Development of a Deep Learning Model for Automated Detection and Diagnosis of Plant Diseases

```
In [ ]:
    import pandas as pd
    import pathlib
 2
    import tensorflow as tf
 4 import matplotlib.pyplot as plt
    import numpy as np
                                                                                 Classified as
 6 import os
 7
    import PIL
 8 import glob
 9 | from tensorflow import keras
10 from tensorflow.keras import layers
11 from tensorflow.keras.models import Sequential
                                                                               Launch Gallery
12 from tensorflow.keras.preprocessing import image dataset from directory
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Activation, Flatten, Dense
14 from tensorflow.keras.losses import SparseCategoricalCrossentropy
15 from tensorflow.keras.regularizers import 12
16
    import os
17
18 %matplotlib inline
19 from glob import glob
20 import seaborn as sns
21 from PIL import Image
22 np.random.seed(11)
23 from sklearn.preprocessing import StandardScaler
24 | from sklearn.model_selection import train_test_split, KFold, cross_val_score, GridSea
25 | from sklearn.metrics import accuracy_score
26
    import itertools
27
28 import keras
29 from keras.utils.np_utils import to_categorical
30 from keras.models import Sequential, Model
31 from keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPool2D
32 from keras import backend as K
33 from tensorflow.keras.layers import BatchNormalization
34 | from tensorflow.keras.optimizers import Adam, RMSprop
35 from keras.preprocessing.image import ImageDataGenerator
36 from keras.callbacks import ReduceLROnPlateau
    from keras.wrappers.scikit learn import KerasClassifier
37
    from keras.applications.inception v3 import InceptionV3
38
    from keras import backend as K
39
    import random
    import urllib.request
41
    import matplotlib.image as mpimg
42
43
44 from skimage.filters import rank, threshold_otsu
45
    from skimage import io
46 from skimage.color import rgb2gray
47
    from sklearn.cluster import KMeans
    from skimage.morphology import closing, square, disk
```

Loading Data

```
In [2]:
```

```
data_dir = pathlib.Path("../input/plantvillage-dataset/color")
train='../input/plantvillage-dataset/color'
```

In [3]:

```
dataset_path_train = os.listdir(data_dir)
print (dataset_path_train)
print("Types of classes labels found: ", len(dataset_path_train))
```

['Tomato___Late_blight', 'Tomato___healthy', 'Grape___healthy', 'Orange___
Haunglongbing_(Citrus_greening)', 'Soybean___healthy', 'Squash___Powdery_m
ildew', 'Potato__healthy', 'Corn_(maize)___Northern_Leaf_Blight', 'Tomato
__Early_blight', 'Tomato___Septoria_leaf_spot', 'Corn_(maize)___Cercospor
a_leaf_spot Gray_leaf_spot', 'Strawberry__Leaf_scorch', 'Peach___health
y', 'Apple__Apple_scab', 'Tomato___Tomato_Yellow_Leaf_Curl_Virus', 'Tomato
__Bacterial_spot', 'Apple__Black_rot', 'Blueberry__healthy', 'Cherry_
(including_sour)__Powdery_mildew', 'Peach__Bacterial_spot', 'Apple__Ced
ar_apple_rust', 'Tomato___Target_Spot', 'Pepper,_bell__healthy', 'Grape__
_Leaf_blight_(Isariopsis_Leaf_Spot)', 'Potato__Late_blight', 'Tomato__To
mato_mosaic_virus', 'Strawberry__healthy', 'Apple__healthy', 'Grape__Bl
ack_rot', 'Potato__Early_blight', 'Cherry_(including_sour)__healthy', 'C
orn_(maize)__Common_rust_', 'Grape__Esca_(Black_Measles)', 'Raspberry__
healthy', 'Tomato__Leaf_Mold', 'Tomato__Spider_mites Two-spotted_spider_
mite', 'Pepper,_bell__Bacterial_spot', 'Corn_(maize)__healthy']
Types of classes labels found: 38

In [4]:

```
image_count_train = len(list(data_dir.glob('*/*.JPG')))
print("The number of Train data:",image_count_train)
```

The number of Train data: 52803

In [5]:

```
# This Parameter we can use it in the network and model
batch_size = 32
img_height = 224
img_width = 224
```

In [6]:

Found 54305 files belonging to 38 classes. Using 43444 files for training.

```
In [7]:
```

Found 54305 files belonging to 38 classes. Using 10861 files for validation.

In [8]:

Found 54305 files belonging to 38 classes.

