Papaya Disease Detector using python tensorflow & SVM

My introduction

 B.S. majoring 'Unmanned Vehicle Engineering' of 'Al robot' department in Sejong univ.

- Experience
 - Mini autonomous car (Xy-car) driving competition 6th place overall, 2021
 - Matlab AI challenge competition Bronze award, 2023
 - Studying in lab, IVPG, 2023~

• Creation of AI fashion recommendation webpage (in charge of backend and CV), 2023

Contents

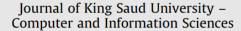
- 1. Introduction
- 2. Explanation
- 3. Flow Chart
- 4. Screenshots of Code
- 5. Result (Demonstration)

Introduction

Journal of King Saud University - Computer and Information Sciences 32 (2020) 300-309

Contents lists available at ScienceDirect





journal homepage: www.sciencedirect.com



Machine vision based papaya disease recognition





^b Department of Computer Science and Engineering, Daffodil International University, Dhaka, Bangladesh

ARTICLE INFO

Article history: Received 22 March 2018 Revised 7 June 2018 Accepted 13 June 2018 Available online 18 June 2018

Keywords:
Papaya disease
Agro-medical expert system
Machine vision
k-means clustering
Support vector machine

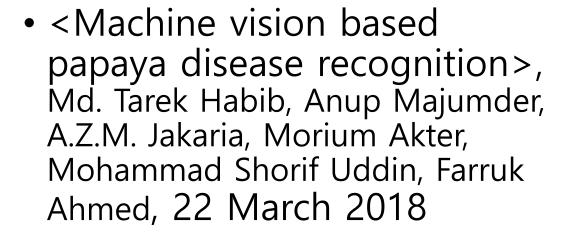
ABSTRACT

Over the years little research has been performed for vision-based papaya disease recognition system in order to help distant farmers, most of whom require proper support for cultivation. Due to advancement of vision-based technology we find a good solution to this problem. Papaya disease recognition mainly involves two challenging problems: one is disease detection and another is disease classification. Considering this scenario, here we present an online machine vision-based agro-medical expert system that processes an image captured through mobile or handheld device and determines the diseases in order to help distant farmers to address the problem. Some experiments are performed to show the dility of the proposed expert system. This, we propose a set of features from the view point of distinguishing attributes. K-means clustering algorithm is used in order to segment out the disease-attacked region from the captured image and then required features are extracted to classify the diseases with the help of support vector machine. More than 90% classification accuracy has been achieved, which appears to be good as well as promising by comparing performances obtained with recently reported relevant weeks.

© 2018 The Authors. Production and hosting by Elsevier B.V. on behalf of King Saud University. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).

1. Introduction

Bangladesh is a developing as well as densely populated country, where the prominent portion of the population depends on agriculture. It is agriculture that has managed employment for quality of the agricultural product must be maintained with proper care. In Bangladesh, the rate of literacy is only 64.6% (List of countries by literacy rate, 2018) and most of the farmers are illiterate. They have hardly any training on farming with advanced technology. Again, there are few problems that come into play. One of



 This paper is written to help farmers in need of support in agriculture, and deals with machine vision technology for papaya disease recognition.

Department of Computer Science and Engineering, Independent University, Bangladesh, Dhaka, Bangladesh

Introduction

- According to this paper, some countries are especially dependent on agriculture.
- In Bangladesh, for an example, 47% of the total population of the country is employed in agriculture.
- However, 39.9% of post-harvest losses were occurred in papaya agriculture, which is immensely popular product in Bangladesh. Moreover, there were two problems in dealing with this loss problems.
- 1) <u>Transportation problem</u>: they cannot reach the agriculture centers easily.
- 2) ignorance of the farmers towards the <u>advanced technology</u>
- So, If they are provided with **proper support** for producing better and disease-free crops, then it will be highly beneficial for them.

Introduction

First, the size of the picture is changed to a fixed size using 'Bicubic interpolation'. And then, the pictures are applied histogram equalization to improve contrast. Then, 'K-means clustering' technology is used to distinguish between defective and non-defective parts. There are the five most frequent papaya diseases, which are called black spot, powdery mildew, brown spot, phytophthora light, and anthracnose.

And then, we extract features from the 'disease part' which is distinguished via clustering step. There are two main features for distinguishing papaya diseases: one is a statistical feature, and the other is a co-occurrence feature. "Statistical features" have mean, standard deviation, variance, kurtosis, and skewness. "co-occurrence features" have contrast, correlation, energy, entropy, and homogeneity. (Each feature has a formula that can be calculated which is referred from previous published papers. Especially, these features were selected by referring to what was concluded to be meaningful in other papers.) After clustering, the vector of these features is extracted from the defective part.

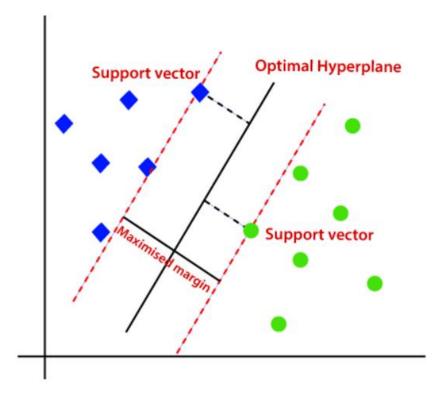
Then, finally, you can solve the multi-class problem by using the feature vector in the SVM, which is one of the way to teach a machine about features.

Performance evaluation was conducted using images that were manually labeled with defects (ground truth images). As a result, the accuracy reached 95.2%. K-means clustering and SVM technologies were selected due to the excellent performance found in other existing papers, and in the case of SVM, the performance of other classification technologies, decision tree and naive bayes, was compared for comparison, but the SVM was finally selected with overwhelming accuracy.

- As this paper figure out the best
 way to make papaya disease
 classifier, I decided to make the
 papaya disease classifier for myself.
- Papaya disease recognition has two challenges:
- 1. Detecting papaya disease
- 2. Classifying the disease

Explanation

- 1. SVM (support vector machine)
- It is a supervised learning model that classifies and regresses based on a decision boundary, that is, a line similar to a linear model.



Explanation

2. Image_dataset_from_directory / Dataset

```
tf.keras.utils.image_dataset_from_directory(
     directory,
     labels='inferred'.
                                                                         Either "inferred" (labels are generated from the directory structure), None (no labels), or a
                                             labels
     label_mode='int',
                                                                         list/tuple of integer labels of the same size as the number of image files found in the directory.
                                                                         Labels should be sorted according to the alphanumeric order of the image file paths (obtained via
     class_names=None,
                                                                         os.walk(directory) in Python).
     color_mode='rgb',
     batch_size=32.
                                             validation_split
                                                                         Optional float between 0 and 1, fraction of data to reserve for validation.
     image_size=(256, 256),
                                             subset
                                                                         Subset of the data to return. One of "training", "validation", or "both". Only used if
     shuffle=True.
                                                                         validation_split is set. When subset="both", the utility returns a tuple of two datasets (the
     seed=None.
                                                                         training and validation datasets respectively).
     validation_split=None,
     subset=None,
     interpolation='bilinear',
                                                                       Returns
     follow_links=False,
     crop_to_aspect_ratio=False,
     **kwargs
                                                                     A tf.data.Dataset object.
```

Explanation

2. Image_dataset_from_directory / Dataset

```
tf.data.Dataset(
    variant_tensor
)
```



Consist of tensor

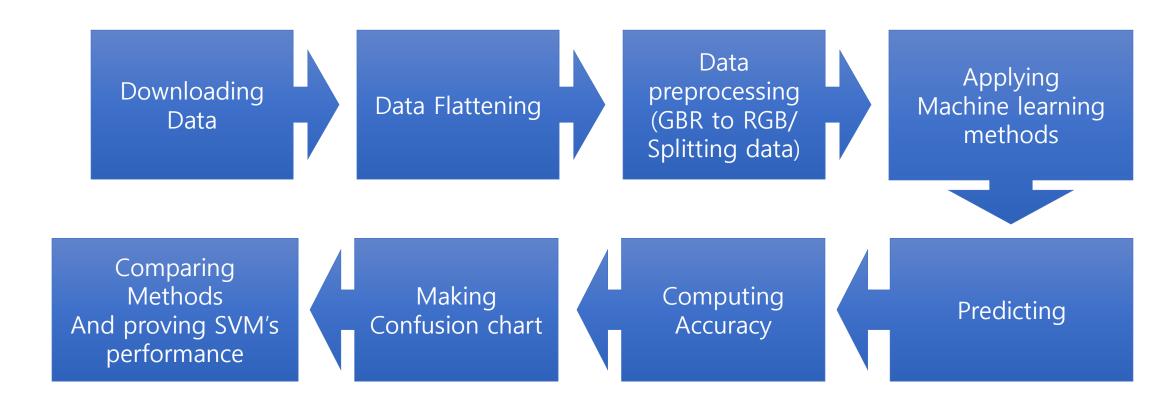
```
take(
   count, name=None
)
```

Creates a Dataset with at most count elements from this dataset.

```
>>> dataset = tf.data.Dataset.range(10)
>>> dataset = dataset.take(3)
>>> list(dataset.as_numpy_iterator())
[0, 1, 2]
```

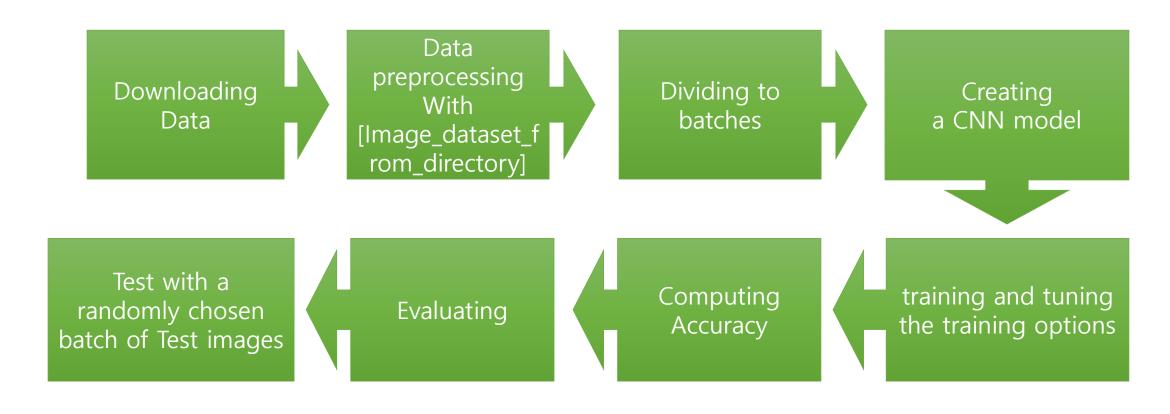
Flow Chart

Part 1) SVM (Machine Learning Part)



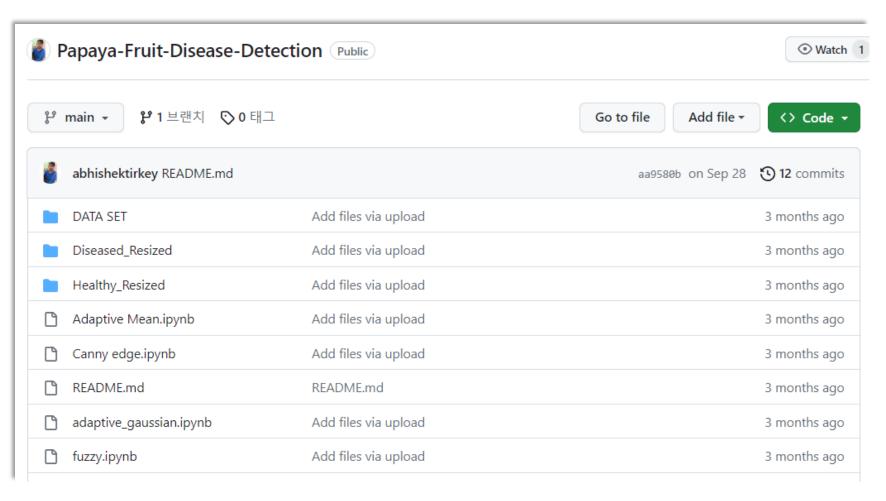
Flow Chart

Part 2) CNN (Deep Learning Part)



Screenshots of Code

1. Downloading Data



A papaya data set in github

266

Name: Target, dtvpe: int64

```
class_names=['Diseased','Healthy']
flat_data_arr=[] #input array
target arr= 1 #output array
data_dir='/content/drive/MyDrive/Colab Notebooks/machine vision system project'
#path which contains all the class names of images
for i in class_names:
    print(f'loading... category : {i}')
    path=os.path.ioin(data dir.i)
    print(path)
    for ima in os.listdir(path):
        img_array=cv2.imread(os.path.ioin(path.img))
        flat_data_arr.append(img_array.flatten())
        target arr.append(class names.index(i))
    print(f'loaded category:{i} successfully')
flat_data=np.array(flat_data_arr)
target=np.arrav(target arr)
df=pd.DataFrame(flat data) #dataframe
df['Target']=target
x=df.iloc[:,:-1] #input data
v=df.iloc[:.-1] #output data
print("x:",x.shape)
print("y:",y.shape)
x train.x test.v train.v test=train test split(x.v.test size=.20.random state=77.stratifv=v)
print('Splitted Successfully')
```

