

```

import pandas as pd
from sklearn.datasets import load_iris

iris = load_iris()
dir(iris)

['DESCR',
 'data',
 'data_module',
 'feature_names',
 'filename',
 'frame',
 'target',
 'target_names']

iris.feature_names

['sepal length (cm)',
 'sepal width (cm)',
 'petal length (cm)',
 'petal width (cm)']

iris.target_names

array(['setosa', 'versicolor', 'virginica'], dtype='<U10')

data = pd.DataFrame(iris.data , columns = iris.feature_names)
data["target"] = iris.target

data

```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	
1	4.9	3.0	1.4	
2	4.7	3.2	1.3	
3	4.6	3.1	1.5	
4	5.0	3.6	1.4	
...	...	...	...	
145	6.7	3.0	5.2	
146	6.3	2.5	5.0	
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

```

```

      target
0         0
1         0
2         0
3         0
4         0
..      ...
145       2
146       2
147       2
148       2
149       2

```

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

```
iris.target_names
```

```
array(['setosa', 'versicolor', 'virginica'], dtype='<U10')
```

```
# if target == 0 then it's setosa else if target == 1 then it's
versicolor else (2) then it's virginica
```

```
data[data.target == 0].head() #setosa
```

```

      sepal length (cm)  sepal width (cm)  petal length (cm)  petal width
(cm) \
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

```

```

      target
0         0
1         0
2         0
3         0
4         0

```

```
data[data.target == 1].head() #versicolor
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
50	7.0	3.2	4.7	1.4
51	6.4	3.2	4.5	1.5
52	6.9	3.1	4.9	1.5
53	5.5	2.3	4.0	1.3
54	6.5	2.8	4.6	1.5

	target
50	1
51	1
52	1
53	1
54	1

```
data[data.target == 1].head() # virginica
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
50	7.0	3.2	4.7	1.4
51	6.4	3.2	4.5	1.5
52	6.9	3.1	4.9	1.5
53	5.5	2.3	4.0	1.3
54	6.5	2.8	4.6	1.5

	target
50	1
51	1
52	1
53	1
54	1

```
data["flower_name"] = data.target.apply(lambda x:
iris.target_names[x])
data.head()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
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			

	target	flower_name
0	0	setosa
1	0	setosa
2	0	setosa
3	0	setosa
4	0	setosa

```
data.to_csv("iris_dataset.csv",index = False)
```

```
dataz = pd.read_csv("iris_dataset.csv")
dataz.head()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
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				

	target	flower_name
0	0	setosa
1	0	setosa
2	0	setosa
3	0	setosa
4	0	setosa

```
data0 =dataz[dataz.target ==0]
data1 =dataz[dataz.target ==1]
data2 =dataz[dataz.target ==2]
```

```
data2.head(5)
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
100	6.3	3.3	6.0	
2.5				

