▼ 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()

```
<class 'pandas.core.frame.DataFrame'>
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
                               float64
3 petal_width 150 non-null
4 species
                                object
             150 non-null
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

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

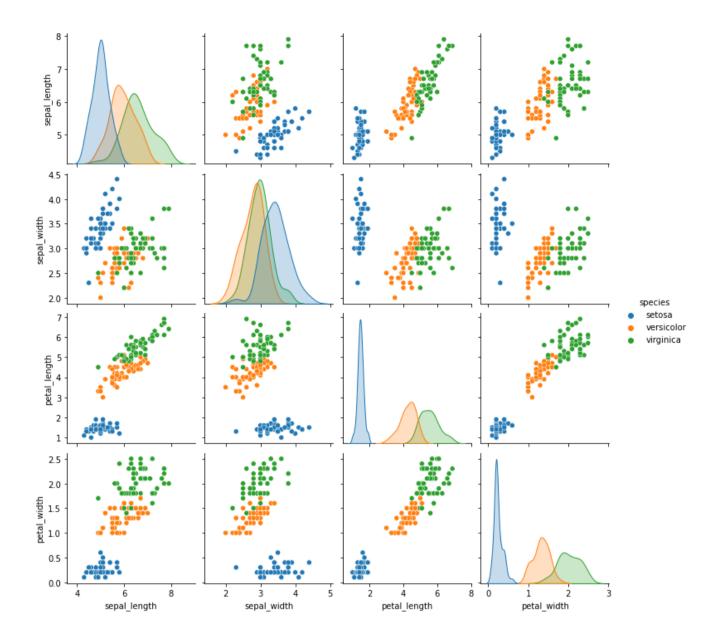
DF.species.value_counts()

```
setosa 50
virginica 50
versicolor 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

```
X = DF[['sepal_length', 'sepal_width', 'petal_length', 'petal_width']]
y = DF['species']
```

→ 2) Train & Test Split

• 7:3

→ III. Modeling

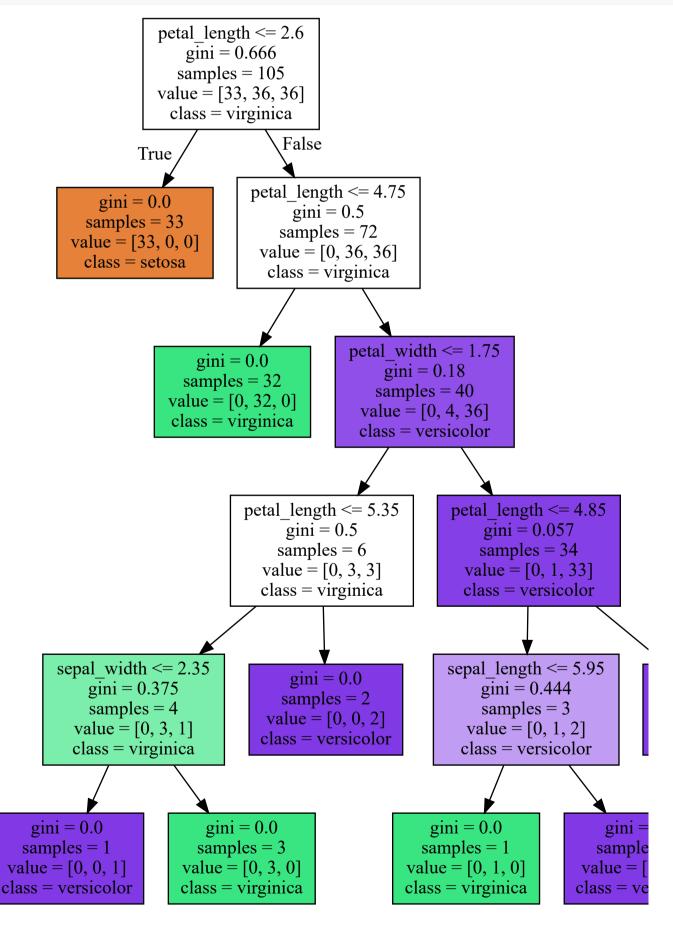
▼ 1) Train_Data로 모델 생성

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



▼ 3) Test_Data에 Model 적용

```
y_hat
```

→ 4) Confusion Matrix

→ 5) Accuracy, Precision, Recall

→ 6) F1_Score

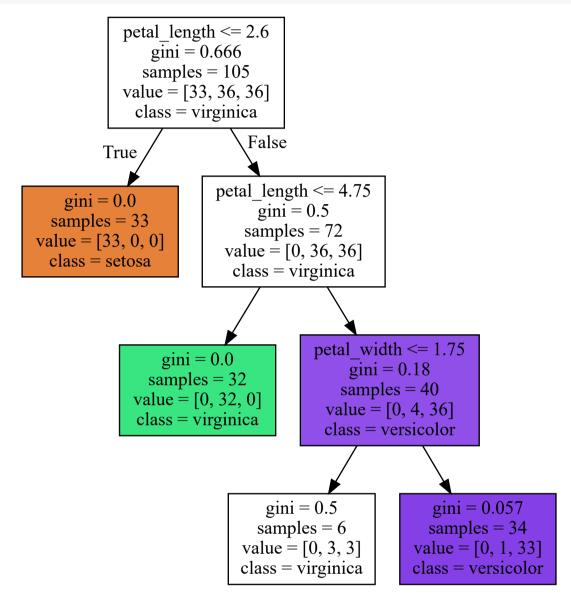
▼ IV. Pruning(가지치기)

- 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,

→ 2) Model Visualization



→ 3) Model Evaluate

Confusion Matrix

```
from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_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

```
f1_score(y_test, y_hat, average = None)

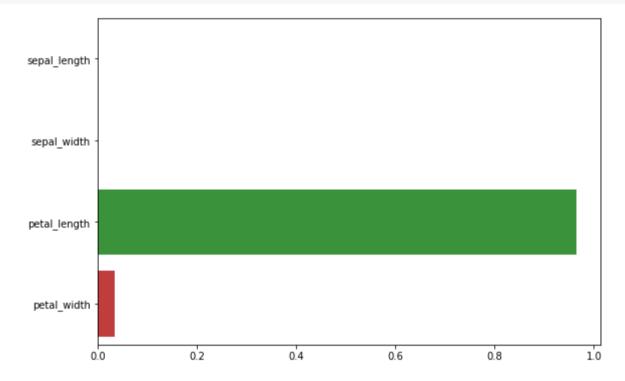
array([1. , 0.93333333, 0.92307692])
```

→ V. Feature Importance

▼ 1) Feature Importance 값 확인

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

▼ 2) Feature Importance 시각화



#

#

The End

#

#

#