▼ Random Forest - 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

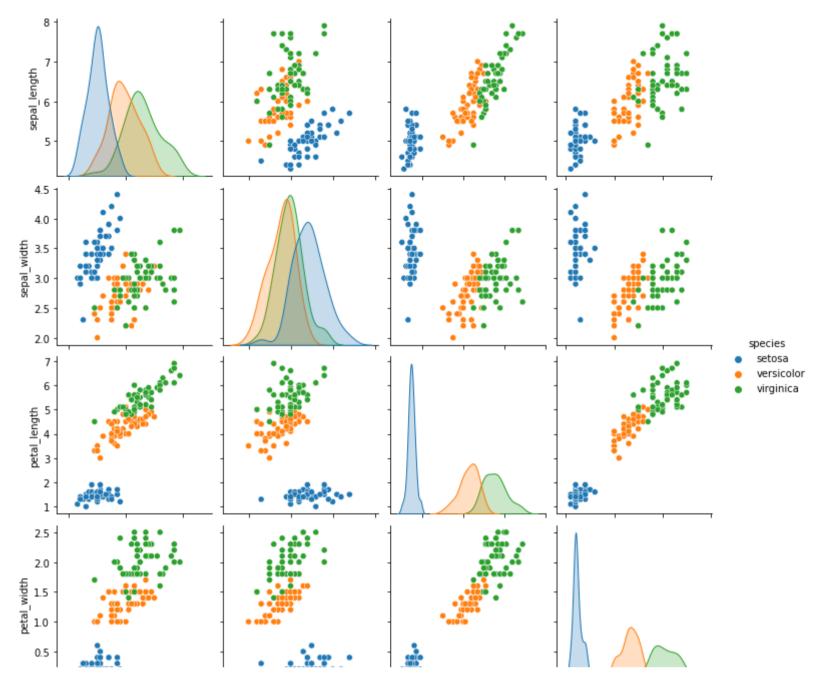
DF.species.value_counts()

```
virginica 50
setosa 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

• random_state: 반복 실행 시 동일한 결과 출력

• n_jobs : 모든 CPU 코어 사용

RandomForestClassifier(bootstrap=True, ccp_alpha=0.0, class_weight=None, criterion='gini', max_depth=None, max_features=2, max_leaf_nodes=None, max_samples=None, min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=1, min_samples_split=2, min_weight_fraction_leaf=0.0, n_estimators=10, n_jobs=-1, oob_score=False, random_state=2045, verbose=0, warm_start=False)

Colab CPU check

```
!cat /proc/cpuinfo | grep 'model name'
```

```
model name : Intel(R) Xeon(R) CPU @ 2.20GHz model name : Intel(R) Xeon(R) CPU @ 2.20GHz
```

→ 2) Test_Data에 Model 적용

```
y_hat = Model_rf.predict(X_test)
```

→ 3) Model Evaluate

```
from sklearn.metrics import confusion_matrix, accuracy_score
print(confusion_matrix(y_test, y_hat))

[[17 0 0]
      [0 14 0]
      [0 2 12]]
```

print(accuracy_score(y_test, y_hat))

0.95555555555556

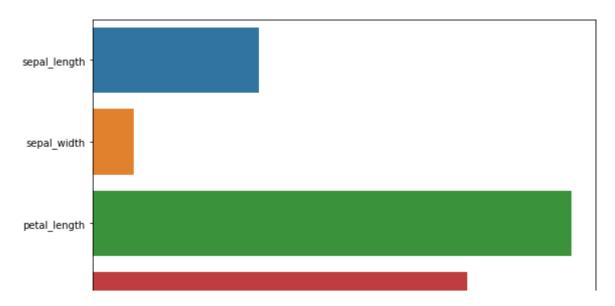
4) Feature Importance

• Feature Importance 값 확인

```
Model_rf.feature_importances_
```

```
array([0.1571031 , 0.03897972, 0.45102744, 0.35288974])
```

Feature Importance 시각화



▼ IV. Hyperparameter Tuning

• n_estimators : 모델에 사용되는 의사결정나무의 개수

• max_features : 분할에 사용되는 Feature의 개수

• max_depth : 트리모델의 최대 깊이를 지정

• max_leaf_nodes : 말단 노드의 최대 개수

• min_samples_split : 분할을 위한 최소한의 샘플데이터 개수

• min_samples_leaf : 말단 노드가 되기 위한 최소한의 샘플데이터 개수

▼ 1) RandomForestClassifier 객체 생성

```
from sklearn.ensemble import RandomForestClassifier

Model_rf = RandomForestClassifier()
```

▼ 2) GridSearchCV Hyperparameters 설정

▼ 3) GridSearchCV 객체 생성

• 5-Fold Cross Validation

▼ 4) GridSearchCV 수행

• 약 3분

```
from datetime import datetime
start_time = datetime.now()
grid cv fit(X train v train)
```

```
end_time = datetime.now()
print('Elapsed Time: ', end_time - start_time)

Elapsed Time: 0:03:16.822999

* 5) 최적 Hyperparameter 확인

• Best Accuracy
```

```
grid_cv.best_score_
```

0.9523809523809523

• Best Hyperparameter

```
grid_cv.best_params_
```

```
▼ 6) 최적 모델 생성 및 평가
```

{'max_depth': 3, 'max_features': 1, 'n_estimators': 100, 'random_state': 2045}

Best Model

```
Model_CV = grid_cv.best_estimator_
```

Evaluation

```
y_hat = Model_CV.predict(X_test)

from sklearn.metrics import confusion_matrix, accuracy_score

print(confusion_matrix(y_test, y_hat))

[[17 0 0]
      [0 13 1]
      [0 1 13]]
```

```
print(accuracy_score(y_test, y_hat))
```

0.95555555555556

#

#

#

The End

#

#

#