pd.read_csv("iris.csv")
```

```
df
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
\					
0	1	5.1	3.5	1.4	0.2
1	2	4.9	3.0	1.4	0.2
2	3	4.7	3.2	1.3	0.2
3	4	4.6	3.1	1.5	0.2
4	5	5.0	3.6	1.4	0.2
..
145	146	6.7	3.0	5.2	2.3
146	147	6.3	2.5	5.0	1.9
147	148	6.5	3.0	5.2	2.0
148	149	6.2	3.4	5.4	2.3
149	150	5.9	3.0	5.1	1.8

	Species
0	setosa
1	setosa
2	setosa
3	setosa
4	setosa
..	...
145	virginica
146	virginica
147	virginica

```
148 virginica
149 virginica
```

```
[150 rows x 6 columns]
```

```
df.count()
```

```
Unnamed: 0      150
Sepal.Length    150
Sepal.Width     150
Petal.Length    150
Petal.Width     150
Species         150
dtype: int64
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 150 entries, 0 to 149
```

```
Data columns (total 6 columns):
```

#	Column	Non-Null Count	Dtype
0	Unnamed: 0	150 non-null	int64
1	Sepal.Length	150 non-null	float64
2	Sepal.Width	150 non-null	float64
3	Petal.Length	150 non-null	float64
4	Petal.Width	150 non-null	float64
5	Species	150 non-null	object

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

```
memory usage: 7.2+ KB
```

```
df.describe()
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length
Petal.Width				
count	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.057333	3.758000
std	43.445368	0.828066	0.435866	1.765298
min	1.000000	4.300000	2.000000	1.000000
25%	38.250000	5.100000	2.800000	1.600000
50%	75.500000	5.800000	3.000000	4.350000
75%	112.750000	6.400000	3.300000	5.100000
max	150.000000	7.900000	4.400000	6.900000

```
df.notnull()
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Species					
0	True	True	True	True	True
1	True	True	True	True	True
2	True	True	True	True	True
3	True	True	True	True	True
4	True	True	True	True	True
..
...					
145	True	True	True	True	True
146	True	True	True	True	True
147	True	True	True	True	True
148	True	True	True	True	True
149	True	True	True	True	True

```
[150 rows x 6 columns]
```

```
df.isnull()
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Species					
0	False	False	False	False	False
1	False	False	False	False	False
2	False	False	False	False	False
3	False	False	False	False	False
4	False	False	False	False	False
..
...					
145	False	False	False	False	False
146	False	False	False	False	False
147	False	False	False	False	False

```
False
148      False      False      False      False      False
False
149      False      False      False      False      False
False
```

```
[150 rows x 6 columns]
```

```
df.dtypes
```

```
Unnamed: 0      int64
Sepal.Length    float64
Sepal.Width     float64
Petal.Length     float64
Petal.Width     float64
Species         object
dtype: object
```

```
df.shape
```

```
(150, 6)
```

```
df.ndim
```

```
2
```

```
df.value_counts()
```

Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Species				
1	5.1	3.5	1.4	0.2
setosa	1			
95	5.6	2.7	4.2	1.3
versicolor	1			
97	5.7	2.9	4.2	1.3
versicolor	1			
98	6.2	2.9	4.3	1.3
versicolor	1			
99	5.1	2.5	3.0	1.1
versicolor	1			
..				
51	7.0	3.2	4.7	1.4
versicolor	1			
52	6.4	3.2	4.5	1.5
versicolor	1			
53	6.9	3.1	4.9	1.5
versicolor	1			
54	5.5	2.3	4.0	1.3
versicolor	1			
150	5.9	3.0	5.1	1.8

```
virginica      1  
Name: count, Length: 150, dtype: int64
```

```
x=df.drop(['Species'],axis=1)
```

```
x
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
0	1	5.1	3.5	1.4	0.2
1	2	4.9	3.0	1.4	0.2
2	3	4.7	3.2	1.3	0.2
3	4	4.6	3.1	1.5	0.2
4	5	5.0	3.6	1.4	0.2
...
145	146	6.7	3.0	5.2	2.3
146	147	6.3	2.5	5.0	1.9
147	148	6.5	3.0	5.2	2.0
148	149	6.2	3.4	5.4	2.3
149	150	5.9	3.0	5.1	1.8

```
[150 rows x 5 columns]
```

```
y=df["Species"]
```

```
y
```