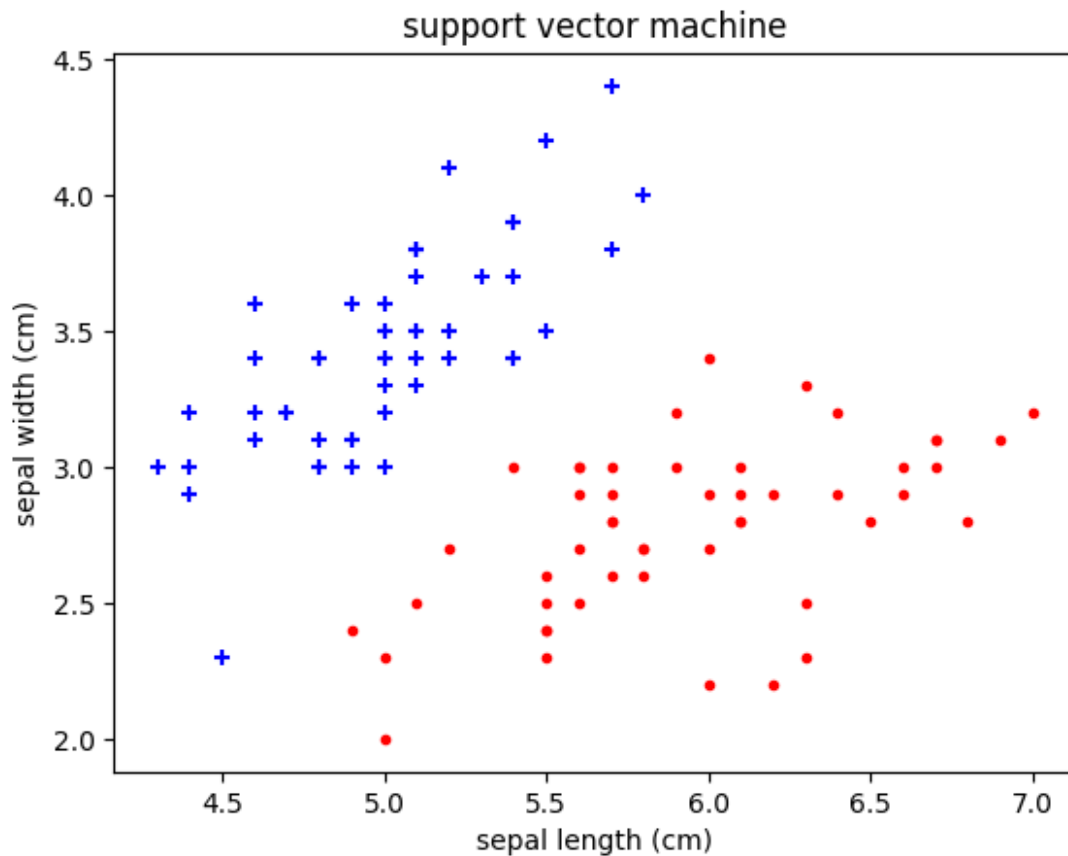
101	5.8	2.7	5.1
1.9			
102	7.1	3.0	5.9
2.1			
103	6.3	2.9	5.6
1.8			
104	6.5	3.0	5.8
2.2			

	target	flower_name
100	2	virginica
101	2	virginica
102	2	virginica
103	2	virginica
104	2	virginica

