# Cassava Leaf Disease Classification

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## **Overview**



## Introduction

**Cassava** is a key food security crop in Africa
But **viral diseases** are major sources of poor yields.
Existing methods of disease detection require experts to visually inspect and diagnose the plants -> labor-intensive, low-supply and costly.

So I want to solve this problem through data science.

-> classify each cassava image into 5 category

## 1. Data Exploration and Visualization

```
BASE_DIR = os.getcwd()
with open(os.path.join(BASE_DIR, "label_num_to_disease_map.json")) as file:
    map_classes = json.loads(file.read())
    map_classes = {int(k) : v for k, v in map_classes.items()}

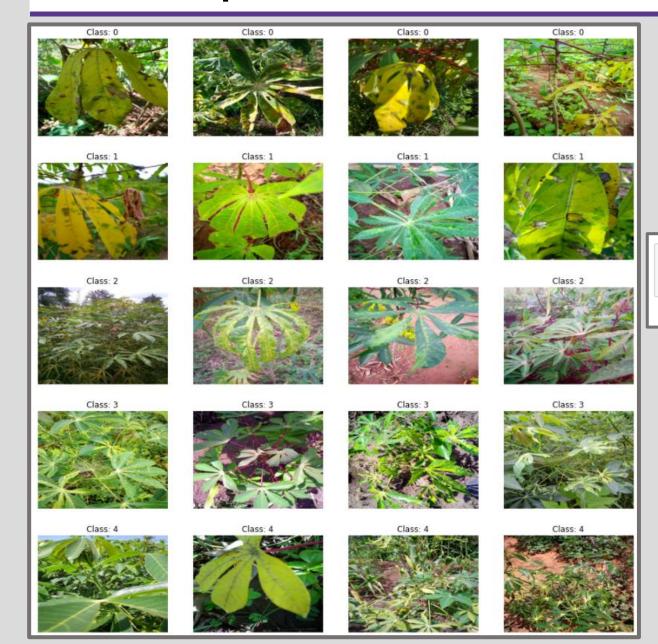
print(json.dumps(map_classes, indent=4))

{
    "O": "Cassava Bacterial Blight (CBB)",
    "1": "Cassava Brown Streak Disease (CBSD)",
    "2": "Cassava Green Mottle (CGM)",
    "3": "Cassava Mosaic Disease (CMD)",
    "4": "Healthy"
}
```

```
img_shapes = {}
for image_name in os.listdir(os.path.join(BASE_DIR, "train_images"))[:300]:
    image = cv2.imread(os.path.join(BASE_DIR, "train_images", image_name))
    img_shapes[image.shape] = img_shapes.get(image.shape, 0) + 1

print(img_shapes)
{(600, 800, 3): 300}
```

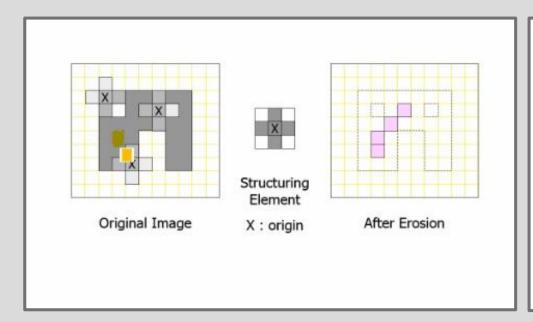
# 1. Data Exploration and Visualization

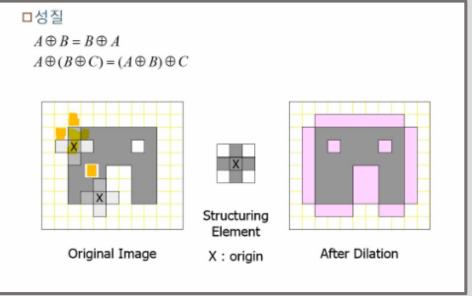


input\_files = os.listdir(os.path.join(BASE\_DIR, "train\_images"))
print(f"Number of train images: {len(input\_files)}")

Number of train images: 21397

## **Segmentation with Opency**





#### **Erosion**

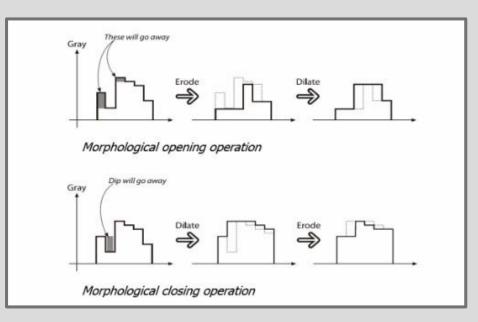
: 각 Pixel에 structuring element를 적용하여 안겹치는 부분이 하나라도 있으면 그 중심 pixel을 제거하는 방법

#### **Dilation**

: 각 pixel에 structuring element를 적용하여 겹치는 부분이 하나라도 있으면 이미지를 확장

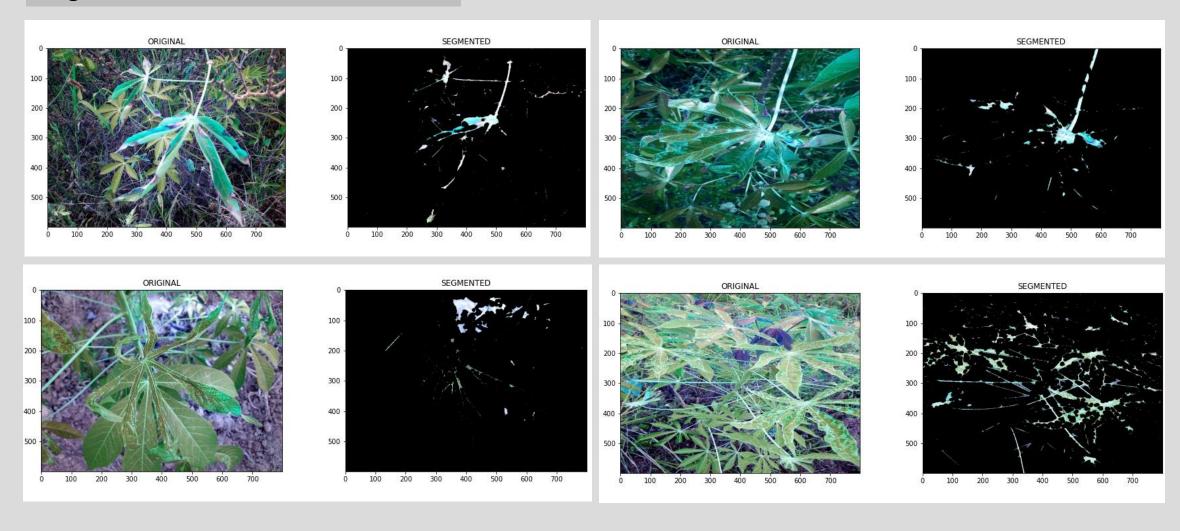
#### **Segmentation with Opency**

```
def read_image(image_id , label):
    plt.figure(figsize=(15, 10))
    image = cv2.imread(os.path.join(BASE_DIR, "train_images", image_id))
    return image
def create_masks(image):
    image_hsv = cv2.cvtColor(image, cv2.COLOR_BGR2HSV)
    lower_hsv = np.array([0,0,250])
    upper_hsv = np.array([250,255,255])
    mask = cv2 inRange(image hsv | lower hsv | unner hsv)
    kernel = cv2.getStructuringElement(cv2.MORPH_ELLIPSE, (11,11))
    mask = cv2.morphologyEx(mask, cv2.MORPH_CLOSE, kernel)
    return mask
def segment_image(image):
    mask = create_masks(image)
    output = cv2.bitwise_and(image, image, mask=mask)
    return output/255
```



**Opening & Closing** 

## Segmentation -> X



## **Image Augmentation + GrayScale**









# 3. Training – Vanilla CNN

max_pooling2d_2 (MaxPooling2	(None, 16, 16, 64)	0
conv2d_3 (Conv2D)	(None, 16, 16, 32)	18464
max_pooling2d_3 (MaxPooling2	(None, 8, 8, 32)	0
dropout_2 (Dropout)	(None, 8, 8, 32)	0
flatten_1 (Flatten)	(None, 2048)	0
dense_1 (Dense)	(None, 256)	524544
dropout_3 (Dropout)	(None, 256)	0
dense_2 (Dense)	(None, 5)	1285
Total params: 546,085		

Output Shape

(None, 32, 32, 64)

Param #

1792

Layer (type)

conv2d\_2 (Conv2D)

Trainable params: 546,085 Non-trainable params: 0

**Test accuracy** 

less than 60%

# 3. Training - EfficientNet

The traditional way to improve CNN performance : increase the depth of model

## -> EfficientNet

: consider not only **depth** but also **width** and **resolution** together

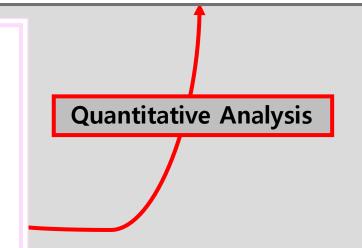
Stage	Operator	Resolution	#Channels	#Layers
i	$\hat{\mathcal{F}}_{\pmb{i}}$	$\hat{H}_i \times \hat{W}_i$	$\hat{C}_{m{i}}$	$\hat{L}_i$
1	Conv3x3	$224 \times 224$	32	1
2	MBConv1, k3x3	$112 \times 112$	16	1
3	MBConv6, k3x3	$112 \times 112$	24	2
4	MBConv6, k5x5	$56 \times 56$	40	2
5	MBConv6, k3x3	$28 \times 28$	80	3
6	MBConv6, k5x5	$14 \times 14$	112	3
7	MBConv6, k5x5	$14 \times 14$	192	4
8	MBConv6, k3x3	7  imes 7	320	1
9	Conv1x1 & Pooling & FC	$7 \times 7$	1280	1

Information:  $128 \times 128 > 64 \times 64$ .

But high-resolution input can result in a large calculation cost,

and does not provide a significant benefit to model performance after certain values.

-> **Balancing** the width, depth, and resolution of a network is very important for improving performance.



# 3. Training - EfficientNet

```
early_stopping = EarlyStopping(monitor='val_accuracy', patience=5)
mc = ModelCheckpoint('best_model.h5', monitor='val_loss', mode='min', save_best_only=True)
model = Sequential()
optimizer = Adam(Ir=0.00105)
b4model = EfficientNetB4(include_top = False,
                          weights = None.
                          pooling='avg')
model.add(b4model)
                                              model.add(Dense(256, activation = 'relu'))
model.add(Dense(5, activation ='softmax'))
model.summary()
Model: "sequential 6"
Layer (type)
                             Output Shape
                                                       Param #
efficientnetb4 (Functional) (None, 1792)
                                                       17673823
dense_11 (Dense)
                             (None, 5)
                                                       8965
Total params: 17,682,788
Trainable params: 17,557,581
Non-trainable params: 125,207
```

# 3. Training - EfficientNet

```
model.compile(loss="categorical_crossentropy", optimizer=optimizer, metrics=["accuracy"])

%%time
hist10 = model.fit(
    train_generator,
    steps_per_epoch = steps_per_epoch,
    epochs = epochs,
    validation_data = validation_generator,
    validation_steps = validation_steps,
    callbacks=[early_stopping, mc],
    verbose = 1)
```

## **Test accuracy**

79~81%

Training on a single epoch is extremely slow... (3 hours)

-> **GPU?** 

# **Kaggle Competition**

[1] Cassava Leaf Disease Classification,

https://www.kaggle.com/c/cassava-leaf-disease-classification/overview

## References

[1] The shortest way to Tensorflow baseline,

https://www.kaggle.com/nozarchos/the-shortest-way-to-tensorflow-baseline

[2] Tensorflow ViT and Image Pre-processing,

https://www.kaggle.com/digvijayyadav/tensorflow-vit-and-image-pre-processing

[3] Convolutional Neural Network의 성능을 높이는 현명한 방법: EfficientNet Google AI, https://ichi.pro/ko/convolutional-neural-networkui-seongneung-eul-nop-ineun-hyeonmyeonghan-bangbeob-efficientnet-google-ai-220426838454404