0	setosa
1	setosa
2	setosa
3	setosa
4	setosa

...	...
145	virginica
146	virginica
147	virginica
148	virginica
149	virginica

```
Name: Species, Length: 150, dtype: object
```

```
from sklearn import linear_model
```

```
from sklearn.model_selection import train_test_split
```

```
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.2)
```

```
x_train
```

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
90	91	5.5	2.6	4.4	1.2
101	102	5.8	2.7	5.1	1.9
14	15	5.8	4.0	1.2	0.2
51	52	6.4	3.2	4.5	1.5
71	72	6.1	2.8	4.0	1.3

130	131	7.4	2.8	6.1	1.9
27	28	5.2	3.5	1.5	0.2
19	20	5.1	3.8	1.5	0.3
145	146	6.7	3.0	5.2	2.3
0	1	5.1	3.5	1.4	0.2

[120 rows x 5 columns]

x_test

	Unnamed: 0	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
43	44	5.0	3.5	1.6	0.6
82	83	5.8	2.7	3.9	1.2
84	85	5.4	3.0	4.5	1.5
98	99	5.1	2.5	3.0	1.1
80	81	5.5	2.4	3.8	1.1
18	19	5.7	3.8	1.7	0.3
94	95	5.6	2.7	4.2	1.3
125	126	7.2	3.2	6.0	1.8
110	111	6.5	3.2	5.1	2.0
55	56	5.7	2.8	4.5	1.3
100	101	6.3	3.3	6.0	2.5
143	144	6.8	3.2	5.9	2.3
119	120	6.0	2.2	5.0	1.5
37	38	4.9	3.6	1.4	0.1
32	33	5.2	4.1	1.5	0.1
91	92	6.1	3.0	4.6	1.4
58	59	6.6	2.9	4.6	1.3
78	79	6.0	2.9	4.5	1.5
87	88	6.3	2.3	4.4	1.3
23	24	5.1	3.3	1.7	0.5
29	30	4.7	3.2	1.6	0.2
111	112	6.4	2.7	5.3	1.9
11	12	4.8	3.4	1.6	0.2
86	87	6.7	3.1	4.7	1.5
142	143	5.8	2.7	5.1	1.9
138	139	6.0	3.0	4.8	1.8
54	55	6.5	2.8	4.6	1.5
139	140	6.9	3.1	5.4	2.1
93	94	5.0	2.3	3.3	1.0
40	41	5.0	3.5	1.3	0.3

y_train

90	versicolor
101	virginica
14	setosa
51	versicolor
71	versicolor

```
...
130    virginica
27     setosa
19     setosa
145    virginica
0      setosa
Name: Species, Length: 120, dtype: object
```

y_test

```
43     setosa
82    versicolor
84    versicolor
98    versicolor
80    versicolor
18     setosa
94    versicolor
125   virginica
110   virginica
55    versicolor
100   virginica
143   virginica
119   virginica
37     setosa
32     setosa
91    versicolor
58    versicolor
78    versicolor
87    versicolor
23     setosa
29     setosa
111   virginica
11     setosa
86    versicolor
142   virginica
138   virginica
54    versicolor
139   virginica
93    versicolor
40     setosa
Name: Species, dtype: object
```

```
len(x_train)
```

```
120
```

```
len(x_test)
```

```
30
```

```
len(y_train)
```

```

120
len(y_test)
30
from sklearn.naive_bayes import GaussianNB,MultinomialNB,BernoulliNB
GNB=GaussianNB()
GNB.fit(x_train,y_train)
GaussianNB()
y_pred=GNB.predict(x_test)
y_pred
array(['setosa', 'versicolor', 'versicolor', 'versicolor',
       'versicolor',
       'setosa', 'versicolor', 'virginica', 'virginica', 'versicolor',
       'virginica', 'virginica', 'virginica', 'setosa', 'setosa',
       'versicolor', 'versicolor', 'versicolor', 'versicolor',
       'setosa',
       'setosa', 'virginica', 'setosa', 'versicolor', 'virginica',
       'virginica', 'versicolor', 'virginica', 'versicolor',
       'setosa'],
      dtype='<U10')

from sklearn.metrics import confusion_matrix,accuracy_score
cm=confusion_matrix(y_test,y_pred)
cm
array([[ 8,  0,  0],
       [ 0, 13,  0],
       [ 0,  0,  9]])

GNB.score(x_train,y_train)
0.9916666666666667

from sklearn.metrics import accuracy_score,classification_report
from sklearn import metrics
accuracy_g=accuracy_score(y_test,y_pred)
accuracy_g
1.0
error_rate=1-accuracy_g

```



```
error_rate
```

```
0.0
```

```
cr_g=classification_report(y_pred,y_test)
```

```
print(cr_g)
```

	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	8
versicolor	1.00	1.00	1.00	13
virginica	1.00	1.00	1.00	9
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
import seaborn as sns
```

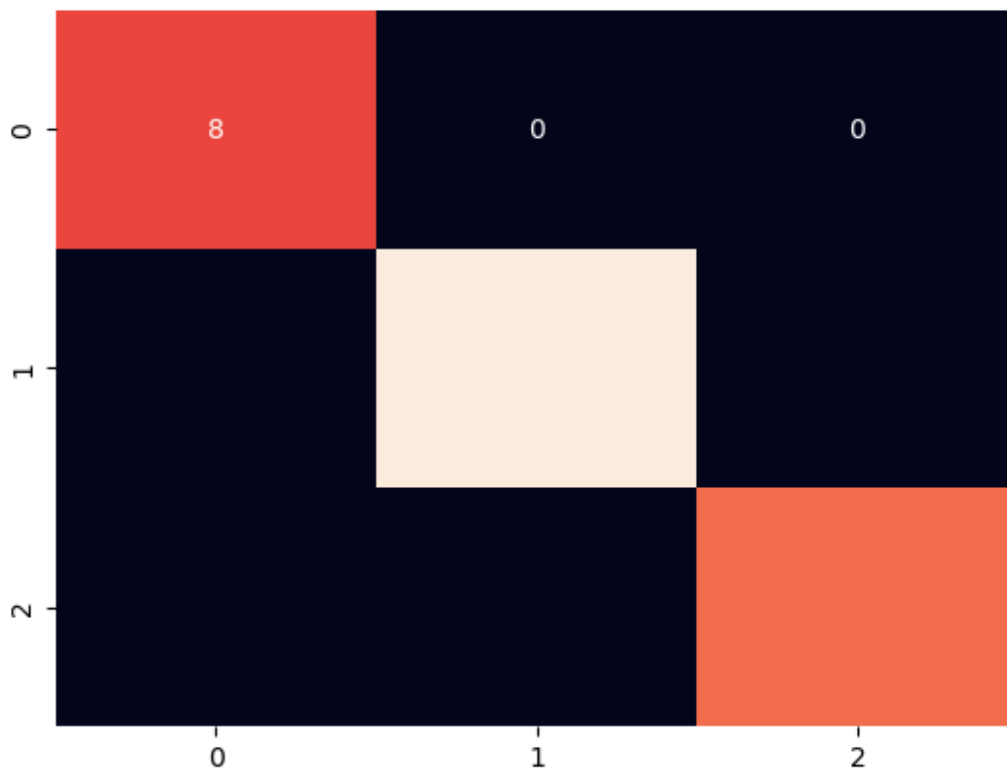
```
import matplotlib.pyplot as plt
```

```
sns.heatmap(cm,annot=True,cbar=False)
```

```
plt.xlabel = ("actual value")
```

```
plt.ylabel = ("predicted value")
```

```
plt.show()
```



```
M=BernoulliNB()
M.fit(x_train,y_train)
BernoulliNB()
M.fit
<bound method _BaseDiscreteNB.fit of BernoulliNB(>
y_pred1=M.predict(x_test)
cm1=confusion_matrix(y_test,y_pred1)
cm1
array([[ 8,  0,  0],
       [13,  0,  0],
       [ 9,  0,  0]])
v=MultinomialNB()
v
MultinomialNB()
v.fit(x_train,y_train)
MultinomialNB()
y_pred2=v.predict(x_test)
cm2=confusion_matrix(y_test,y_pred2)
cm2
array([[7, 1, 0],
       [0, 6, 7],
       [0, 1, 8]])
```