

```
In [1]: import numpy as np
import pandas as pd
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
```

```
In [2]: df = sns.load_dataset('iris')
df
```

```
Out[2]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa
...	...	...	...	...	...
145	6.7	3.0	5.2	2.3	virginica
146	6.3	2.5	5.0	1.9	virginica
147	6.5	3.0	5.2	2.0	virginica
148	6.2	3.4	5.4	2.3	virginica
149	5.9	3.0	5.1	1.8	virginica

150 rows × 5 columns

```
In [3]: x = df[['sepal_length', 'sepal_width', 'petal_length', 'petal_width']]
y = df[['species']]
print(x)
print(y)

# x = df.iloc[:, :4]
# y = df['species']
# print(x)
# print(y)
```

	sepal_length	sepal_width	petal_length	petal_width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
..	...	...	...	...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

[150 rows x 4 columns]

	species
0	setosa
1	setosa
2	setosa
3	setosa
4	setosa
..	...
145	virginica
146	virginica
147	virginica
148	virginica
149	virginica

[150 rows x 1 columns]

```
In [4]: from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.25, random_st
```

```
In [5]: from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
x_test = sc.fit_transform(x_test)
x_train = sc.fit_transform(x_train)

# print(x_test)
# print(x_train)
```

```
In [6]: from sklearn.naive_bayes import GaussianNB
nb = GaussianNB()
nb.fit(x_train, y_train)
```

```
Out[6]: GaussianNB
GaussianNB()
```

```
In [7]: y_predict = nb.predict(x_test)
print(y_predict)
```

```
['virginica' 'versicolor' 'setosa' 'versicolor' 'versicolor' 'setosa'
'versicolor' 'versicolor' 'setosa' 'versicolor' 'virginica' 'versicolor'
'setosa' 'virginica' 'setosa' 'virginica' 'virginica' 'virginica'
'setosa' 'setosa' 'versicolor' 'virginica' 'versicolor' 'versicolor'
'virginica' 'virginica' 'versicolor' 'versicolor' 'virginica' 'virginica'
'virginica' 'versicolor' 'setosa' 'virginica' 'versicolor' 'setosa'
'setosa' 'setosa']
```

In [8]: `from sklearn.metrics import accuracy_score, confusion_matrix, precision_score, reca`

```
acc = accuracy_score(y_test, y_predict)
print(acc)

pr = precision_score(y_test, y_predict, average='macro') # Using average because we
print(pr)

rc = recall_score(y_test, y_predict, average='macro')
print(rc)

cm = confusion_matrix(y_test, y_predict)
print(cm)

cr = classification_report(y_test, y_predict)
print(cr)
```

0.8947368421052632

0.9010989010989011

0.9010989010989011

[[11 0 0]

[ 0 12 2]

[ 0 2 11]]

	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	11
versicolor	0.86	0.86	0.86	14
virginica	0.85	0.85	0.85	13
accuracy			0.89	38
macro avg	0.90	0.90	0.90	38
weighted avg	0.89	0.89	0.89	38

In [9]: `w = [[2.7,4.6,3.3,1.9]]`  
`w = sc.fit_transform(w)`

```
q = nb.predict(w)
print(q)
```

['versicolor']