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```
#library import - tensorflow/keras/sklearn/matplotlib
import os
from os.path import join

import numpy as np
import pandas as pd

import tensorflow as tf
import keras
from keras import layers, Input, models
from tensorflow.keras.utils import to_categorical
from keras.wrappers.scikit_learn import KerasClassifier

from sklearn.model_selection import KFold
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import train_test_split
import matplotlib.pyplot as plt
```

```
In []:
#dataset 구성
datapath = join('data', 'wafer')
url = "E:/WaferMap"
#print(os.listdir(url)) #dir list

#경고 메시지 끄기(ignore), 켜기(default)
import warnings
warnings.filterwarnings("ignore")

#read pickle 파일
#https://www.kaggle.com/code/kcs93023/keras-wafer-classification-cnn2d-with-augmentation/data
df=pd.read_pickle(url+"/LSWMD.pkl")
```

pickle 파일구성 (이미 학습할 수 있도록, 분류가 된 파일)

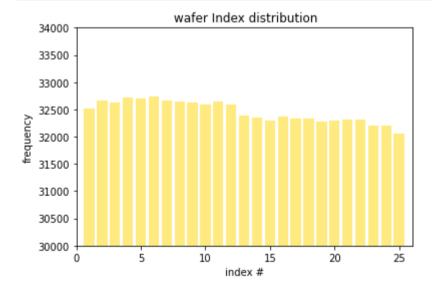
• waferMap file 개수: 811.457 rows(2차원 행렬 data, 행*열=waferMapDim에 있음)

```
데이터 구성
           # none label: 638,507 - 78.7%
           # label: 172,950 - 21.3%
              - ng: 25,519
                            14.8%(3.1%) --> 불량 항목 별로도 데이터 불균형
              - ok: 147,431 85.2%(18.2%)
In [ ]:
         #data file 정보
         df.info()
In [ ]:
        #해더 내용
         df.head()
                             waferMap dieSize lotName trianTestLabel failureType waferMapDim
Out[ ]:
                                                 lot1
        [[Training]]
                                                                    [[none]]
                                                                                 (45, 48)
        lot1
                                                        [[Training]]
                                                                                 (45, 48)
                                                                    [[none]]
        lot1
                                                        [[Training]]
                                                                    [[none]]
                                                                                 (45, 48)
        lot1
                                                        [[Training]]
                                                                    [[none]]
                                                                                 (45, 48)
        lot1
                                                        [[Training]]
                                                                    [[none]]
                                                                                 (45, 48)
In [ ]:
        #tail 내용
         df.tail()
                                  waferMap dieSize lotName trianTestLabel
Out[ ]:
                                                                      failureType waferMapDim
                                            600.0 lot47542
                                                                                      (26, 30)
        811452 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 2, 1, 1,...
                                                                [[Test]]
                                                                      [[Edge-Ring]]
                                            600.0 lot47542
        811453 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 2, 1, 1,...
                                                                [[Test]]
                                                                      [[Edge-Loc]]
                                                                                      (26, 30)
        811454 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 2, 1, 1,...
                                            600.0 lot47542
                                                                [[Test]] [[Edge-Ring]]
                                                                                      (26, 30)
                                            600.0 lot47543
                                                                   П
                                                                                      (26, 30)
        811455 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, ...
```

• 6개 컬럼(wafer map, die size, lot name, index, T/T type, failType) --> failuretype: 불량 유형 (none, edge-ring...)

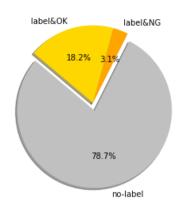
```
In [ ]:
         #csv 파일로도 한번 받아 봄(583mb)
         df.to_csv(path_or_buf = 'wafermap.csv', \text{\text{\text{$\psi}}}
                    sep=str(','), header=True, index=True, mode='w', encoding='CP949')
In [ ]:
         # 1lot = 25장 wafer인데(1lot / wafer index = 1~25)
         # lot의 wafer index 별로 보면, 없는 데이터도 존재 함
         import matplotlib.pyplot as plt
         uni_Index=np.unique(df.waferIndex, return_counts=True)
         plt.bar(uni_Index[0],uni_Index[1], color='gold', align='center', alpha=0.5)
         plt.title("wafer Index distribution")
         plt.xlabel("index #")
         plt.ylabel("frequency")
         plt.xlim(0.26)
         plt.ylim(30000,34000)
         plt.show()
```

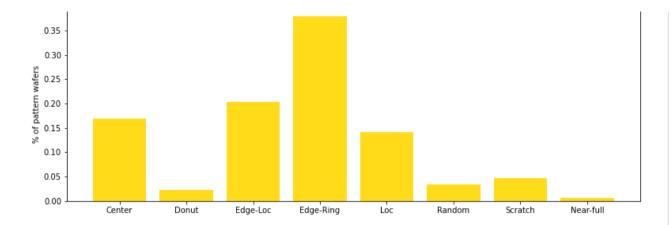
(26, 30)



```
In []: #데이터 분포 라벨링율 = 21.3%
#데이터 분포 ok : ng = 85.2:14.8
#데이터 분포 pg(detection) 내 의 code병 분포
```

```
#41014 44 114146166110117 41 4 60462 44
df['failureNum']=df.failureType #컬럼 추가
df['trainTestNum']=df.trianTestLabel
                                     #컬럼 추가
mapping type={'Center':0.'Donut':1.'Edge-Loc':2.'Edge-Ring':3.'Loc':4.'Random':5.'Scratch':6.'Near-full':7.'none':8}
                                                                                                                     #불
mapping traintest={'Training':0.'Test':1} #train:0. test:1 로 라벨링
df=df.replace({'failureNum':mapping_type, 'trainTestNum':mapping_traintest})
                                                                              #replace
#failureType의 label이 존재
df_withlabel = df[(df['failureNum']>=0) & (df['failureNum']<=8)]</pre>
df withlabel = df withlabel.reset index()
#failureType의 label이 존재 && 불량z코드 존재 (0~7번 코드)
df_withpattern = df[(df['failureNum']>=0) & (df['failureNum']<=7)]</pre>
df withpattern = df withpattern.reset index()
#failureType의 label이 존재 && OK품(8번 코드. none)
df_nonpattern = df[(df['failureNum']==8)]
df_withlabel.shape[0], df_withpattern.shape[0], df_nonpattern.shape[0]
#graph - pie
from matplotlib import gridspec
fig = plt.figure(figsize=(20, 4.5))
gs = gridspec.GridSpec(1, 2, width_ratios=[1, 2.5])
ax1 = plt.subplot(gs[0])
ax2 = plt.subplot(gs[1])
tol_wafers = df.shape[0]
no_wafers=[tol_wafers-df_withlabel.shape[0], df_withpattern.shape[0], df_nonpattern.shape[0]]
colors = ['silver', 'orange', 'gold']
explode = (0.1, 0, 0) # pie에서 sclicing해서 차트에서 벗어나게
labels = ['no-label', 'label&NG', 'label&OK']
ax1.pie(no_wafers, explode=explode, labels=labels, colors=colors, autopct='%1.1f%%', shadow=True, startangle=140)
#graph - bar
uni_pattern=np.unique(df_withpattern.failureNum, return_counts=True)
labels2 = [''.'Center'.'Donut'.'Edge-Loc'.'Edge-Ring'.'Loc'.'Random'.'Scratch'.'Near-full']
ax2.bar(uni_pattern[0],uni_pattern[1]/df_withpattern.shape[0], color='gold', align='center', alpha=0.9)
ax2.set_title("failure type frequency")
ax2.set_ylabel("% of pattern wafers")
ax2.set_xticklabels(labels2)
plt.show()
```





```
In [ ]: #df head()
    df[(df['failureNum']==8)].head()
```

Out[]:		waferMap	dieSize	lotName	trianTestLabel	failureType	waferMapDim	failureNum	trainTestNum	
	0	[[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0	1683.0	lot1	[[Training]]	[[none]]	(45, 48)	8	0	
	1	[[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0	1683.0	lot1	[[Training]]	[[none]]	(45, 48)	8	0	
	2	[[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0	1683.0	lot1	[[Training]]	[[none]]	(45, 48)	8	0	
	3	[[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0	1683.0	lot1	[[Training]]	[[none]]	(45, 48)	8	0	
	4	[[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0	1683.0	lot1	[[Training]]	[[none]]	(45, 48)	8	0	

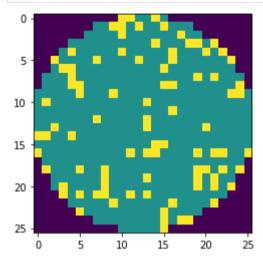
```
In []: # 구분별 개수 찍어 봄
print("total: ", tol_wafers)
print("none-label: ", tol_wafers -df_withlabel.shape[0])
print("label: ", df_withlabel.shape[0])
print("ng: ", df_withpattern.shape[0])
print("ok: ", df_nonpattern.shape[0])
```

```
In []: #wafer index 제외 df = df.drop(['waferIndex'], axis = 1)
```

```
비니다 #각 wafer map의 해상도 size가 다름 -- 해상도 size에 대한(행렬size) 컬럼과 데이터 생성
        def find_dim(x):
            dim0=np.size(x.axis=0)
            dim1=np.size(x.axis=1)
            return dim0.dim1
        df['waferMapDim']=df.waferMap.apply(find_dim)
         df.sample(5)
                                 waferMap dieSize lotName trianTestLabel failureType waferMapDim
Out[ ]:
        158981 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 1, 2, 1,... 515.0 lot10237
                                                             [[Training]]
                                                                       [[Center]]
                                                                                     (25, 27)
        487489 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 2,... 710.0 lot30167
                                                                   Π
                                                                                     (32, 29)
                                                                             П
        П
                                                                             П
                                                                                     (49, 39)
        502224 [[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, ... 710.0 lot31209
                                                                   П
                                                                                     (32, 29)
        П
                                                                             П
                                                                                     (64, 72)
In [ ]:
        #26*26 해상도의 wafer data 추출
        sub_df = df.loc[df['waferMapDim'] == (26, 26)]
        sub_wafer = sub_df['waferMap'].values
         sw = np.ones((1, 26, 26))
         label = list()
        for i in range(len(sub_df)):
            # 불량코드에 null 라벨의 skip 처리
            if len(sub_df.iloc[i,:]['failureType']) == 0:
                continue
            sw = np.concatenate((sw, sub_df.iloc[i,:]['waferMap'].reshape(1, 26, 26)))
            label.append(sub_df.iloc[i,:]['failureType'][0][0])
In [ ]:
        x = sw[1:]
        y = np.array(label).reshape((-1,1))
         # x, y shape
        print('x shape : {}, y shape : {}'.format(x.shape, y.shape))
```

```
In []: # plot 1st data 보여주고
plt.imshow(x[0])
plt.show()

# check faulty case: failure type
print('Faulty case : {} '.format(y[0]))
```



```
In []: #add channel
    x = x.reshape((-1, 26, 26, 1))

faulty_case = np.unique(y)
    print('Faulty case list : {}'.format(faulty_case))
```

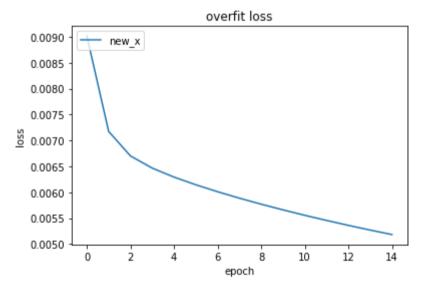
```
In []: #26*26 size map의 failure type별 데이터 개수 for f in faulty_case : print('{} : {}'.format(f, len(y[y==f])))
```

```
# One-hot-Encoding faulty categorical variable as channel
new_x = np.zeros((len(x), 26, 26, 3))
for w in range(len(x)):
```

```
for i in range(26):
                 for j in range(26):
                     new_x[w, i, j, int(x[w, i, j])] = 1
In [ ]:
         #check new x dimension
         new_x.shape
Out[ ]:
In [ ]:
          # parameter
         epoch=15
         batch_size=1024
         # Encoder
         input\_shape = (26, 26, 3)
         input_tensor = Input(input_shape)
         encode = layers.Conv2D(64, (3,3), padding='same', activation='relu')(input_tensor)
         latent_vector = layers.MaxPool2D()(encode)
         # Decoder
         decode_layer_1 = layers.Conv2DTranspose(64, (3,3), padding='same', activation='relu')
         decode_layer_2 = layers.UpSampling2D()
         output_tensor = layers.Conv2DTranspose(3, (3,3), padding='same', activation='sigmoid')
         # connect decoder layers
         decode = decode_layer_1(latent_vector)
         decode = decode_layer_2(decode)
         ae = models.Model(input_tensor, output_tensor(decode))
         ae.compile(optimizer = 'Adam',
                        loss = 'mse',
         ae.summary()
```

```
epochs=epoch,
verbose=2)
```

```
In []:
# loss chart - 데이터 불균형 그대로 .. 과적합 검증
# loss plot
plt.plot(ae_history.history['loss'])
plt.title('overfit loss')
plt.ylabel('loss')
plt.xlabel('epoch')
plt.legend(['new_x'], loc='upper left')
plt.show()
```



```
In []: # Make encoder model with part of autoencoder model layers
    encoder = models.Model(input_tensor, latent_vector)

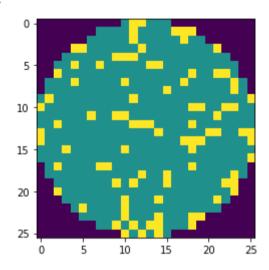
# Make decoder model with part of autoencoder model layers
    decoder_input = Input((13, 13, 64))
    decode = decode_layer_1(decoder_input)
    decode = decode_layer_2(decode)
    decoder = models.Model(decoder_input, output_tensor(decode))

# Encode original faulty wafer
    encoded x = encoder_predict(new_x)
```

```
# Add noise to encoded latent faulty wafers vector.
noised_encoded_x = encoded_x + np.random.normal(loc=0, scale=0.1, size = (len(encoded_x), 13, 13, 64))
```

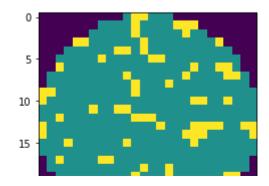
```
# check original faulty wafer data
plt.imshow(np.argmax(new_x[3], axis=2))
```

Out[]:



```
# check new noised faulty wafer data
noised_gen_x = np.argmax(decoder.predict(noised_encoded_x), axis=3)
plt.imshow(noised_gen_x[3])
```

Out[]:

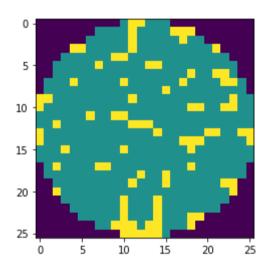


```
25 - 0 5 10 15 20 25
```

```
# check reconstructed original faulty wafer data
gen_x = np.argmax(ae.predict(new_x), axis=3)
plt.imshow(gen_x[3])

# ------ Data augmentation ------
```

Out[]:



```
In []: # augment function define
    def gen_data(wafer, label):
        # Encode input wafer
        encoded_x = encoder.predict(wafer)

# dummy array for collecting noised wafer
        gen_x = np.zeros((1, 26, 26, 3))

# Make wafer until total # of wafer to 2000
        for i in range((2000//len(wafer)) + 1):
            noised_encoded_x = encoded_x + np.random.normal(loc=0, scale=0.1, size = (len(encoded_x), 13, 13, 64))
            noised_encoded_x = decoder_predict(paised_encoded_x)
```

