

```
In [26]: # following steps
# step 1 Prepare A & b
# step 2 Import ML Algorithm
# step 3 train ML model
# step 4 new test predic
```

```
In [5]: # Library
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn import datasets

from sklearn.tree import DecisionTreeClassifier
from sklearn import metrics
%pylab inline
%matplotlib inline
```

%pylab is deprecated, use %matplotlib inline and import the required libraries.  
Populating the interactive namespace from numpy and matplotlib

```
In [6]: # data Load
iris = pd.read_csv("./Downloads/Iris.csv")
iris.head()
```

```
Out[6]:
```

	<b>Id</b>	<b>SepalLengthCm</b>	<b>SepalWidthCm</b>	<b>PetalLengthCm</b>	<b>PetalWidthCm</b>	<b>Species</b>
<b>0</b>	1	5.1	3.5	1.4	0.2	Iris-setosa
<b>1</b>	2	4.9	3.0	1.4	0.2	Iris-setosa
<b>2</b>	3	4.7	3.2	1.3	0.2	Iris-setosa
<b>3</b>	4	4.6	3.1	1.5	0.2	Iris-setosa
<b>4</b>	5	5.0	3.6	1.4	0.2	Iris-setosa

```
In [7]: iris["Species"].value_counts()
```

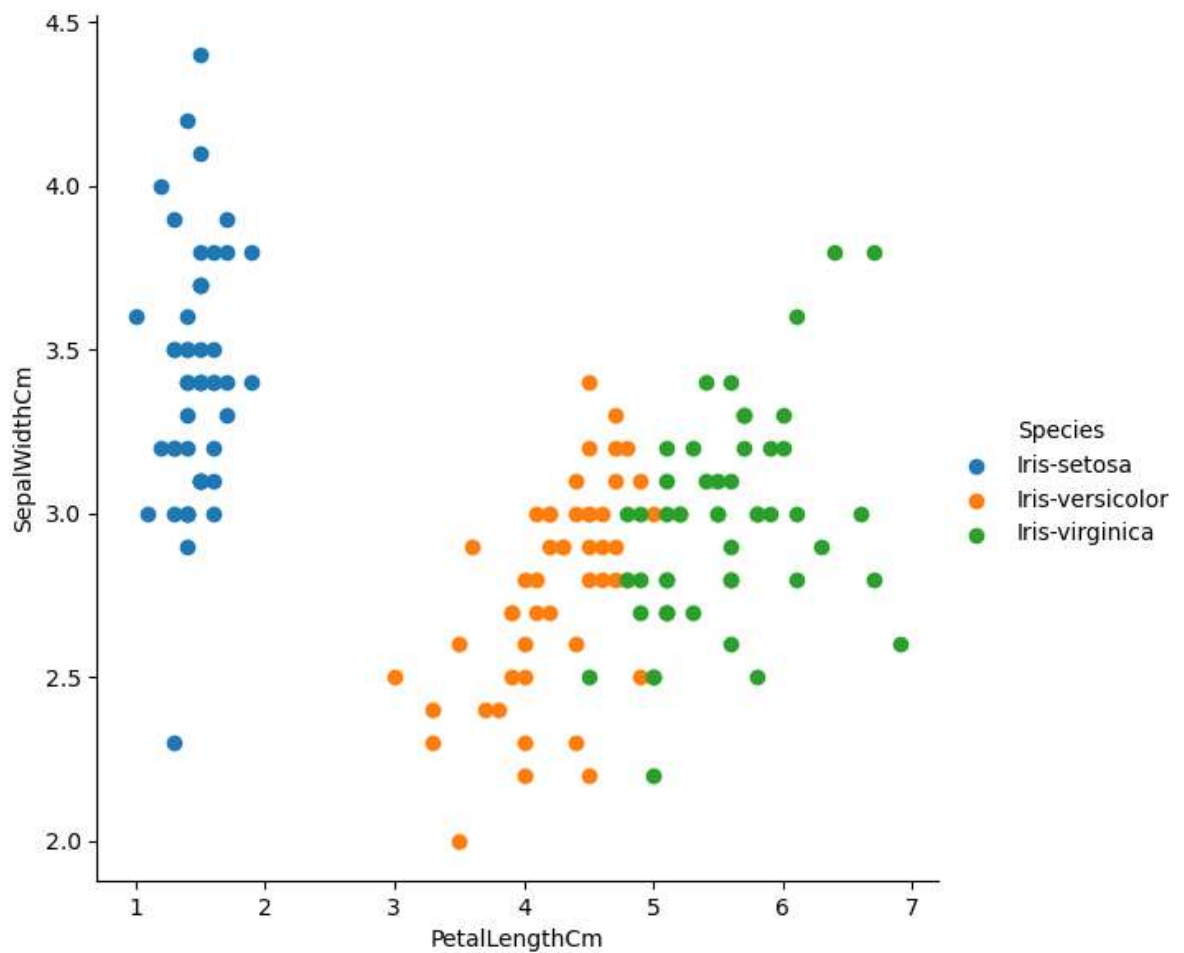
```
Out[7]: Iris-setosa      50
Iris-versicolor      50
Iris-virginica       50
Name: Species, dtype: int64
```

```
In [8]: print(iris.shape)

(150, 6)
```

```
In [9]: # Scatter Plot
sns.FacetGrid(iris, hue="Species",height=6).map(plt.scatter, "PetalLengthCm", "SepalLengthCm")
```

```
Out[9]: <seaborn.axisgrid.FacetGrid at 0x1be2de4ed50>
```



```
In [10]: # categorical variables into numbers
flower_mapping = {'Iris-setosa' : 0, 'Iris-versicolor' : 1, 'Iris-virginica' : 2}
iris["Species"] = iris["Species"].map(flower_mapping)
```

```
In [11]: iris.head()
```

```
Out[11]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	0
1	2	4.9	3.0	1.4	0.2	0
2	3	4.7	3.2	1.3	0.2	0
3	4	4.6	3.1	1.5	0.2	0
4	5	5.0	3.6	1.4	0.2	0

```
In [12]: #inputs and outputs
A = iris[['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm']].values
b = iris[['Species']].values
```

```
In [13]: # import algorithm
from sklearn.linear_model import LogisticRegression
```

```
In [14]: model = LogisticRegression()
```

```
In [15]: model.fit(A, b)
```

```
C:\ProgramData\anaconda3\Lib\site-packages\sklearn\utils\validation.py:1143: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n_samples, ), for example using ravel().
  y = column_or_1d(y, warn=True)
```

```
Out[15]: ▾ LogisticRegression
LogisticRegression()
```

```
In [16]: # Accuracy
model.score(A,b)
```

```
Out[16]: 0.9733333333333334
```

```
In [17]: # same input make prediction
expected = b
predicted = model.predict(A)
predicted
```

```
Out[17]: 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, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
        1, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 2, 1, 1, 1, 1, 1, 2, 1, 1, 1, 1,
        1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 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], dtype=int64)
```

```
In [18]: from sklearn import metrics
```

```
In [19]: # from the scatter plot the difference
print(metrics.classification_report(expected,predicted))
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	50
1	0.98	0.94	0.96	50
2	0.94	0.98	0.96	50
accuracy			0.97	150
macro avg	0.97	0.97	0.97	150
weighted avg	0.97	0.97	0.97	150

```
In [20]: # prediction 'Iris-setosa':100%, 'Iris-versicolor' : 97%, 'Iris-virginica' : 99% accuracy
print(metrics.confusion_matrix(expected,predicted))
```

```
[[50  0  0]
 [ 0 47  3]
 [ 0  1 49]]
```

```
In [25]: # This is test
A_new = np.array([[3,2,1,0.2],[4.9,2.2,3.8,1.1],[5.3,2.5,4.6,1.9]])
predicted = model.predict(A_new)
print("Predicted of Species: {}".format(predicted))
```

```
Predicted of Species: [0 1 2]
```