

붓꽃(Iris) 품종 분류

데이터 입력

https://en.wikipedia.org/wiki/Iris_flower_data_set

```
import seaborn as sns
iris = sns.load_dataset("iris")
# data.to_csv("dataset.csv")
print(iris.head())
# print(iris.head(n = 3))
# print(iris.tail(n = 3))
# print(iris.shape)
```

Setosa



Versicolor



Verginica



붓꽃 품종

| | 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 |

seaborn.load_dataset

`seaborn.load_dataset` (*name*, *cache=True*, *data_home=None*, ***kws*)

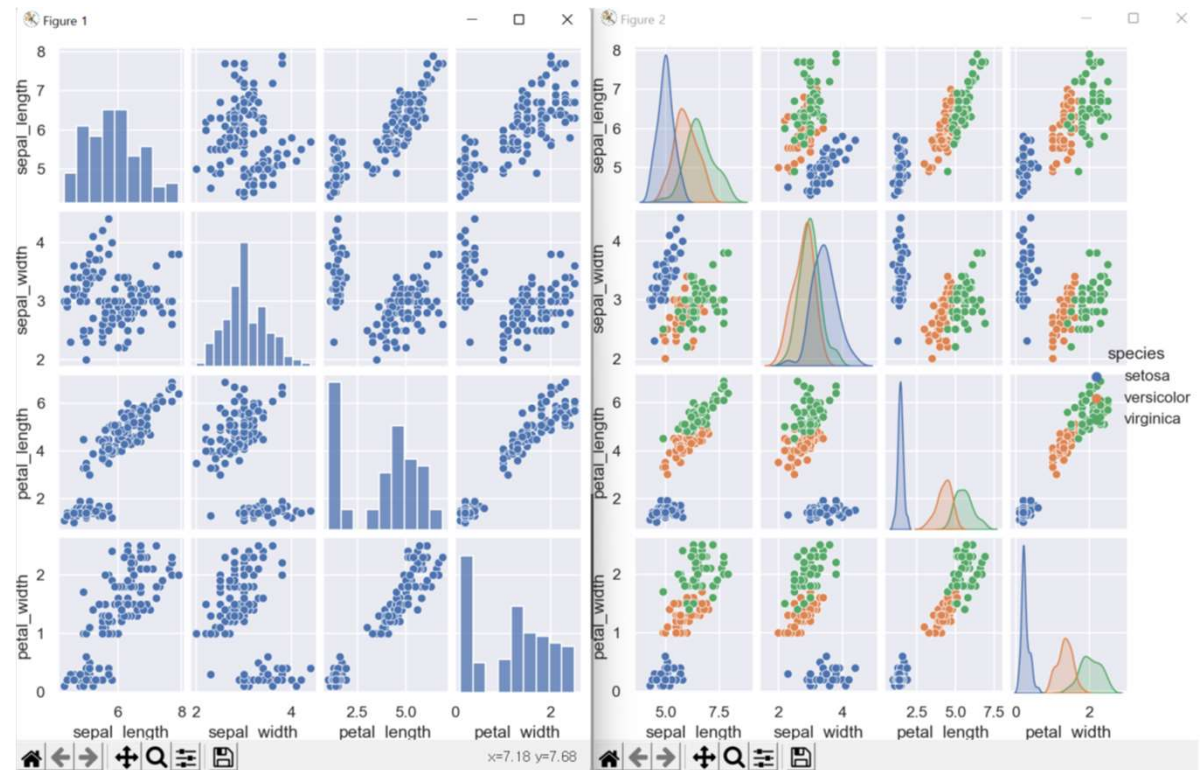
Load an example dataset from the online repository (requires internet).

https://seaborn.pydata.org/generated/seaborn.load_dataset.html#seaborn.load_dataset

데이터 시각화

```
sns.set_theme()
sns.pairplot(iris)
sns.pairplot(iris, hue="species")

import matplotlib.pyplot as plt
plt.show()
```



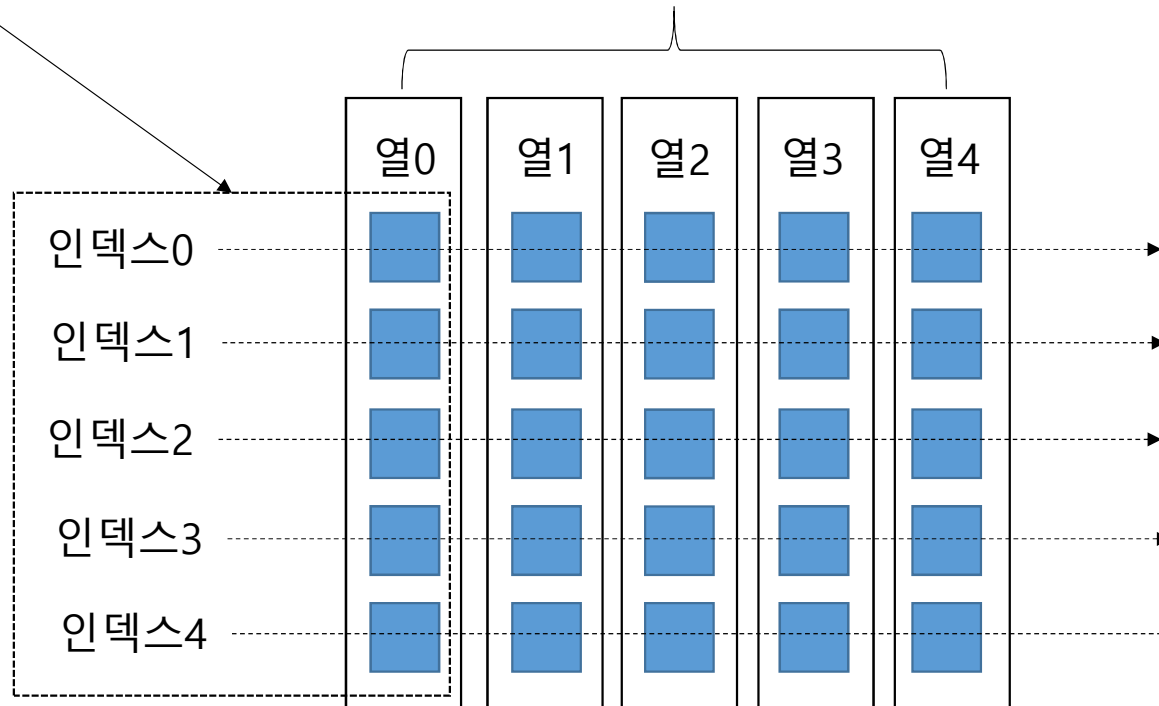
seaborn.pairplot

`seaborn.pairplot` (*data*, *, *hue=None*, *hue_order=None*, *palette=None*, *vars=None*, *x_vars=None*, *y_vars=None*, *kind='scatter'*, *diag_kind='auto'*, *markers=None*, *height=2.5*, *aspect=1*, *corner=False*, *dropna=False*, *plot_kws=None*, *diag_kws=None*, *grid_kws=None*, *size=None*)

Plot pairwise relationships in a dataset.

판다스 자료구조

- 행 방향으로는 행 인덱스, 열 방향으로는 열 이름으로 구성하는 데이터프레임(DataFrame)
- 시리즈(Series)는 데이터 값의 1차원 벡터, 데이터프레임은 2차원 구조를 의미



pandas.DataFrame

```
class pandas.DataFrame(data=None, index=None, columns=None, dtype=None, copy=None)
```

Two-dimensional, size-mutable, potentially heterogeneous tabular data.

[\[source\]](#)

Data structure also contains labeled axes (rows and columns). Arithmetic operations align on both row and column labels. Can be thought of as a dict-like container for Series objects. The primary pandas data structure.

<https://numpy.org/doc/stable/reference/arrays.ndarray.html>

```
>>> d = {'col1': [1, 2], 'col2': [3, 4]}
>>> df = pd.DataFrame(data=d)
>>> df
   col1  col2
0     1     3
1     2     4
```

```
>>> df2 = pd.DataFrame(np.array([[1, 2, 3], [4, 5, 6], [7, 8, 9]]),
...                     columns=['a', 'b', 'c'])
>>> df2
   a  b  c
0  1  2  3
1  4  5  6
2  7  8  9
```

```
>>> d = {'col1': [0, 1, 2, 3], 'col2': pd.Series([2, 3], index=[2, 3])}
>>> pd.DataFrame(data=d, index=[0, 1, 2, 3])
   col1  col2
0     0   NaN
1     1   NaN
2     2   2.0
3     3   3.0
```

<https://pandas.pydata.org/docs/reference/api/pandas.DataFrame.html#pandas.DataFrame>

데이터프레임 접근

```
▶ player= ["Mbappe", "Haaland", "Salah", "Messi"]  
country = ["France", "Norway", "Egypt", "Argentina"]  
dict_data = {"player": player, "country": country}
```

```
import pandas as pd  
df = pd.DataFrame(dict_data)  
print(df)
```

| | player | country |
|---|---------|-----------|
| 0 | Mbappe | France |
| 1 | Haaland | Norway |
| 2 | Salah | Egypt |
| 3 | Messi | Argentina |

```
[18] df.columns = ["Top-Player", "Nationality"]  
df.index = ["1st", "2nd", "3rd", "4th"]  
print(df)
```

| | Top-Player | Nationality |
|-----|------------|-------------|
| 1st | Mbappe | France |
| 2nd | Haaland | Norway |
| 3rd | Salah | Egypt |
| 4th | Messi | Argentina |

```
▶ print(df.Nationality)
```

```
↳ 1st      France  
   2nd      Norway  
   3rd      Egypt  
   4th      Argentina  
   Name: Nationality, dtype: object
```

```
▶ print(df.loc["2nd", "Nationality"])  
print(df.loc[:, "Top-Player"])
```

```
Norway  
1st      Mbappe  
2nd      Haaland  
3rd      Salah  
4th      Messi  
   Name: Top-Player, dtype: object
```

```
[23] print(df.loc["2nd"])
```

```
Top-Player      Haaland  
Nationality      Norway  
   Name: 2nd, dtype: object
```

```
[24] print(df.iloc[1])
```

```
Top-Player      Haaland  
Nationality      Norway  
   Name: 2nd, dtype: object
```

```
▶ print(df.iloc["2nd"])
```


