In [4]:
```
sklearn.datasets import load_iris6
iris= load_iris()
dir(iris)
iris
```

Out[4]:
```
{'data': array([[5.1, 3.5, 1.4, 0.2],
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                [5.2, 2.7, 3.9, 1.4],
```

```
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[6.8, 3. , 5.5, 2.1],
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[6. , 2.2, 5. , 1.5],
```

```
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        [6.9, 3.1, 5.4, 2.1],
        [6.7, 3.1, 5.6, 2.4],
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        [6.8, 3.2, 5.9, 2.3],
        [6.7, 3.3, 5.7, 2.5],
        [6.7, 3. , 5.2, 2.3],
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        [6.2, 3.4, 5.4, 2.3],
        [5.9, 3. , 5.1, 1.8]]),
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 'frame': None,
 'target_names': array(['setosa', 'versicolor', 'virginica'], dtype='<U10'),
 'DESCR': '.. _iris_dataset:\n\nIris plants dataset\n--------------------\n\n**
Data Set Characteristics:**\n\n    :Number of Instances: 150 (50 in each of thr
ee classes)\n    :Number of Attributes: 4 numeric, predictive attributes and th
e class\n    :Attribute Information:\n        - sepal length in cm\n        - s
epal width in cm\n        - petal length in cm\n        - petal width in cm\n
- class:\n                - Iris-Setosa\n                - Iris-Versicolour\n
- Iris-Virginica\n                \n    :Summary Statistics:\n\n    ===========
=== ==== ==== ======= ===== ====================\n                    Min  Max
Mean    SD   Class Correlation\n    ============== ==== ==== ======= ===== ====
================\n    sepal length:   4.3  7.9   5.84   0.83    0.7826\n    sep
al width:    2.0  4.4   3.05   0.43   -0.4194\n    petal length:   1.0  6.9
3.76   1.76    0.9490  (high!)\n    petal width:    0.1  2.5   1.20   0.76
0.9565  (high!)\n    ============== ==== ==== ======= ===== ====================
=\n\n    :Missing Attribute Values: None\n    :Class Distribution: 33.3% for ea
ch of 3 classes.\n    :Creator: R.A. Fisher\n    :Donor: Michael Marshall (MARS
HALL%PLU@io.arc.nasa.gov)\n    :Date: July, 1988\n\nThe famous Iris database, f
irst used by Sir R.A. Fisher. The dataset is taken\nfrom Fisher\'s paper. Note
that it\'s the same as in R, but not as in the UCI\nMachine Learning Repositor
y, which has two wrong data points.\n\nThis is perhaps the best known database
to be found in the\npattern recognition literature.  Fisher\'s paper is a class
```

```
ic in the field and\nis referenced frequently to this day.  (See Duda & Hart, f
or example.)  The\ndata set contains 3 classes of 50 instances each, where each
class refers to a\ntype of iris plant.  One class is linearly separable from th
e other 2; the\nlatter are NOT linearly separable from each other.\n\n.. topi
c:: References\n\n   - Fisher, R.A. "The use of multiple measurements in taxono
mic problems"\n     Annual Eugenics, 7, Part II, 179-188 (1936); also in "Contr
ibutions to\n     Mathematical Statistics" (John Wiley, NY, 1950).\n   - Duda,
R.O., & Hart, P.E. (1973) Pattern Classification and Scene Analysis.\n     (Q32
7.D83) John Wiley & Sons.  ISBN 0-471-22361-1.  See page 218.\n   - Dasarathy,
B.V. (1980) "Nosing Around the Neighborhood: A New System\n     Structure and C
lassification Rule for Recognition in Partially Exposed\n     Environments".  I
EEE Transactions on Pattern Analysis and Machine\n     Intelligence, Vol. PAMI-
2, No. 1, 67-71.\n   - Gates, G.W. (1972) "The Reduced Nearest Neighbor Rule".
IEEE Transactions\n     on Information Theory, May 1972, 431-433.\n   - See als
o: 1988 MLC Proceedings, 54-64.  Cheeseman et al"s AUTOCLASS II\n     conceptua
l clustering system finds 3 classes in the data.\n   - Many, many more ...',
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  'petal length (cm)',
  'petal width (cm)'],
 'filename': 'iris.csv',
 'data_module': 'sklearn.datasets.data'}
```