In [9]:

```
# Here We give the name of the data the we found the Benign dataset and malignant
class_names = train_ds.class_names
print(len(class_names))
print( class_names)
```

38

['Apple__Apple_scab', 'Apple__Black_rot', 'Apple__Cedar_apple_rust', 'A

pple__healthy', 'Blueberry__healthy', 'Cherry_(including_sour)__Powdery
mildew', 'Cherry(including_sour)__healthy', 'Corn_(maize)__Cercospora_
leaf_spot Gray_leaf_spot', 'Corn_(maize)__Common_rust_', 'Corn_(maize)__
Northern_Leaf_Blight', 'Corn_(maize)__healthy', 'Grape__Black_rot', 'Gra

pe__Esca_(Black_Measles)', 'Grape__Leaf_blight_(Isariopsis_Leaf_Spot)',
'Grape__healthy', 'Orange__Haunglongbing_(Citrus_greening)', 'Peach__Ba

cterial_spot', 'Peach__healthy', 'Pepper,_bell__Bacterial_spot', 'Peppe

r,_bell__healthy', 'Potato__Early_blight', 'Potato__Late_blight', 'Pota

to__healthy', 'Raspberry__healthy', 'Soybean__healthy', 'Squash__Powde

ry_mildew', 'Strawberry__Leaf_scorch', 'Strawberry__healthy', 'Tomato__
Bacterial_spot', 'Tomato__Early_blight', 'Tomato__Late_blight', 'Tomato__
Leaf_Mold', 'Tomato__Septoria_leaf_spot', 'Tomato__Spider_mites Two-sp

otted_spider_mite', 'Tomato__Target_Spot', 'Tomato__healthy']

In [10]:

```
for image_batch, labels_batch in train_ds.take(1):
    print(image_batch.shape)
    print(labels_batch.shape)
```

```
(32, 224, 224, 3)
(32,)
```

Creat Model

In [11]:

```
num classes = 38
2
   model = Sequential()
   model.add(layers.experimental.preprocessing.Rescaling(1./255, input_shape=(img_height
   model.add(Conv2D(16, 3, padding='same'))
 5
   model.add(Activation('relu'))
7
   model.add(MaxPooling2D())
9
   model.add(Conv2D(32, 3, padding='same'))
   model.add(Activation('relu'))
   model.add(MaxPooling2D())
11
12
   model.add(Conv2D(64, 3, padding='same'))
13
   model.add(Activation('relu'))
   model.add(MaxPooling2D())
15
16
   model.add(Dropout(0.15))
17
   model.add(Flatten())
18
   model.add(Dense(128))
19
   model.add(Activation('Softmax'))
20
21 model.add(Dense(num_classes))
```

In [12]:

In [13]:

1 model.summary()

Model: "sequential"

Layer (type)	Output Shape	Param #
rescaling (Rescaling)	(None, 224, 224, 3)	0
conv2d (Conv2D)	(None, 224, 224, 16)	448
activation (Activation)	(None, 224, 224, 16)	0
max_pooling2d (MaxPooling2D)	(None, 112, 112, 16)	0
conv2d_1 (Conv2D)	(None, 112, 112, 32)	4640
activation_1 (Activation)	(None, 112, 112, 32)	0
max_pooling2d_1 (MaxPooling2	(None, 56, 56, 32)	0
conv2d_2 (Conv2D)	(None, 56, 56, 64)	18496
activation_2 (Activation)	(None, 56, 56, 64)	0
max_pooling2d_2 (MaxPooling2	(None, 28, 28, 64)	0
dropout (Dropout)	(None, 28, 28, 64)	0
flatten (Flatten)	(None, 50176)	0
dense (Dense)	(None, 128)	6422656
activation_3 (Activation)	(None, 128)	0
dense_1 (Dense)	(None, 38)	4902

Total params: 6,451,142 Trainable params: 6,451,142 Non-trainable params: 0

```
In [15]:
```

```
epochs = 10
 2
 3 history = model.fit(
    train_ds,
 4
 5
    validation_data=val_ds,
 6
     epochs=epochs
 7 )
Epoch 1/10
1186/1358 [=====================>....] - ETA: 1:06 - loss: 1.0078 - ac
curacy: 0.7149
Cleanup called...
1358/1358 [============= ] - 567s 417ms/step - loss: 0.947
9 - accuracy: 0.7303 - val_loss: 0.4355 - val_accuracy: 0.8683
Epoch 2/10
1186/1358 [=====================>....] - ETA: 1:04 - loss: 0.3279 - ac
curacy: 0.8966
Cleanup called...
1358/1358 [================ ] - 543s 400ms/step - loss: 0.320
6 - accuracy: 0.8987 - val_loss: 0.3058 - val_accuracy: 0.9056
Epoch 3/10
curacy: 0.9454
Cleanup called...
1358/1358 [=============== ] - 544s 400ms/step - loss: 0.167
8 - accuracy: 0.9460 - val_loss: 0.3885 - val_accuracy: 0.8882
Epoch 4/10
curacy: 0.9613
Cleanup called...
1358/1358 [================= ] - 540s 398ms/step - loss: 0.109
3 - accuracy: 0.9631 - val_loss: 0.3530 - val_accuracy: 0.9100
Epoch 5/10
curacy: 0.9697
Cleanup called...
1358/1358 [================= ] - 542s 399ms/step - loss: 0.089
9 - accuracy: 0.9698 - val_loss: 0.4175 - val_accuracy: 0.8948
1186/1358 [=====================>....] - ETA: 1:03 - loss: 0.0747 - ac
curacy: 0.9751
Cleanup called...
1358/1358 [================= ] - 540s 397ms/step - loss: 0.075
6 - accuracy: 0.9754 - val_loss: 0.3449 - val_accuracy: 0.9175
Epoch 7/10
1186/1358 [=====================>....] - ETA: 1:04 - loss: 0.0560 - ac
curacy: 0.9820
Cleanup called...
```

```
1358/1358 [============ ] - 541s 398ms/step - loss: 0.057
4 - accuracy: 0.9816 - val_loss: 0.3413 - val_accuracy: 0.9188
Epoch 8/10
curacy: 0.9823
Cleanup called...
1358/1358 [============= ] - 546s 402ms/step - loss: 0.052
7 - accuracy: 0.9830 - val_loss: 0.3569 - val_accuracy: 0.9200
Epoch 9/10
curacy: 0.9843
Cleanup called...
1358/1358 [================ ] - 547s 403ms/step - loss: 0.051
1 - accuracy: 0.9839 - val_loss: 0.4819 - val_accuracy: 0.9010
Epoch 10/10
curacy: 0.9847
Cleanup called...
1358/1358 [============= ] - 544s 401ms/step - loss: 0.042
3 - accuracy: 0.9860 - val_loss: 0.4520 - val_accuracy: 0.9125
```

In [16]:

```
acc = history.history['accuracy']
   val_acc = history.history['val_accuracy']
 3
4 loss = history.history['loss']
 5
   val_loss = history.history['val_loss']
 6
7
   epochs_range = range(epochs)
8
9
   plt.figure(figsize=(8, 8))
10
   plt.subplot(1, 2, 1)
plt.plot(epochs_range, acc, label='Training Accuracy')
   plt.plot(epochs_range, val_acc, label='Validation Accuracy')
   plt.legend(loc='lower right')
13
   plt.title('Training and Validation Accuracy')
14
15
16
   plt.subplot(1, 2, 2)
17
   plt.plot(epochs_range, loss, label='Training Loss')
18 plt.plot(epochs_range, val_loss, label='Validation Loss')
   plt.legend(loc='upper right')
19
   plt.title('Training and Validation Loss')
   plt.show()
21
```



In [17]:

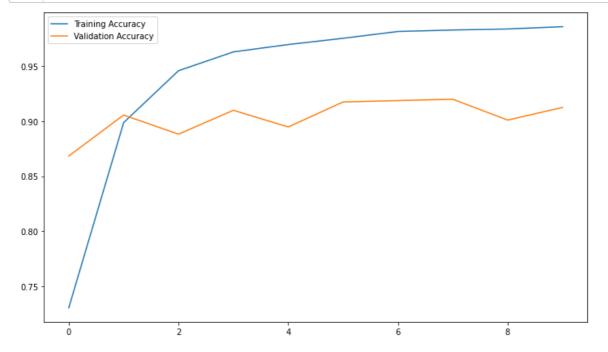
```
preformance = model.evaluate(val_ds)
```

```
In [ ]:
```

1 preformance

In [18]:

```
acc = history.history['accuracy']
 2
   val_acc = history.history['val_accuracy']
 4 loss = history.history['loss']
   val_loss = history.history['val_loss']
   no_of_epochs = history.params['epochs']
 7
   len(history.history['accuracy']) == no_of_epochs
   plt.figure(figsize=(12, 7))
 8
 9
   plt.plot(range(no_of_epochs), acc, label='Training Accuracy')
10
11
   plt.plot(range(no_of_epochs), val_acc, label='Validation Accuracy')
12
13 plt.legend()
14 plt.show()
```



