

Decision_tree_16_09_2025

September 16, 2025

```
[69]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
from sklearn.preprocessing import LabelEncoder
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import \
    classification_report, confusion_matrix, accuracy_score
from sklearn.tree import plot_tree
```

```
[37]: df = sns.load_dataset('iris')
df.head()
```

```
[37]:
```

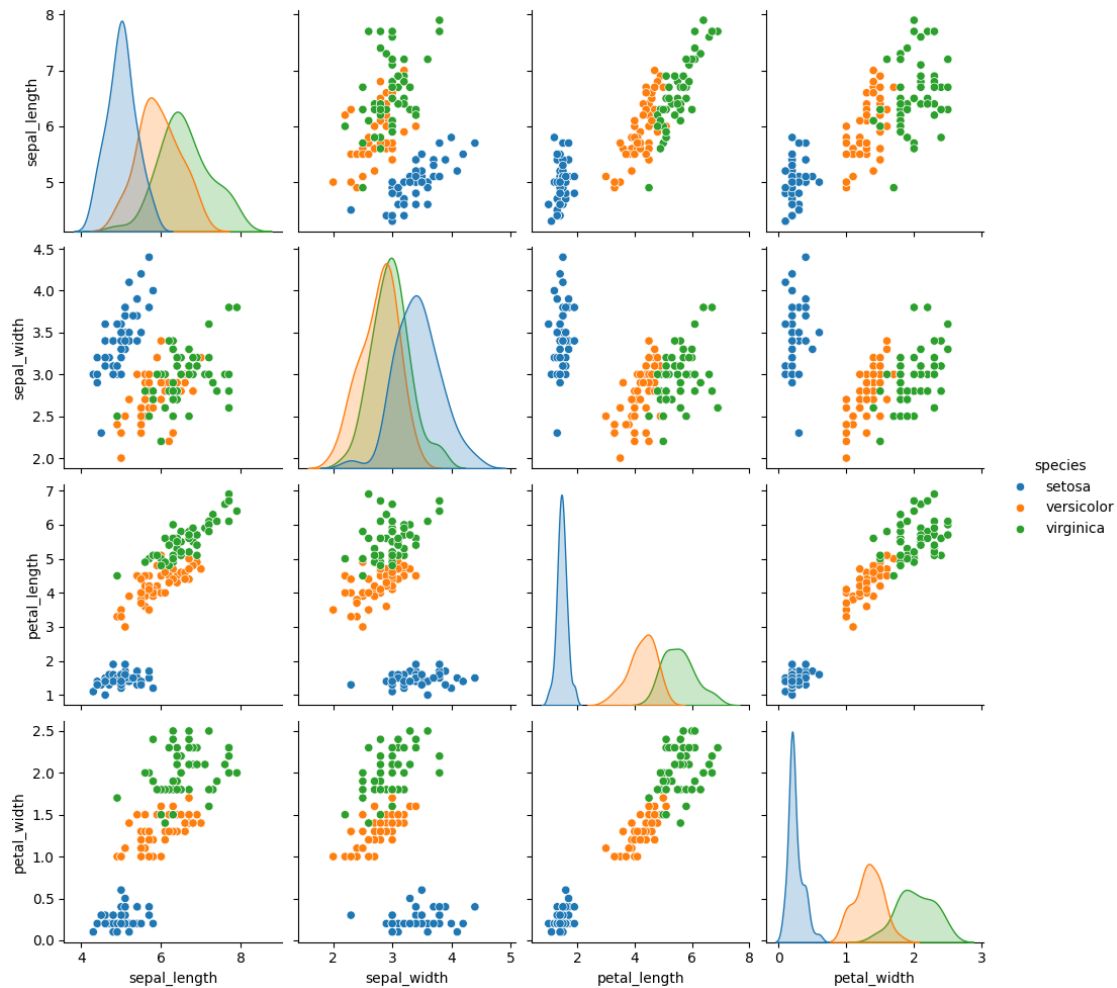
	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

```
[38]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype
---  -
0   sepal_length    150 non-null   float64
1   sepal_width     150 non-null   float64
2   petal_length    150 non-null   float64
3   petal_width     150 non-null   float64
4   species         150 non-null   object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

```
[39]: sns.pairplot(data = df ,hue='species')
```

[39]: <seaborn.axisgrid.PairGrid at 0x1e80e47a520>