In [3]:
```
iris.feature_names
```

Out[3]:
```
['sepal length (cm)',
 'sepal width (cm)',
 'petal length (cm)',
 'petal width (cm)']
```

In [5]:
```
iris.data
```

Out[5]:
```
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       [5.4, 3.9, 1.7, 0.4],
       [4.6, 3.4, 1.4, 0.3],
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       [4.8, 3.4, 1.6, 0.2],
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       [5.5, 3.5, 1.3, 0.2],
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       [5. , 3.5, 1.6, 0.6],
       [5.1, 3.8, 1.9, 0.4],
       [4.8, 3. , 1.4, 0.3],
       [5.1, 3.8, 1.6, 0.2],
       [4.6, 3.2, 1.4, 0.2],
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       [7. , 3.2, 4.7, 1.4],
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       [5.5, 2.3, 4. , 1.3],
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       [5.2, 2.7, 3.9, 1.4],
```

```
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[6. , 2.2, 5. , 1.5],
```

```
       [6.9, 3.2, 5.7, 2.3],
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       [6.3, 2.5, 5. , 1.9],
       [6.5, 3. , 5.2, 2. ],
       [6.2, 3.4, 5.4, 2.3],
       [5.9, 3. , 5.1, 1.8]])
```

In [12]:
```
import pandas as pd
df =pd.DataFrame(iris.data ,columns= iris.feature_names)
df.head()
```

Out[12]:

	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

In [13]:
```
iris.target_names
```

Out[13]:
```
array(['setosa', 'versicolor', 'virginica'], dtype='<U10')
```

In [16]:
```
df['target'] =iris. target
df.head
```

Out[16]:
```
<bound method NDFrame.head of      sepal length (cm)  sepal width (cm)  petal l
ength (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
..                 ...               ...                ...               ...
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

     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]>
```

In [25]:
```
df[df.target==2].head()
```

Out[25]:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
100	6.3	3.3	6.0	2.5	2
101	5.8	2.7	5.1	1.9	2
102	7.1	3.0	5.9	2.1	2
103	6.3	2.9	5.6	1.8	2
104	6.5	3.0	5.8	2.2	2

In [48]:
```
df0 =df[df.target==0]
df1 =df[df.target==1]
df2 =df[df.target==2]

df0.head(),df1.head(),df2.head()
```

Out[48]:
```
(   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 flowers_names Flowers_names
 0       0        setosa        setosa
 1       0        setosa        setosa
 2       0        setosa        setosa
 3       0        setosa        setosa
 4       0        setosa        setosa  ,
     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 flowers_names Flowers_names
 50       1    versicolor    versicolor
 51       1    versicolor    versicolor
 52       1    versicolor    versicolor
 53       1    versicolor    versicolor
 54       1    versicolor    versicolor  ,
      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 flowers_names Flowers_names
 100       2     virginica     virginica
 101       2     virginica     virginica
 102       2     virginica     virginica
 103       2     virginica     virginica
 104       2     virginica     virginica  )
```

In [45]:
```
df['Flowers_names'] = df.target.apply( lambda x: iris.target_names[x])
df
```

Out[45]:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	flowers_names	Flowers_names
0	5.1	3.5	1.4	0.2	0	setosa	setosa
1	4.9	3.0	1.4	0.2	0	setosa	setosa
2	4.7	3.2	1.3	0.2	0	setosa	setosa
3	4.6	3.1	1.5	0.2	0	setosa	setosa
4	5.0	3.6	1.4	0.2	0	setosa	setosa
...	...	...	...	...	...	...	...
145	6.7	3.0	5.2	2.3	2	virginica	virginica
146	6.3	2.5	5.0	1.9	2	virginica	virginica
147	6.5	3.0	5.2	2.0	2	virginica	virginica
148	6.2	3.4	5.4	2.3	2	virginica	virginica
149	5.9	3.0	5.1	1.8	2	virginica	virginica

150 rows × 7 columns

In [57]:
```
X=df.drop(['flowers_names','target','Flowers_names']  ,axis = 'columns')
X
```

Out[57]:

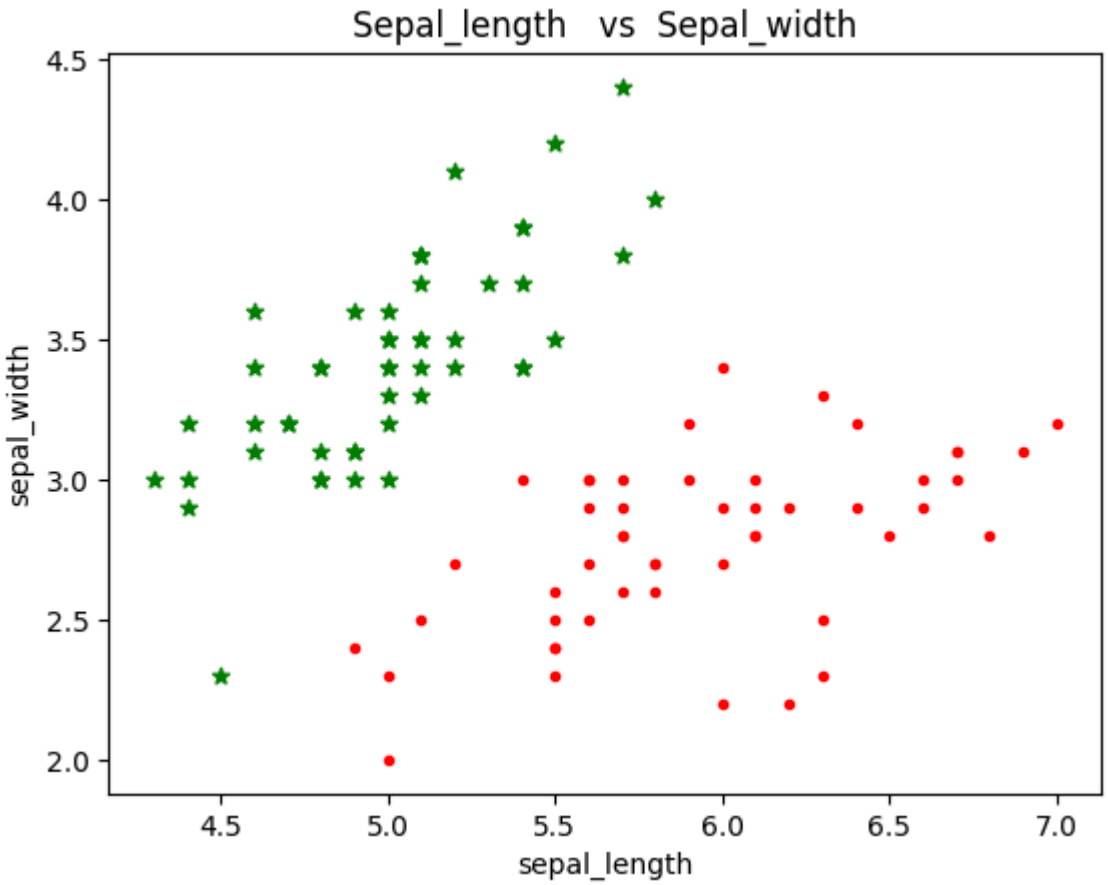
	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
...	...	...	...	...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