```
ποτοσα_χοπ_Λ ασσοάστιρισατοι (ποτοσα_σπόσασα_Λ/
                 gen_x = np.concatenate((gen_x, noised_gen_x), axis=0)
             # also make label vector with same length
             gen v = np.full((len(gen x), 1), label)
             # return date without 1st dummy data.
             return gen_x[1:], gen_y[1:]
In [ ]:
         # Augmentation for all faulty case.
         for f in faulty_case:
             # skip none case
             if f == 'none' :
                 continue
             gen_x, gen_y = gen_data(new_x[np.where(y==f)[0]], f)
             new_x = np.concatenate((new_x, gen_x), axis=0)
             y = np.concatenate((y, gen_y))
         print('After Generate new_x shape : {}, new_y shape : {}'.format(new_x.shape, y.shape))
         for f in faulty_case:
             print('\{\}: \{\}'.format(f, len(y[y==f])))
         # choice index without replace.
         none_idx = np.where(y=='none')[0][np.random.choice(len(np.where(y=='none')[0]), size=11000, replace=False)]
In [ ]:
         # delete choiced index data.
         new_x = np.delete(new_x, none_idx, axis=0)
         new_y = np.delete(y, none_idx, axis=0)
         print('After Delete "none" class new_x shape : {}, new_y shape : {}'.format(new_x.shape, new_y.shape))
```

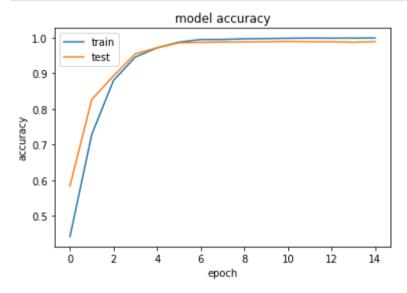
```
print('\{\}: \{\}', format(f, len(new v[new v==f])))
In [ ]:
         # make string label data to numerical data
         for i, I in enumerate(faulty_case):
             new_y[new_y==1] = i
         # one-hot-encoding
         new_v = to_categorical(new_v)
         # split data train, test
         x_train, x_test, y_train, y_test = train_test_split(new_x, new_y,
                                                              test size=0.33.
                                                              random_state=2019)
         print('Train x : {}, y : {}'.format(x_train.shape, y_train.shape))
         print('Test x: {}, y : {}'.format(x_test.shape, y_test.shape))
In [ ]:
         # The data is ready. As wafer data is image, simply use cnn for classification.
In [ ]:
         # The data is ready. As wafer data is image. simply use cnn for classification.
         # Make model
         # define create model function, because we will validate model with sklearn kfold cross validation.
         def create_model():
             input\_shape = (26, 26, 3)
             input_tensor = Input(input_shape)
             conv_1 = layers.Conv2D(16, (3,3), activation='relu', padding='same')(input_tensor)
             conv_2 = layers.Conv2D(64, (3,3), activation='relu', padding='same')(conv_1)
             conv_3 = layers.Conv2D(128, (3,3), activation='relu', padding='same')(conv_2)
             flat = layers.Flatten()(conv_3)
             dense 1 = layers. Dense(512, activation='relu')(flat)
             dense_2 = layers.Dense(128, activation='relu')(dense_1)
             output tensor = layers.Dense(9, activation='softmax')(dense 2)
             model = models. Model(input tensor, output tensor)
              model compile on timizer = 1 Adom!
```

TOT I III TAUTLY_Case .

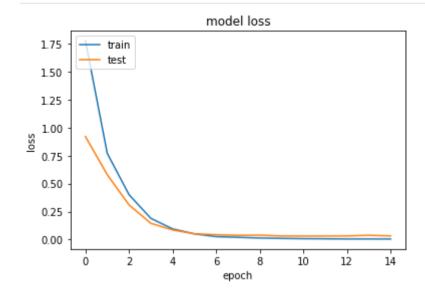
```
moder.comprie(optimizer- Adam ,
                          loss='categorical_crossentropy',
                          metrics=['accuracy'])
             return model
In [ ]:
         # Cross validate model: Using sklearn KFold Cross validation, we validate our simple cnn.
         # Make keras model to sklearn classifier.
         model = KerasClassifier(build_fn=create_model, epochs=10, batch_size=1024, verbose=2)
         # 3-Fold Crossvalidation
         kfold = KFold(n_splits=3, shuffle=True, random_state=2019)
         results = cross_val_score(model, x_train, y_train, cv=kfold)
         # Check 3-fold model's mean accuracy
         print('Simple CNN Cross validation score : {:.4f}'.format(np.mean(results)))
         batch size
Out[ ]:
         print(x_test.shape)
         print(y_test.shape)
         print(x_train.shape)
         print(y_train.shape)
         y_train.dtype
Out[]:
         # Our model seems quite a good model.
         history = model.fit(x_train, y_train,
                  epochs=epoch,
```

```
batch_size=batch_size,
validation_data=(x_test, y_test)
)
```

```
# accuracy chart
# accuracy plot
plt.plot(history.history['accuracy'])
plt.plot(history.history['val_accuracy'])
plt.title('model accuracy')
plt.ylabel('accuracy')
plt.xlabel('epoch')
plt.legend(['train', 'test'], loc='upper left')
plt.show()
```



```
In []: # loss plot
plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])
plt.title('model loss')
plt.ylabel('loss')
plt.xlabel('epoch')
plt.legend(['train', 'test'], loc='upper left')
plt.show()
```



In []: