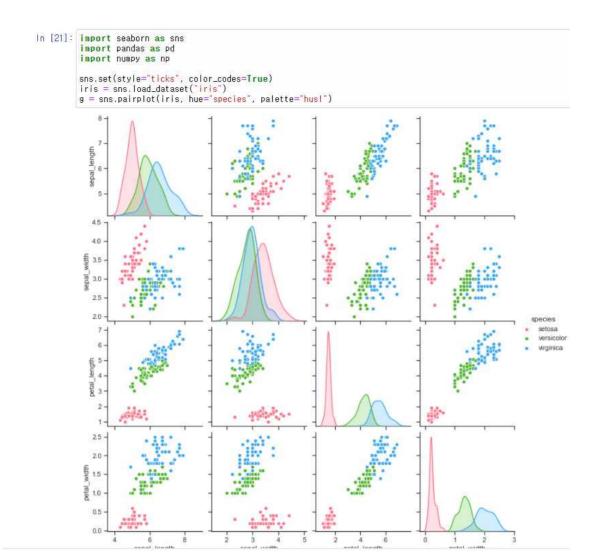
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케라스를 이용한 붓꽃 분류

```
In [30]: #import libraries
                import os
                import pandas as pd
                import matplotlib.pyplot as plt
               #Set working directory and load data os.ohdir('C:/PycharmProjects/')
                iris =pd.read_osv('iris.osv')
               print(iris.head())
               #Create numerio olasses for species(0,1,2)
iris.loo[iris['species']='virginioa', 'species']=0
iris.loo[iris['species']='versicolor', 'species']=1
iris.loo[iris['species']='setosa', 'species']=2
iris = iris[iris['species']!=2]
                #Create input and output oclums
               X = iris[['sepal_length', 'sepal_width']].values.T
Y = iris[['species']].values.T
Y=Y.astype('uint8')
              #Make a souther plot
plt.souther(X[0,:], X[1,:],o=Y[0,:],s=40, omap=plt.om.Spectral);
plt.title("IRIS DATA | Blue - Yersicolor, Red - Virginica")
plt.xlabel("petal_length")
plt.ylabel("petal_width")
               plt.show()
                     sepal_length sepal_width petal_length petal_width species
                                                                                                   0.2 setosa
0.2 setosa
               0
                                   5.1
                                                        3.5
                                                                                1.4
                                   4.9
                                                         3.0
                                                                                1.4
                                                                                                            setosa
               23
                                                        3.2
                                                                               1.3
                                                                                                    0.2 setosa
                                                                                                    0.2 setosa
0.2 setosa
                                                        3.1
                                                                                1.5
                                   4.6
                                   5.0
                                                        3.6
                                                                                1.4
                                       IRIS DATA | Blue - Yersicolor, Red - Virginica
                     3.75 -
                     3.50
                     3.25
                     3.00 -
                 275
                     2.50
                     2.25
                     2.00 -
                                5.0
                                            5.5
                                                        6.0
                                                                   6.5
                                                                               7.0
                                                                                            7.5
                                                                                                        8.0
                                                            petal_length
```



```
In [22]: Iris.info()

cclass 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149

Data columns (total 5 columns):

# Column Non-Null Count Dtype

0 sepal.elength 150 non-null float64
1 sepal.withh 150 non-null float64
2 sepal.withh 150 non-null float64
4 species 150 non-null object diverse: float64(4), object(1)
memory usage: 6,0 kB

In [23]: Iris['species'].unique()

Out[23]: array(['setosa', 'versicolor', 'virginica'], dtype=object)

In [24]: from sklearn.preprocessing import LabelEncoder

X = iris.iloc[:,0:4].values
encoder = LabelEncoder()
y1 = encoder.fit.transform(y)
y = encoder.fit.transform(y)
y = d.get.dumies(y1).values

v = d.get.dumies(y1).values

10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10, 0, 11, 10
```

```
In [25]: from sklearn.model_selection import train_test_split
          X_train, X_test, y_train, y_test = train_test_split(X, Y,
                                                               test_size=0.2.
                                                               random_state=1)
         X_train.shape, X_test.shape, y_train.shape, y_test.shape
Out [25]: ((120, 4), (30, 4), (120, 3), (30, 3))
In [26]: from keras, models import Sequential
          from keras.layers import Dense
          from keras.optimizers import Adam
          model = Sequential()
          model.add(Dense(64,input_shape=(4,),activation='relu'))
          model.add(Dense(64,activation='relu'))
          model.add(Dense(3,activation='softmax'))
          model.compile(loss='categorical_crossentropy',
                        optimizer='Adam',
metrics=['accuracy'])
          model.summary()
```

Model: "sequential_2"

Layer (type)	Output Shape	Param #
dense_4 (Dense)	(None, 64)	320
dense_5 (Dense)	(None, 64)	4160
dense_6 (Dense)	(None, 3)	195

Total params: 4,675 Trainable params: 4,675 Non-trainable params: 0

```
In [29]: import matplotlib.pyplot as plt

plt.figure(figs[ze=(12,8))
    plt.plot(hist.history['loss'])
    plt.plot(hist.history['val_loss'])
    plt.plot(hist.history['val_accuracy'])
    plt.legend(['loss', 'val_loss', 'accuracy', 'val_accuracy'])
    plt.grid()
    plt.show()

14

12

10

08
```

60

In [1:

0.6

0.4

0.2