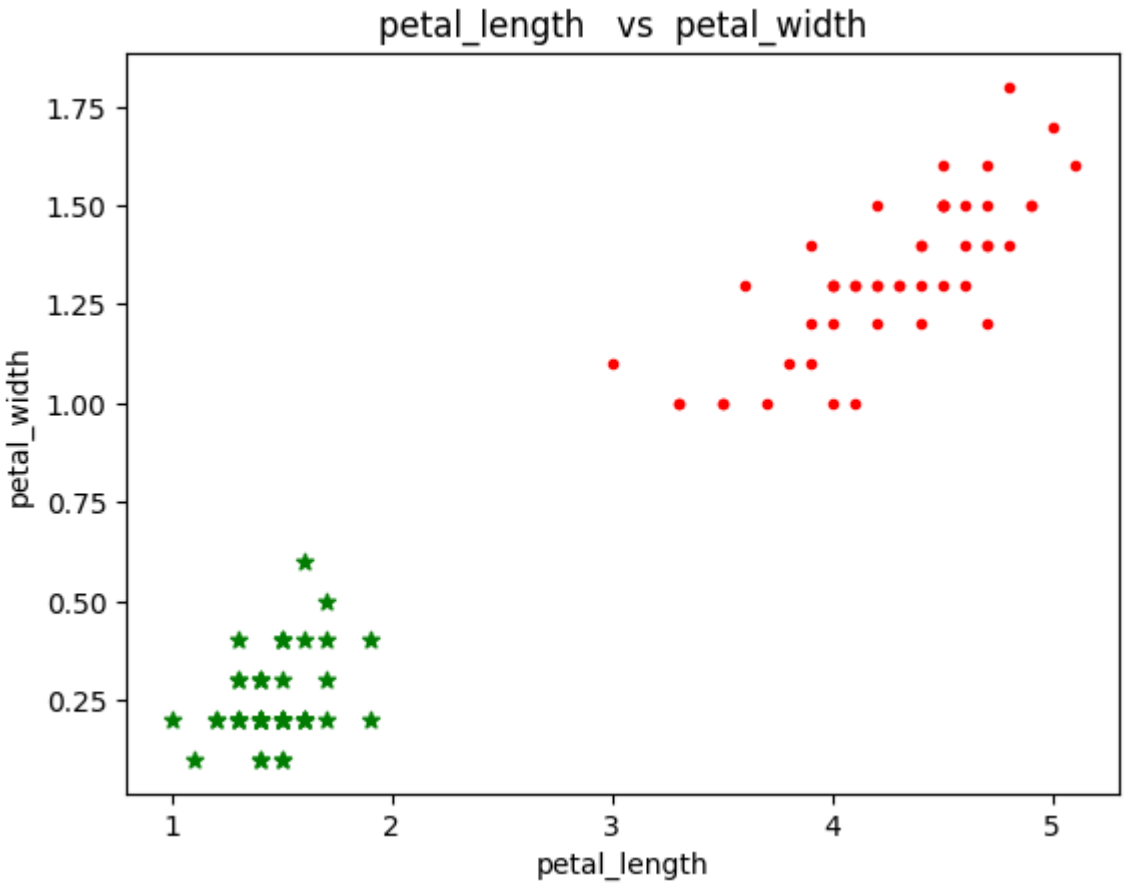
In [50]:
```
y= iris.target
y
```

Out[50]:
```
array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
       2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
       2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

In [73]:
```
from matplotlib import pyplot as pp
pp.scatter(df0['sepal length (cm)'],df0['sepal width (cm)'], color= "green",mark
pp.scatter(df1['sepal length (cm)'],df1['sepal width (cm)'], color= "red"  , mar
pp.title('Sepal_length  vs  Sepal_width')
pp.xlabel("sepal_length")
pp.ylabel("sepal_width")
pp.show()
```



In [76]:
```
pp.scatter(df0['petal length (cm)'],df0['petal width (cm)'], color= "green",mark
pp.scatter(df1['petal length (cm)'],df1['petal width (cm)'], color= "red",marker
pp.title('petal_length  vs  petal_width')
pp.xlabel("petal_length")
pp.ylabel("petal_width")
pp.show()
```



In [122…
```
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)
```

In [123…
```
len(X_train)
```

Out[123…
```
120
```

In [124…
```
len(X_test)
```

Out[124…
```
30
```

In [125…
```
from sklearn.svm import SVC
classifier = SVC()
```

In [126…
```
classifier.fit(X_train,y_train)
```

Out[126…
```
SVC()
```

In [127…
```
y_pred =classifier . predict(X_test)
(y_pred)
```

Out[127…
```
array([1, 2, 0, 1, 2, 0, 1, 0, 0, 2, 0, 0, 2, 2, 0, 0, 2, 0, 0, 2, 1, 1,
       1, 2, 0, 1, 2, 1, 0, 1])
```

In [130…
```
from sklearn.metrics import accuracy_score,confusion_matrix
accuracy = accuracy_score(y_test, y_pred)
cm = confusion_matrix(y_test, y_pred)
print(f'Accuracy: {accuracy}')
print('Confusion Matrix:')
print(cm)
```

```
Accuracy: 1.0
Confusion Matrix:
[[12  0  0]
 [ 0  9  0]
 [ 0  0  9]]
```

In [131…

```
#  evaluation model
classifier.score(X_test,y_test)
```

Out[131…

```
1.0
```