▼ Decision Tree - 분류

```
import warnings
warnings.filterwarnings('ignore')
```

▼ 실습용 데이터 설정

iris.csv

```
import seaborn as sns

DF = sns.load_dataset('iris')
```

• pandas DataFrame

DF.info()

```
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
                  Non-Null Count Dtype
    Column
 0
   sepal_length 150 non-null
                                  float64
 1 sepal_width 150 non-null
                                  float64
 2 petal_length 150 non-null
                                  float64
    petal_width 150 non-null
                                  float64
    species
                  150 non-null
                                  object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

<class 'pandas.core.frame.DataFrame'>

DF.head(3)

	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

▼ I. 탐색적 데이터 분석

▼ 1) 빈도분석

DF.species.value_counts()

versicolor 50 virginica 50 setosa 50

Name: species, dtype: int64

▼ 2) 분포 시각화

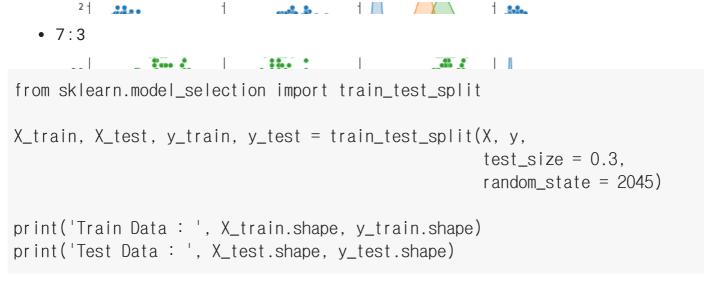
```
import matplotlib.pyplot as plt
import seaborn as sns
sns.pairplot(hue = 'species', data = DF)
plt.show()
```



II. Data Preprocessing

```
→ 1) Data Set
    y ·
                       X = DF[['sepal_length', 'sepal_width', 'petal_length', 'petal_width']]
y = DF['species']
      6 -
```

→ 2) Train & Test Split



Train Data: (105, 4) (105.) Test Data: (45, 4) (45,)

→ III. Modeling

▼ 1) Train_Data로 모델 생성

```
from sklearn.tree import DecisionTreeClassifier
Model_dt = DecisionTreeClassifier(random_state = 2045)
Model_dt.fit(X_train, y_train)
```

DecisionTreeClassifier(ccp_alpha=0.0, class_weight=None, criterion='gini', max_depth=None, max_features=None, max_leaf_nodes=None,

```
min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=1, min_samples_split=2, min_weight_fraction_leaf=0.0, presort='deprecated', random_state=2045, splitter='best')
```

→ 2) Visualization

```
petal length \leq 2.6
             gini = 0.666
            samples = 105
         value = [33, 36, 36]
           class = virginica
                           False
       True
                      petal length \leq 4.75
 gini = 0.0
                            gini = 0.5
samples = 33
                          samples = 72
```

▼ 3) Test_Data에 Model 적용

```
y_hat = Model_dt.predict(X_test)
y_hat
      array(['setosa', 'setosa', 'versicolor', 'virginica', 'setosa',
               'versicolor', 'virginica', 'setosa', 'virginica', 'setosa', 'versicolor', 'virginica', 'versicolor', 'setosa', 'setosa', 'versicolor', 'versicolor', 'setosa', 'versicolor', 'setosa',
                'setosa', 'setosa', 'virginica', 'setosa', 'virginica',
                'versicolor', 'setosa', 'versicolor', 'versicolor', 'virginica',
               'setosa', 'versicolor', 'setosa', 'versicolor', 'versicolor',
                'versicolor', 'virginica', 'versicolor', 'setosa', 'versicolor',
                'virginica', 'setosa', 'virginica', 'virginica', 'virginica'],
              dtype=object)
```

→ 4) Confusion Matrix

```
from sklearn.metrics import confusion_matrix
confusion_matrix(y_test, y_hat)
    array([[17, 0, 0],
          [0, 14, 0].
          [0, 2, 12]])
```

▼ 5) Accuracy, Precision, Recall

```
from sklearn.metrics import accuracy_score, precision_score, recall_score
print(accuracy_score(y_test, y_hat))
print(precision_score(y_test, y_hat, average = None))
nrint(recall score(v test v hat average = None))
```

printtroodii_oooroty_toot, y_nat, avorago nono,

```
0.955555555555556
[1. 0.875 1. ]
[1. 1. 0.85714286]
```

→ 6) F1_Score

▼ IV. Pruning(가지치기)

- min_samples_split : 분할을 위한 최소한의 샘플데이터 개수
- min_samples_leaf: 말단 노드가 되기 위한 최소한의 샘플데이터 개수
- max_leaf_nodes : 말단 노드의 최대 개수
- max_depth : 트리모델의 최대 깊이를 지정

→ 1) Model Pruning

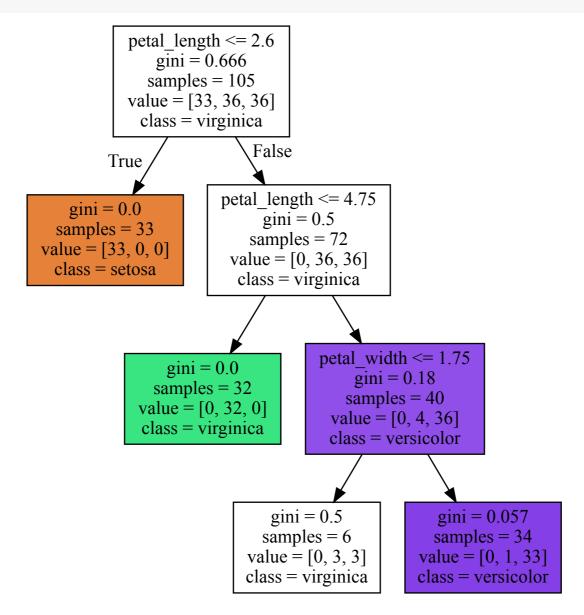
```
DecisionTreeClassifier(ccp_alpha=0.0, class_weight=None, criterion='gini', max_depth=3, max_features=None, max_leaf_nodes=None, min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=1, min_samples_split=2, min_weight_fraction_leaf=0.0, presort='deprecated', random_state=2045, splitter='best')
```

→ 2) Model Visualization

```
from sklearn.tree import export_graphviz import graphviz
```

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```
class_names = (['setosa', 'virginica', 'versi feature_names = (['sepal_length', 'sepal_widt filled = True))
```



→ 3) Model Evaluate

Confusion Matrix

```
from sklearn.metrics import confusion_matrix, accuracy_score, precision_score
y_hat = Model_pr.predict(X_test)
print(confusion_matrix(y_test, y_hat))

[[17 0 0]
       [0 14 0]
```

[0 2 12]]

· Accuracy, Precision, Recall

```
print(accuracy_score(y_test, y_hat))
print(precision_score(y_test, y_hat, average = None))
print(recall_score(y_test, y_hat, average = None))

0.955555555555556
[1. 0.875 1. ]
[1. 1. 0.85714286]
```

• F1-Score

```
f1_score(y_test, y_hat, average = None)
array([1. , 0.93333333, 0.92307692])
```

▼ V. Feature Importance

▼ 1) Feature Importance 값 확인

```
Model_pr.feature_importances_
array([0. , 0. , 0.96524977, 0.03475023])
```

▼ 2) Feature Importance 시각화

sepal_length -	
sepal_width -	

#

#

#

The End

#

#

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