- 1) Import Data
- 2) Data Flattening
- 3) Data split (train/test)

```
loading... category: Diseased
/content/drive/MyDrive/Colab Notebooks/machine vision system project/Diseased
loaded category:Diseased successfully
loading... category: Healthy
/content/drive/MyDrive/Colab Notebooks/machine vision system project/Healthy
loaded category:Healthy successfully
x: (421, 30000)
y: (421,)
Splitted Successfully
```

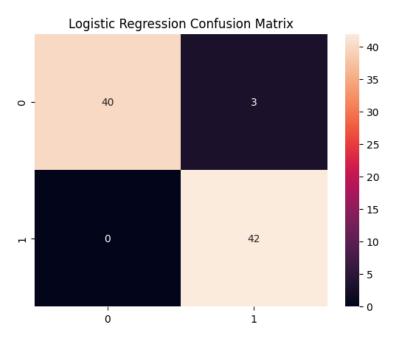
	0	1	2	3	4	5	6	7	8	9	 29990	29991	29992	29993	29994	29995	29996	29997	29998	29999
223	248	222	206	249	223	207	253	221	208	253	 139	177	148	139	177	148	139	177	148	139
266	253	253	253	253	253	253	253	253	253	253	 253	253	253	253	253	253	253	253	253	253
352	183	160	152	183	160	152	183	160	152	183	 192	233	197	191	234	198	192	237	201	195
325	172	138	132	172	138	132	172	138	132	172	 176	224	185	176	223	184	175	223	184	175
386	188	158	153	188	158	153	190	158	153	190	 187	239	196	187	239	196	187	239	196	187
rows	× 30	000 с	olumn	ns																
y_tra	in .hea	ad()																		

1) Logistic Regression

```
Ir = LogisticRegression(penalty='I1', solver = 'liblinear', random_state=42)
Ir.fit(x_train, y_train)

y_pred = Ir.predict(x_test)
Ir_train_acc = round(accuracy_score(y_train, Ir.predict(x_train)) *100, 2) # 소숫점 2자리로 반올림
Ir_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
print('\maccuracy = ', Ir_test_acc, ' \mathcal{w}\mathrix')

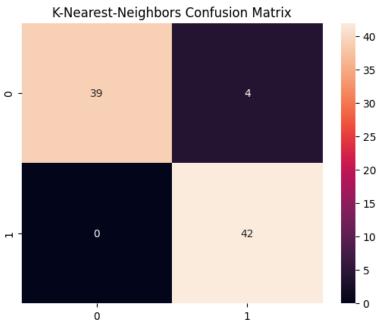
cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True)
plt.title('Logistic Regression Confusion Matrix')
```



Accuracy = 96.47 %

2) KNN

```
| knn = KNeighborsClassifier(n_neighbors=5)
| knn.fit(x_train, y_train)
| y_pred = knn.predict(x_test)
| knn_train_acc = round(accuracy_score(y_train, |r.predict(x_train)) *100, 2) # 소숫점 2자리로 반올림
| knn_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
| print('\mathref{w}nAccuracy = ', knn_test_acc, ' \mathref{w}m')
| cm = confusion_matrix(y_test, y_pred)
| sns.heatmap(cm, annot=True)
| plt.title('K-Nearest-Neighbors Confusion Matrix')
```



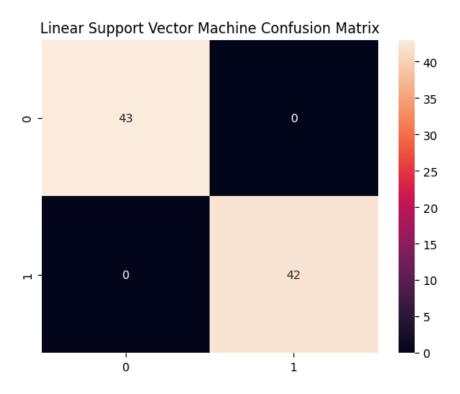
Accuracy = 95.29 %

3) SVM

Accuracy = 100.0 %

```
In_svm = SVC(kernel='linear')
In_svm.fit(x_train, y_train)
y_pred = In_svm.predict(x_test)
In_svm_train_acc = round(accuracy_score(y_train, In_svm.predict(x_train)) *100, 2)
In_svm_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
print('\mathraceuracy = ', In_svm_test_acc, ' \mathraceuracy')

cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True)
plt.title('Linear Support Vector Machine Confusion Matrix')
```



4) Decision Tree

```
tree = DecisionTreeClassifier(max_depth=5)
tree.fit(x_train, y_train)

y_pred = tree.predict(x_test)
tree_train_acc = round(accuracy_score(y_train, tree.predict(x_train)) *100, 2)
tree_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
print('\max_depth=2 confusion_matrix(y_test_acc, ' \max_mn')

cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True)
plt.title('Decision Tree Confusion Matrix')
```

```
Decision Tree Confusion Matrix

- 40
- 35
- 30
- 25
- 20
- 15
- 10
- 5
```

Accuracy = 96.47 %

Random Forest Confusion Matrix

5) Random Forest

Accuracy = 96.47 %

```
rdm_frst = RandomForestClassifier(n_estimators=5, random_state=10)
rdm_frst.fit(x_train, y_train)

y_pred = rdm_frst.predict(x_test)
rdm_train_acc = round(accuracy_score(y_train, rdm_frst.predict(x_train)) *100, 2)
rdm_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
print('\text{\text{\text{MAccuracy}} = ', rdm_test_acc, ' \text{\text{\text{\text{\text{M}n'}}}})

cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True)
plt.title('Random Forest Confusion Matrix')
```

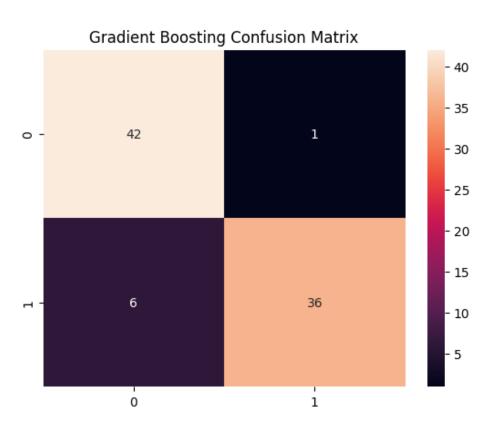
6) Gradient Boosting

```
gb = GradientBoostingClassifier(n_estimators=5, random_state=10)
gb.fit(x_train, y_train)

y_pred = gb.predict(x_test)
gb_train_acc = round(accuracy_score(y_train, gb.predict(x_train)) *100, 2)
gb_test_acc = round(accuracy_score(y_test, y_pred)*100, 2)
print('\maccuracy = ', gb_test_acc, ' \mathcal{m}n')

cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True)
plt.title('Gradient Boosting Confusion Matrix')
```

Accuracy = 91.76 %



Comparing Classification Models

```
models = pd.DataFrame({
    'Model': [
        'Logistic Regression', 'K Nearest Neighbors', 'Linear Support Vector Machines',
        'Decision Tree', 'Random Forest', 'Gradient Boosting'
],
    'Model Accuracy Score': [
        Ir_test_acc,knn_test_acc,In_svm_test_acc,tree_test_acc,rdm_test_acc,gb_test_acc
]
})
models.sort_values(by='Model Accuracy Score', ascending=False)
```

	Mode I	Model Accuracy Score
2	Linear Support Vector Machines	100.00
0	Logistic Regression	96.47
3	Decision Tree	96.47
4	Random Forest	96.47
1	K Nearest Neighbors	95.29
5	Gradient Boosting	91.76

Then, finally, you can solve the multi-class problem by using the feature vector in the SVM, which is one of the way to teach a machine about features.

Performance evaluation was conducted using images that were manually labeled with defects (ground truth images). As a result, the accuracy reached 95.2%. K-means clustering and SVM technologies were selected due to the excellent performance found in other existing papers, and in the case of SVM, the performance of other classification technologies, decision tree and naive bayes, was compared for comparison, but the SVM was finally selected with overwhelming accuracy.

As stated in the paper, SVM shows the highest performance.

With SVM

```
img=cv2.imread("/content/drive/MyDrive/Colab Notebooks/machine vision system project/Diseased/a0_a49.jpg")
img = cv2.cvtColor(img, cv2.CoLOR_BGR2RGB) #brg->rgb
plt.imshow(img)
plt.axis('off')
plt.show()

sample=[img.flatten()]
probability_svm=ln_svm.predict(sample)
print("The predicted image is : "+class_names[probability_svm[0]])
```



The predicted image is: Diseased

With SVM

```
img=cv2.imread("/content/drive/MyDrive/Colab Notebooks/machine vision system project/Healthy/r3_1 (25).jpg")
img = cv2.cvtColor(img, cv2.CoLOR_BGR2RGB) #brg->rgb
plt.imshow(img)
plt.axis('off')
plt.show()

sample=[img.flatten()]
probability_svm=In_svm.predict(sample)
print("The predicted image is : "+class_names[probability_svm[0]])
```



The predicted image is : Healthy

Importing Data

```
PATH = '_/content/drive/MyDrive/Colab Notebooks/machine vision system project'

#diseased_dir = os.path.join(PATH, 'Diseased')
#healthy_dir = os.path.join(PATH, 'Healthy')

data_dir = PATH

batch_size = 3
sz=(100,100)
```

데이터 split

```
train dataset = tf.keras.utils.image dataset from directory(data dir.
                                                             shuffle=True.
                                                             batch_size=batch_size,
                                                             validation_split=0.3,
                                                             subset="training",
                                                             interpolation='bicubic',
                                                             image_size=sz,
                                                             seed=1204)
test and val dataset = tf.keras.utils.image dataset from directory(data dir,
                                                             shuffle=True.
                                                             batch_size=batch_size,
                                                             validation_split=0.3,
                                                             subset="validation".
                                                             image_size=sz,
                                                             interpolation='bicubic',
                                                                    seed=1204)
```

Found 421 files belonging to 2 classes. Using 295 files for training. Found 421 files belonging to 2 classes. Using 126 files for validation.

split validation set

```
test batches num = tf.data.experimental.cardinality(test and val dataset)
 val_dataset = test_and_val_dataset.take(test_batches_num // 5)
 test_dataset = test_and_val_dataset.skip(test_batches_num // 5)
 print('Number of available test and validation batches: %d' %test batches num)
Number of available test and validation batches: 42
print('Number of validation batches: %d' % tf.data.experimental.cardinality(val_dataset))
print('Number of test batches: %d' % tf.data.experimental.cardinality(test dataset))
Number of validation batches: 8
Number of test batches: 34
print(train dataset.class names)
print(test and val dataset.class names)
class_names = train_dataset.class_names
['Diseased', 'Healthy']
['Diseased', 'Healthy']
```

Showing dataset

```
plt.figure(figsize=(10, 10))
for images, labels in train_dataset.take(1):
    for i in range(3):
        ax = plt.subplot(3, 3, i + 1)
        #plt.imshow(images[i].numpy().astype("uint8"))
        plt.imshow(images[i]/255)
        plt.title(class_names[labels[i]])
        plt.axis("off")
```

Healthy





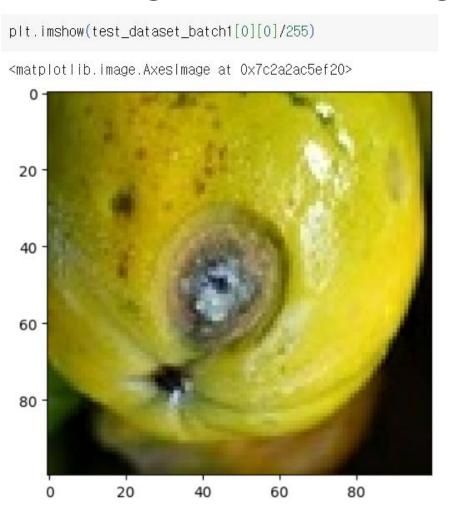
Dividing to batches

```
test_dataset_batch1 = list(test_dataset.take(1))[0]
test_dataset_batch1
```

```
[144., 148., 177.],
[143., 147., 176.],
...,
[152., 156., 191.],
[152., 156., 191.]],
[152., 156., 191.]],

[[123., 127., 156.],
[123., 127., 156.],
[123., 127., 156.],
[151., 155., 190.],
[151., 155., 190.],
[151., 155., 190.]]]], dtype=float32)>,
<tf.Tensor: shape=(3,), dtype=int32, numpy=array([1, 0, 1], dtype=int32)>)
```

• Showing one of the image in a Batch



Creating a CNN model

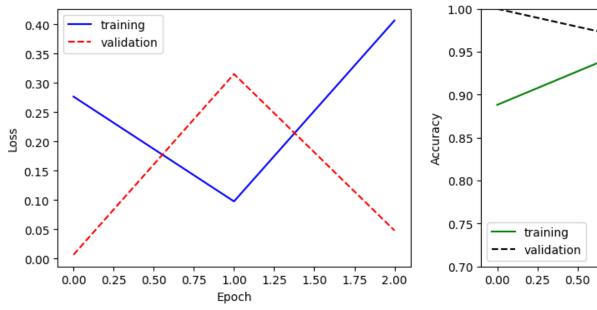
```
num_classes = 2
model = tf.keras.Sequential([
    tf.keras.layers.Rescaling(1./255), # lightness to range in [0,1]
    tf.keras.layers.Conv2D(32, 3, activation='relu',padding='same',input_shape=sz),
    tf.keras.layers.MaxPooling2D(strides=(2,2)),
    tf.keras.layers.Conv2D(32, 3, activation='relu',padding='same'),
    tf.keras.layers.MaxPooling2D(strides=(2,2)),
    tf.keras.layers.Conv2D(32, 3, activation='relu',padding='same'),
    tf.keras.layers.MaxPooling2D(strides=(2,2)).
    tf.keras.layers.Flatten(),
    tf.keras.layers.Dense(64, activation='relu'),
    tf.keras.layers.Dense(num_classes),
    tf.keras.layers.Softmax()
```

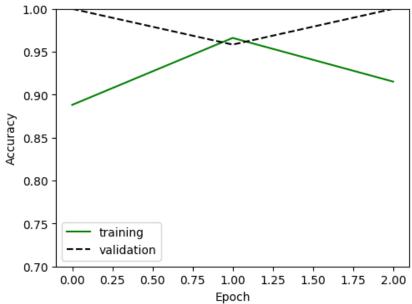
Compiling and fitting

```
model.compile(
 optimizer='adam'.
 loss=tf.keras.losses.SparseCategoricalCrossentropy(from logits=False), # loss=tf.keras.losses.BinaryCrossentropy(from logits=True)
 metrics=['accuracy'])
history = model.fit(
  train dataset.
  validation_data=val_dataset,
  epochs=3
Epoch 1/3
                       - 14s 99ms/step - loss: 0.2768 - accuracy: 0.8881 - val_loss: 0.0067 - val_accuracy: 1.0000
Epoch 2/3
Epoch 3/3
```

```
plt.figure(figsize=(12, 4)), plt.subplot(1, 2, 1)
plt.plot(history.history['loss'], 'b-', label='training')
plt.plot(history.history['val_loss'], 'r--', label='validation')
plt.xlabel('Epoch'), plt.ylabel('Loss'), plt.legend()
plt.subplot(1, 2, 2)
plt.plot(history.history['accuracy'], 'g-', label='training')
plt.plot(history.history['val_accuracy'], 'k--', label='validation')
plt.xlabel('Epoch'), plt.ylabel('Accuracy'), plt.ylim(0.7, 1), plt.legend()
plt.show()
```

Checking progress with plots





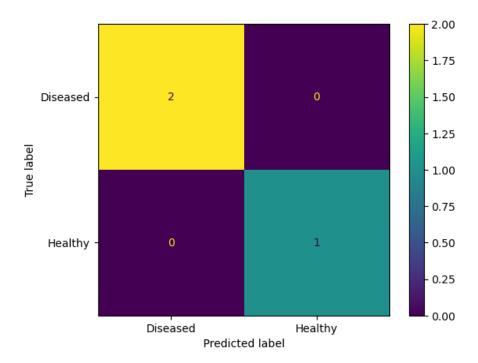
Evaluating

Test with a randomly chosen batch of Test images

```
test_dataset_batch1 = list(test_dataset.take(1))[0]
                                                                  Label:Healthy, Prediction:Healthy
                                                                                                      Label:Diseased, Prediction:Diseased
                                                                                                                                          Label:Diseased, Prediction:Diseased
#image batch, label batch = next(iter(test dataset))
prediction = model.predict(test_dataset_batch1[0])
                                                          20 -
predicted_labels = np.argmax(prediction, axis=1)
#test labels = np.argmax(test dataset, axis=1)
                                                          40
test_labels = test_dataset_batch1[1]
                                                          60
count = 0
plt.figure(figsize=(12,8))
for n in range(3):
     count += 1
     plt.subplot(1, 3, count)
     plt.imshow(test_dataset_batch1[0][n]/255)
                                                                                                                   prediction
     #plt.imshow(test dataset batch1[0][n].reshape(28, 28, 3)/255, interpolation='bicubic')
     tmp = "Label:" + class_names[test_labels[n]] + ", Prediction:" + class_names[predicted_labels[n]]
                                                                                                                   array([[0.03853271, 0.9614673],
     plt.title(tmp.fontdict = {'fontsize' : 10})
                                                                                                                          [0.04772344, 0.9522765 ],
                                                                                                                          [0.02524399, 0.974756 ]], dtype=float32)
plt.tight layout()
plt.show()
                                                                                                                                    ▲ score
```

Making confusion chart

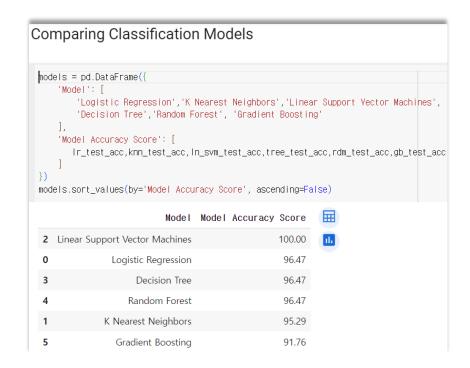
```
cm = confusion_matrix(test_labels.numpy(), predicted_labels)
disp = ConfusionMatrixDisplay(confusion_matrix=cm, display_labels=class_names)
disp.plot()
plt.show()
```

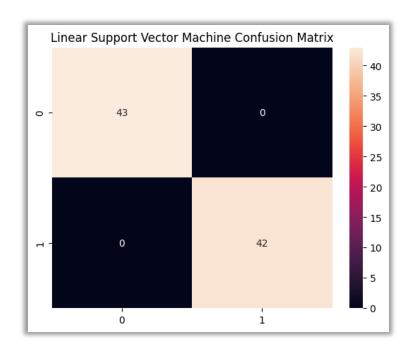


Screenshots of Code

4. Evaluating

- As this paper,
- "K-means clustering and SVM technologies were selected due to the excellent performance found in other existing papers." and "SVM was finally selected with overwhelming accuracy."
- It was confirmed that the contents of the paper were true with this try.



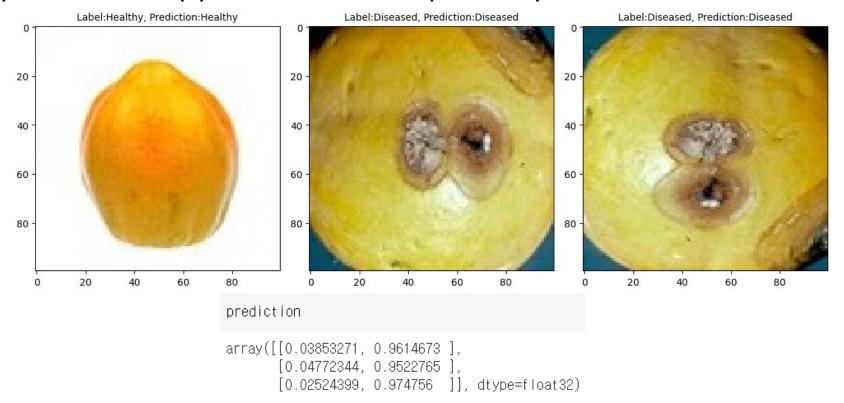


Screenshots of Code

4. Evaluating

 But Because machine learning involves the inconvenience of having to manually extract features,

I think using deep learning, which has sufficiently good performance, will be more helpful in the application development process.



Result (Demonstration)

https://colab.research.google.com/drive/1I8VxTbtETkzZR 2y7DphHx8UhnLWFpLu5?usp=sharing

https://colab.research.google.com/drive/1dybkeLY0U1dD GTd17wsH_58cllUJBR3w?usp=sharing