데이터 전처리

```
data = iris.drop("species", axis = 1)
# data = iris.drop(columns = "species")
# print(data.head())
# print(data.shape)
# print(data.dtypes)
t = iris[["species"]].copy()
# t = iris[["species"]]
# print(t["species"].unique())
t[t["species"] == "setosa"] = 0
# t.loc[t.species == "setosa", "species"] = 0
# t["species"] = t["species"].replace(["setosa"], 0)
t[t["species"] == "versicolor"] = 1
t[t["species"] == "virginica"] = 2
# print(t["species"].unique())
t = t["species"].astype("int")
# print(t.dtypes)
```

```
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
(150, 4)
sepal_length    float64
sepal_width     float64
petal_length    float64
petal_width     float64
```



| Setosa | Versicolor | Verginica |
|--------|------------|-----------|
| 0 | 1 | 2 |

```
['setosa' 'versicolor' 'virginica']
[0 1 2]
```

pandas.DataFrame.drop

```
DataFrame.drop(labels=None, axis=0, index=None, columns=None, level=None, inplace=False, errors='raise')
```

[source]

Drop specified labels from rows or columns.

Parameters: labels : single label or list-like

Index or column labels to drop. A tuple will be used as a single label and not treated as a list-like.

axis : {0 or 'index', 1 or 'columns'}, default 0

Whether to drop labels from the index (0 or 'index') or columns (1 or 'columns').

index : single label or list-like

Alternative to specifying axis (labels, axis=0 is equivalent to index=labels).

columns : single label or list-like

Alternative to specifying axis (labels, axis=1 is equivalent to columns=labels).

```
>>> df = pd.DataFrame(np.arange(12).reshape(3, 4),
...                    columns=['A', 'B', 'C', 'D'])
>>> df
   A  B  C  D
0  0  1  2  3
1  4  5  6  7
2  8  9 10 11
```

```
>>> df.drop(['B', 'C'], axis=1)
   A  D
0  0  3
1  4  7
2  8 11

>>> df.drop(columns=['B', 'C'])
   A  D
0  0  3
1  4  7
2  8 11
```


KNN

```
from sklearn.model_selection import train_test_split
train_data, test_data, train_target, test_target = train_test_split(
    data, t, test_size = 0.3, random_state = 42, stratify = t)
# print(train_data.shape)      (105, 4)
# print(test_data.shape)       (45, 4)
# print(train_target.shape)    (105,)
# print(test_target.shape)     (45,)

from sklearn.neighbors import KNeighborsClassifier
kn = KNeighborsClassifier()
kn.fit(train_data, train_target)
print("Train-Eval:", kn.score(train_data, train_target)) } Train-Eval: 0.9714285714285714
print("Test-Eval :", kn.score(test_data, test_target)) } Test-Eval : 0.9777777777777777
```

```
from sklearn.metrics import confusion_matrix
conf = confusion_matrix(test_target, kn.predict(test_data))
print(conf)
```

```
[[15  0  0]
 [ 0 15  0]
 [ 0  1 14]]
```

| | 예측0 | 예측1 | 예측2 |
|-----|-----|-----|-----|
| 정답0 | 15개 | 0개 | 0개 |
| 정답1 | 0개 | 15개 | 0개 |
| 정답2 | 0개 | 1개 | 14개 |

sklearn.metrics.confusion_matrix

```
sklearn.metrics.confusion_matrix(y_true, y_pred, *, labels=None, sample_weight=None, normalize=None)
```

Compute confusion matrix to evaluate the accuracy of a classification.

Parameters:

- y_true : array-like of shape (n_samples,)**
Ground truth (correct) target values.
- y_pred : array-like of shape (n_samples,)**
Estimated targets as returned by a classifier.

혼합행렬(Confusion Matrix):

원 클래스와 예측 클래스의
일치 여부를 나타내는 행렬

```
>>> from sklearn.metrics import confusion_matrix
>>> y_true = [2, 0, 2, 2, 0, 1]
>>> y_pred = [0, 0, 2, 2, 0, 2]
>>> confusion_matrix(y_true, y_pred)
array([[2, 0, 0],
       [0, 0, 1],
       [1, 0, 2]])
```

```
>>> y_true = ["cat", "ant", "cat", "cat", "ant", "bird"]
>>> y_pred = ["ant", "ant", "cat", "cat", "ant", "cat"]
>>> confusion_matrix(y_true, y_pred, labels=["ant", "bird", "cat"])
array([[2, 0, 0],
       [0, 0, 1],
       [1, 0, 2]])
```

참고자료

- 지능기전공학부 최유경 교수님 자료, <https://github.com/sejongresearch/2021.MachineLearning>
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