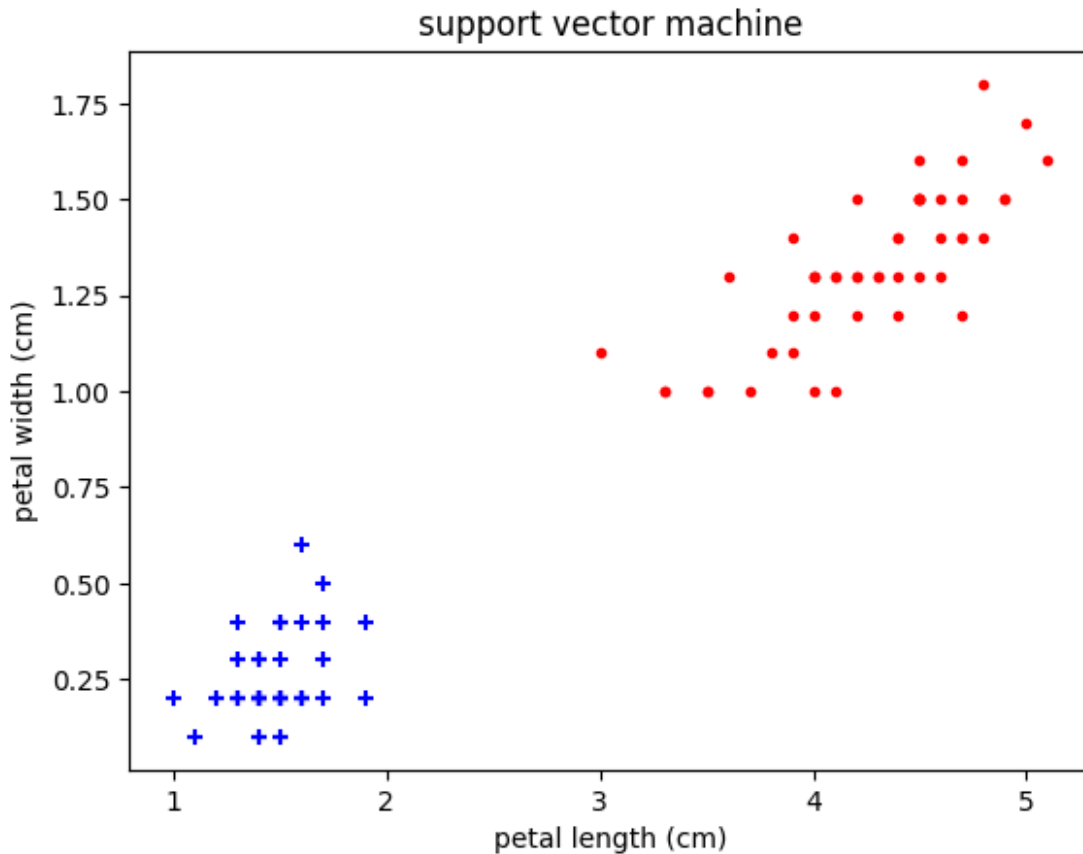
```
import matplotlib.pyplot as plt
%matplotlib inline

plt.title("support vector machine")
plt.xlabel("sepal length (cm)")
plt.ylabel("sepal width (cm)")
plt.scatter(data0["sepal length (cm)"],data0["sepal width (cm)"],marker="+",color="blue")
plt.scatter(data1["sepal length (cm)"],data1["sepal width (cm)"],marker=".",color="red")

<matplotlib.collections.PathCollection at 0x19ba6eaffe0>
```



```
plt.title("support vector machine")
plt.xlabel("petal length (cm)")
plt.ylabel("petal width (cm)")
plt.scatter(data0["petal length (cm)"],data0["petal width (cm)"],marker="+",color="blue")
plt.scatter(data1["petal length (cm)"],data1["petal width (cm)"],marker=".",color="red")
<matplotlib.collections.PathCollection at 0x19b8229e9f0>
```



```
## training the data using sklearn module
```

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

```
x= dataz.drop(["target","flower_name"],axis ="columns")
x.head()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
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

```
y = dataz.target
y.head()
```

```

0      0
1      0
2      0
3      0
4      0
Name: target, dtype: int64

x_train,x_test,y_train,y_test = train_test_split(x,y,test_size = 0.2)
len(x_train)
120

from sklearn.svm import SVC
model = SVC()
model.fit(x_train,y_train)
SVC()

y_pred = model.predict(x_test)
y_pred
array([0, 1, 1, 0, 0, 2, 2, 1, 0, 2, 0, 1, 1, 2, 0, 2, 2, 2, 0, 2, 0,
1,
      0, 1, 1, 1, 2, 0, 1, 2])

from sklearn.metrics import confusion_matrix
confusion_matrix(y_test,y_pred)
array([[10,  0,  0],
       [ 0, 10,  0],
       [ 0,  0, 10]])

from sklearn.metrics import accuracy_score
accuracy_score(y_test,y_pred)*100
100.0

##### manual prediction
model.predict([[5.9 ,3.0 ,5.1 ,1.8 ]])

C:\Users\suvas\AppData\Local\Programs\Python\Python312\Lib\site-
packages\sklearn\base.py:493: UserWarning: X does not have valid
feature names, but SVC was fitted with feature names
  warnings.warn(
array([2])

```



```
import numpy as np
x = int(input("tell me the above prdicted value 1 or 2 or 3 : "))

if np.array([x]) == 0 :
    print("setosa")
elif np.array([x]) == 1 :
    print("versicolor")
elif np.array([x]) == 2:
    print("virginica")

tell me the above prdicted value 1 or 2 or 3 : 2
virginica
```