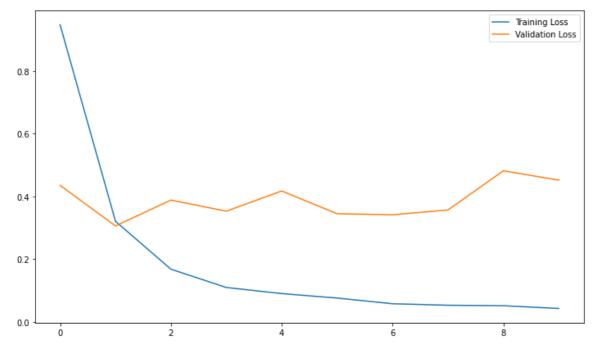
In [19]:

```
plt.figure(figsize=(12, 7))

plt.plot(range(no_of_epochs), loss, label='Training Loss')

plt.plot(range(no_of_epochs), val_loss, label='Validation Loss')

plt.legend()
plt.show()
```



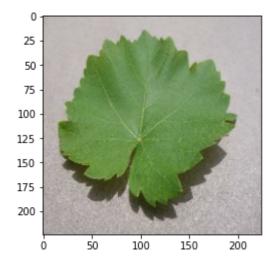
In []:

1

In [20]:

```
labels = train_ds.class_names
 2
   labels
 3
   for images_batch, labels_batch in test.take(1):
4
 5
        image = images_batch[0].numpy().astype('uint8')
        label = labels_batch[0].numpy()
 6
 7
        plt.imshow(image)
 8
9
        print("actual label:", labels[label])
10
        batch_prediction = model.predict(images_batch)
11
        print("predicted label:", labels[np.argmax(batch_prediction[0])])
12
```

actual label: Grape___healthy
predicted label: Grape___healthy



In [22]:

```
from tensorflow import expand_dims, newaxis
1
2
   def predict(model, img):
 3
4
 5
       img_array = img.numpy()
 6
       img_array = expand_dims(img_array, 0)
7
8
       predictions = model.predict(img_array)
9
       predicted_class = labels[np.argmax(predictions[0])]
10
       confidence = round( (np.max(predictions[0])), 2)
11
12
       return predicted_class, confidence
13
```

In [23]:

```
plt.figure(figsize=(15, 15))
 2
   for images, lbs in val_ds.take(1):
 3
 4
        for i in range(9):
 5
 6
            plt.subplot(3, 3, i + 1)
 7
            plt.imshow(images[i].numpy().astype("uint32"))
 8
9
            predicted_class, confidence = predict(model, images[i])
            actual_class = labels[lbs[i]]
10
11
            plt.title(f"Actual: {actual_class},\n Predicted: {predicted_class}.\n Confide
12
13
           plt.axis("off")
14
```

Actual: Tomato__Tomato_Yellow_Leaf_Curl_Virus, Predicted: Tomato__Tomato_Yellow_Leaf_Curl_Virus. Confidence: 36.45000076293945%



Actual: Tomato__healthy, Predicted: Tomato__healthy. Confidence: 45.4900016784668%



Actual: Corn_(maize)___healthy,

Actual: Pepper_bell healthy, Predicted: Pepper, bell healthy. Confidence: 20.56999969482422%



Actual: Orange__Haunglongbing_(Citrus_greening), Predicted: Orange__Haunglongbing_(Citrus_greening). Confidence: 27.690000534057617%



Actual: Potato Early_blight, Actual: Orange_ Predicted: Pepper,_bell_Bacterial_spot. Predicted: Orange_ Confidence: 29.489999771118164% Confidence



Actual: Orange__Haunglongbing_(Citrus_greeningActual: Tomato__Tomato_Yellow_Leaf_Curl_Virus,
Predicted: Orange__Haunglongbing_(Citrus_greeningActual: Tomato__Tomato_Yellow_Leaf_Curl_Virus.
Confidence: 21.469999313354492%
Confidence: 27.040000915527344%





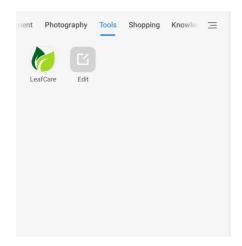


In [25]:

1 model.save("leafModel.h5")

In []:

1



App Logo



Classified as: Pepper,_bell__Bacterial_spot

Take Picture

Launch Gallery