```
[35]: df.isnull().sum()
```

```
[35]: sepal_length    0
      sepal_width    0
      petal_length   0
      petal_width    0
      species        0
      dtype: int64
```

```
[30]: sns.heatmap(df.drop('species',axis = 1).corr(),annot = True,cmap = 'Reds')
```

[30]: <Axes: >



```
[42]: trarget = df['species']
      df1 = df.copy()
      df1.drop('species',axis =1, inplace=True)
```

```
[45]: a = df1
      print(a.head())
```

```

      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
0      setosa
1      setosa
2      setosa
3      setosa
4      setosa
Name: species, dtype: object
```

```
[52]: le = LabelEncoder()
      trarget = le.fit_transform(trarget)
```

traget

```
[52]: 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, 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, 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])
```

```
[60]: b = traget
      b
```

```
[60]: 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, 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, 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])
```

```
[6]: x = df.iloc[:,1:4]
      x
```

```
[6]:      sepal_width  petal_length  petal_width
0           3.5           1.4           0.2
1           3.0           1.4           0.2
2           3.2           1.3           0.2
3           3.1           1.5           0.2
4           3.6           1.4           0.2
..      ...           ...           ...
145         3.0           5.2           2.3
146         2.5           5.0           1.9
147         3.0           5.2           2.0
148         3.4           5.4           2.3
149         3.0           5.1           1.8
```

```
[150 rows x 3 columns]
```

```
[7]: y = df.iloc[:, -1]
      y
```

```
[7]: 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

```

```

[61]: le = LabelEncoder()
      y = le.fit_transform(y)
      y

```

```

[61]: 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, 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, 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])

```

```

[64]: x_train,x_test,y_train,y_test = train_test_split(a,b,test_size = 0.
      ↪2,random_state = 42)#random_state is the produces the same sequence of the
      ↪numbers every time it is runned

```

```

[47]: x_train

```

```

[47]:      sepal_length  sepal_width  petal_length  petal_width
48           5.3           3.7           1.5           0.2
23           5.1           3.3           1.7           0.5
94           5.6           2.7           4.2           1.3
139          6.9           3.1           5.4           2.1
102          7.1           3.0           5.9           2.1
..          ...           ...           ...           ...
120          6.9           3.2           5.7           2.3
143          6.8           3.2           5.9           2.3
126          6.2           2.8           4.8           1.8
142          5.8           2.7           5.1           1.9
91           6.1           3.0           4.6           1.4

```

```

[120 rows x 4 columns]

```

```

[63]: y_train

```

```

[63]: array([0, 0, 1, 0, 0, 2, 1, 0, 0, 0, 2, 1, 1, 0, 0, 1, 2, 2, 1, 2, 1, 2,
           1, 0, 2, 1, 0, 0, 0, 1, 2, 0, 0, 0, 1, 0, 1, 2, 0, 1, 2, 0, 2, 2,
           1, 1, 2, 1, 0, 1, 2, 0, 0, 1, 1, 0, 2, 0, 0, 1, 1, 2, 1, 2, 2, 1,
           0, 0, 2, 2, 0, 0, 0, 1, 2, 0, 2, 2, 0, 1, 1, 2, 1, 2, 0, 2, 1, 2,
           1, 1, 1, 0, 1, 1, 0, 1, 2, 2, 0, 1, 2, 2, 0, 2, 0, 1, 2, 2, 1, 2,

```

```
1, 1, 2, 2, 0, 1, 2, 0, 1, 2])
```

```
[49]: x_test
```

```
[49]:
```

	sepal_length	sepal_width	petal_length	petal_width
101	5.8	2.7	5.1	1.9
63	6.1	2.9	4.7	1.4
106	4.9	2.5	4.5	1.7
22	4.6	3.6	1.0	0.2
115	6.4	3.2	5.3	2.3
95	5.7	3.0	4.2	1.2
104	6.5	3.0	5.8	2.2
56	6.3	3.3	4.7	1.6
90	5.5	2.6	4.4	1.2
59	5.2	2.7	3.9	1.4
111	6.4	2.7	5.3	1.9
130	7.4	2.8	6.1	1.9
110	6.5	3.2	5.1	2.0
80	5.5	2.4	3.8	1.1
149	5.9	3.0	5.1	1.8
4	5.0	3.6	1.4	0.2
41	4.5	2.3	1.3	0.3
122	7.7	2.8	6.7	2.0
3	4.6	3.1	1.5	0.2
46	5.1	3.8	1.6	0.2
62	6.0	2.2	4.0	1.0
45	4.8	3.0	1.4	0.3
38	4.4	3.0	1.3	0.2
105	7.6	3.0	6.6	2.1
31	5.4	3.4	1.5	0.4
73	6.1	2.8	4.7	1.2
69	5.6	2.5	3.9	1.1
67	5.8	2.7	4.1	1.0
40	5.0	3.5	1.3	0.3
136	6.3	3.4	5.6	2.4

```
[50]: y_test
```

```
[50]:
```

101	virginica
63	versicolor
106	virginica
22	setosa
115	virginica
95	versicolor
104	virginica
56	versicolor
90	versicolor

```

59     versicolor
111     virginica
130     virginica
110     virginica
80     versicolor
149     virginica
4       setosa
41      setosa
122     virginica
3       setosa
46      setosa
62     versicolor
45      setosa
38      setosa
105     virginica
31      setosa
73     versicolor
69     versicolor
67     versicolor
40      setosa
136     virginica
Name: species, dtype: object

```

```
[51]: dc = DecisionTreeClassifier()
```

```
[65]: dc.fit(x_train,y_train)
```

```
[65]: DecisionTreeClassifier()
```

```
[66]: y_pred = dc.predict(x_test)
```

```
[67]: print("Classification report : ",classification_report(y_test,y_pred))
```

```

Classification report :                precision    recall  f1-score   support

      0      1.00      1.00      1.00      10
      1      1.00      1.00      1.00       9
      2      1.00      1.00      1.00      11

 accuracy      1.00      1.00      1.00      30
 macro avg      1.00      1.00      1.00      30
weighted avg      1.00      1.00      1.00      30

```

```
[70]: print("Accuracy Report : ",accuracy_score(y_test,y_pred))
```

```
Accuracy Report : 1.0
```

```
[72]: cm=confusion_matrix
```

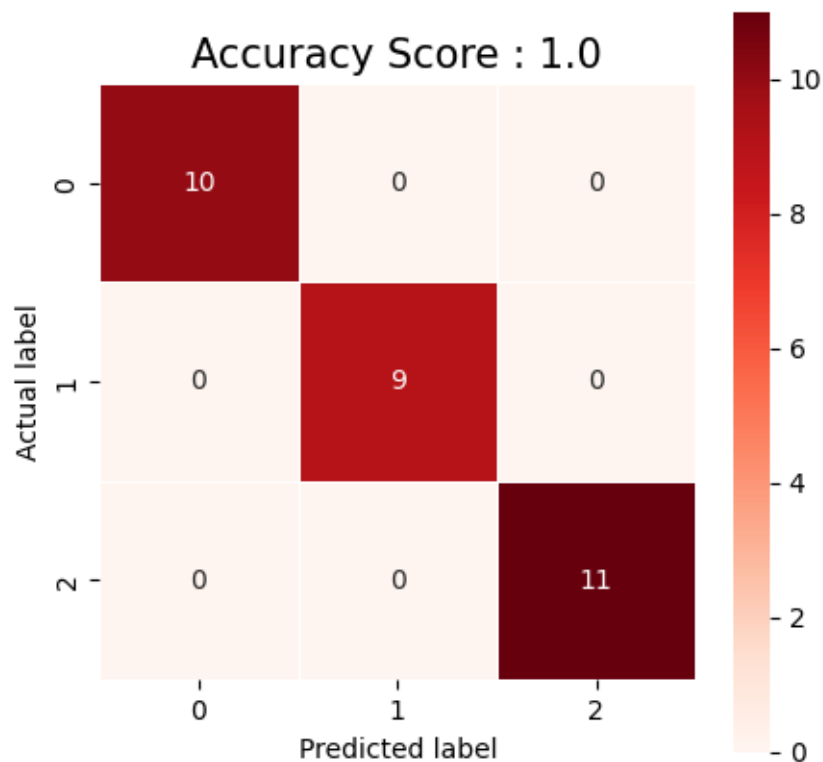
```
[73]: c = cm(y_test,y_pred)
```

```
[74]: c
```

```
[74]: array([[10,  0,  0],
          [ 0,  9,  0],
          [ 0,  0, 11]])
```

```
[76]: plt.figure(figsize=(5,5))
sns.heatmap(data = c,linewidth = .5,annot = True,square = True, cmap = 'Reds')
plt.ylabel('Actual label')
plt.xlabel('Predicted label')
all_sample_title = 'Accuracy Score : {0}'.format(dc.score(x_test,y_test))
plt.title(all_sample_title,size = 15)
```

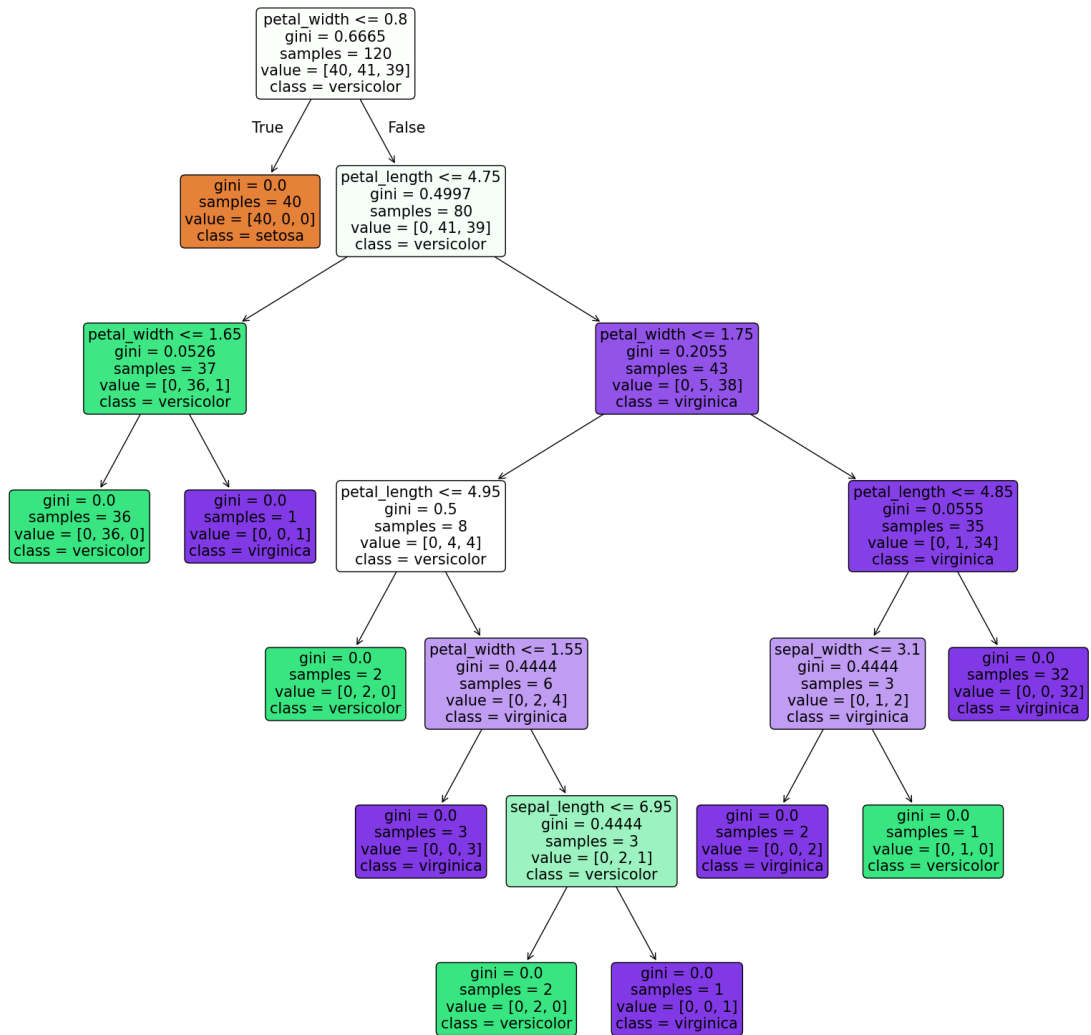
```
[76]: Text(0.5, 1.0, 'Accuracy Score : 1.0')
```



```
[77]: plt.figure(figsize = (20,20))
```



```
dec_tree = plot_tree(decision_tree=dc,feature_names = df1.columns,class_names =
↳['setosa','versicolor','virginica'],filled = True, precision = 4, rounded =
↳True)
```



[]: