```
from google.colab import drive
drive.mount('/content/drive')

Mounted at /content/drive

#https://www.kaggle.com/datasets/vbookshelf/rice-leaf-diseases
#https://www.kaggle.com/datasets/emmarex/plantdisease

# Imkdir -p ~/.kaggle
# Icp kaggle.json ~/.kaggle/

# !kaggle datasets download -d vbookshelf/rice-leaf-diseases
# # !kaggle datasets download -d emmarex/plantdisease

import zipfile
zip.ref = zipfile.Zipfile('/content/drive/MyDrive/Crop-Disease/Plant_leaf_diseases_dataset_without_augmentation.zip', 'r')
zip.ref.close()
```

→ Importing the Library

```
import matplotlib.pyplot as plt
import numpy as np
import cv2
import os
import PIL

import tensorflow as tf
from tensorflow import keras
from keras import Sequential
from keras.layers import Dense, Conv2D, MaxPooling2D, Flatten, BatchNormalization, Dropout
```

▼ Labeling the Data

```
data_dir = "/content/Plant_leave_diseases_dataset_without_augmentation"
import pathlib
data_dir=pathlib.Path(data_dir)
data_dir
     PosixPath('/content/Plant_leave_diseases_dataset_without_augmentation')
# dict={
     "bacteria":list(data_dir.glob("Bacterial leaf blight/*")),
     "brown":list(data_dir.glob("Brown spot/*")),
     "smut":list(data_dir.glob("Leaf smut/*"))
#
dict={
   "appleScab":list(data_dir.glob("Apple__Apple_scab/*")),
   "appleBlackRot":list(data_dir.glob("Apple__Black_rot/*")),
   "appleRust":list(data_dir.glob("Apple__Cedar_apple_rust/*")),
   "appleHealthy":list(data_dir.qlob("Apple__Healthy/*")),
   "backWithoutLeaves":list(data_dir.glob("Background_without_leaves/*")),
   "blueberryHealthy":list(data_dir.glob("Blueberry_healthy/*")),
   "cherryHealthy":list(data_dir.glob("Cherry_healthy/*")),
   "cherryPowderyMildew":list(data_dir.glob("Cherry__Powdery_mildew/*")),
   "cornCercospora":list(data_dir.glob("Corn___Cercospora_leaf_spot Gray_leaf_spot/*")),
   "cornRust":list(data_dir.glob("Corn__Common_rust/*")),
   "cornHealthy":list(data_dir.glob("Corn_healthy/*")),
   "cornBlight":list(data_dir.glob("Corn__Northern_Leaf_Blight/*")),
   "grapeBlackRot":list(data_dir.glob("Grape___Black_rot/*")),
   "grapeEsca":list(data_dir.glob("Grape___Esca_(Black_Measles)/*")),
   "grapeHealthy":list(data_dir.glob("Grape__healthy/*")),
   "grapeBlight":list(data_dir.glob("Grape__Leaf_blight_(Isariopsis_Leaf_Spot)/*")),
   "orangeHaun":list(data_dir.glob("Orange___Haunglongbing_(Citrus_greening)/*")),
   "peachBacteria":list(data_dir.qlob("Peach__Bacterial_spot/*")),
   "peachHealthy":list(data_dir.glob("Peach__Healthy/*")),
   "pepperBacteria":list(data_dir.glob("Pepper_bell_Bacterial_spot/*")),
   "pepperHealthy":list(data_dir.glob("Pepper,_bell__healthy/*")),
   "potatoEarlyBlight":list(data_dir.glob("Potato___Early_blight/*")),
   "potatoHealthy":list(data_dir.glob("Potato__healthy/*")),
   "potatoLateBlight":list(data_dir.glob("Potato__Late_blight/*")),
   "raspberryHealthy":list(data_dir.glob("Raspberry_healthy/*")),
   "soybeanHealthy":list(data_dir.glob("Soybean_healthy/*")),
   "squashPowderyMildew":list(data_dir.glob("Squash__Powdery_mildew/*")),
   "strawberryHealthy":list(data_dir.glob("Strawberry_healthy/*")),
   "strawberryLeafScorch":list(data_dir.glob("Strawberry__Leaf_scorch/*")),
   "tomatoBacteria":list(data_dir.glob("Tomato___Bacterial_spot/*")),
   "tomatoEarlyBlight":list(data_dir.glob("Tomato___Early_blight/*")),
   "tomatoHealthy":list(data_dir.glob("Tomato_healthy/*")),
   "tomatoLateBlight":list(data_dir.glob("Tomato__Late_blight/*")),
   "tomatoLeafMold":list(data_dir.glob("Tomato__Leaf_Mold/*")),
   "tomatoSeptoria":list(data_dir.glob("Tomato__Septoria_leaf_spot/*")),
   "tomatoSpiderMites":list(data_dir.qlob("Tomato__Spider_mites Two-spotted_spider_mite/*")),
```

```
"tomatoTargetSpot":list(data_dir.glob("Tomato__Target_Spot/*")),
   "tomatoMosaic":list(data_dir.glob("Tomato__Tomato_mosaic_virus/*")),
   "tomatoYellowLeaf":list(data_dir.glob("Tomato__Tomato_Yellow_Leaf_Curl_Virus/*"))
# labels_dict = {
     'bacteria': 0,
     'brown': 1,
     'smut': 2,
# }
labels_dict = {
   'appleScab': O,
   'appleBlackRot': 1,
   'appleRust': 2,
   'appleHealthy': 3,
   'backWithoutLeaves': 4,
   'blueberryHealthy': 5,
   'cherryHealthy': 6,
   'cherryPowderyMildew': 7,
   'cornCercospora': 8,
   'cornRust': 9,
   'cornHealthy': 10,
   'cornBlight': 11,
   'grapeBlackRot': 12,
   'grapeEsca': 13,
   'grapeHealthy': 14,
   'grapeBlight': 15,
   'orangeHaun': 16,
   'peachBacteria': 17,
   'peachHealthy': 18,
   'pepperBacteria': 19,
   'pepperHealthy': 20,
   'potatoEarlyBlight': 21,
   'potatoHealthy': 22,
   'potatoLateBlight': 23,
   'raspberryHealthy': 24,
   'soybeanHealthy': 25,
   'squashPowderyMildew': 26,
   'strawberryHealthy': 27,
   'strawberryLeafScorch': 28,
   'tomatoBacteria': 29,
   'tomatoEarlyBlight': 30,
   'tomatoHealthy': 31,
   'tomatoLateBlight': 32,
   'tomatoLeafMold': 33,
   'tomatoSeptoria': 34,
   'tomatoSpiderMites': 35,
   'tomatoTargetSpot': 36,
   'tomatoMosaic': 37,
```

```
'tomatoYellowLeaf': 38,
}
```

→ Splitting the Data

```
X, y = [], []
for name, images in dict.items():
   for image in images:
      img = cv2.imread(str(image))
      resized_img = cv2.resize(img,(64,64))
      X.append(resized_img)
      y.append(labels_dict[name])
X = np.array(X)
y = np.array(y)
# # Store X and y in a file using numpy's save function
# np.save('X64.npy', X)
# np.save('y64.npy', y)
# # Load X and y from file
# X = np.load('/content/drive/MyDrive/Crop-Disease/CNN/X.npy')
# y = np.load('/content/drive/MyDrive/Crop-Disease/CNN/y.npy')
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=0)
del X
del y
X_train_scaled = X_train / 255
X_test_scaled = X_test / 255
del X_train
del X_test
```

```
X_train_scaled.shape
(40082, 64, 64, 3)

X_test_scaled.shape
(13361, 64, 64, 3)

# np.save('X64_train_scaled.npy', X_train_scaled)
# np.save('y64_train_npy', y_train)

# np.save('Y64_test_scaled.npy', X_test_scaled)
# np.save('Y64_test_scaled.npy', y_test)
```

Creating the Model

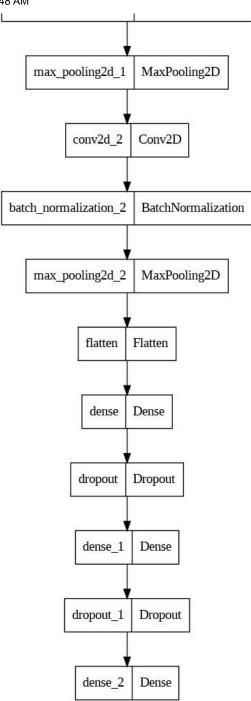
```
num_classes = 39
model = Sequential()
model.add(Conv2D(32,kernel_size=(3,3),padding='valid',activation='relu',input_shape=(64,64,3)))
model.add(BatchNormalization())
model.add(MaxPooling2D(pool_size=(2,2),strides=2,padding='valid'))
model.add(Conv2D(64,kernel_size=(3,3),padding='valid',activation='relu'))
model.add(BatchNormalization())
model.add(MaxPooling2D(pool_size=(2,2),strides=2,padding='valid'))
model.add(Conv2D(128,kernel_size=(3,3),padding='valid',activation='relu'))
model.add(BatchNormalization())
model.add(MaxPooling2D(pool_size=(2,2),strides=2,padding='valid'))
model.add(Flatten())
model.add(Dense(128,activation='relu'))
model.add(Dropout(0.1))
model.add(Dense(64,activation='relu'))
model.add(Dropout(0.1))
model.add(Dense(num_classes))
model.summary()
```

Model: "sequential"

Layer (type)	Output Shape	Param #	
conv2d (Conv2D)	(None, 62, 62, 3	2) 896	
batch_normalization (E ormalization)	BatchN (None, 62, 62	2, 32) 128	
max_pooling2d (MaxPo)	poling2D (None, 31, 3	31, 32) <i>O</i>	
conv2d_1 (Conv2D)	(None, 29, 29, 6	54) 18496	
batch_normalization_1 hNormalization)	(Batc (None, 29, 29	. 64) 256	
max_pooling2d_1 (Max 2D)	Pooling (None, 14, 1	4, 64) 0	
conv2d_2 (Conv2D)	(None, 12, 12,	128) 73856	
batch_normalization_2 hNormalization)	(Batc (None, 12, 12	, 128)	
max_pooling2d_2 (Max 2D)	Pooling (None, 6, 6,	128) 0	
flatten (Flatten)	(None, 4608)	0	
dense (Dense)	(None, 128)	589952	
dropout (Dropout)	(None, 128)	0	
dense_1 (Dense)	(None, 64)	8256	
dropout_1 (Dropout)	(None, 64)	0	
dense_2 (Dense)	(None, 39)	2535	
Total params: 694,887 Trainable params: 694 Non-trainable params:	,439	=======================================	=======

from tensorflow.keras.utils import plot_model

plot_model(model)



Compiling and Training

model.compile(optimizer='adam',loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),metrics=['accuracy'])

history = model.fit(X_train_scaled, y_train, epochs=20, validation_data=(X_test_scaled, y_test))

```
Epoch 1/20
1253/1253 [============================  - 28s 12ms/step - loss: 1.1292 - accuracy: 0.6807 - val_loss: 1.8164 - val_accuracy: 0.5831
Epoch 2/20
Epoch 3/20
Epoch 4/20
Epoch 5/20
Epoch 6/20
1253/1253 [========================== ] - 12s 10ms/step - loss: 0.1847 - accuracy: 0.9425 - val_loss: 0.3560 - val_accuracy: 0.8996
Epoch 7/20
Epoch 8/20
Epoch 9/20
Epoch 10/20
Epoch 11/20
Epoch 12/20
Epoch 13/20
```

model.save('CNNModel9894.h5')

from tensorflow.keras.models import load_model

Load the saved model model = load_model('/content/drive/MyDrive/Crop-Disease/CNNModel9894.h5')

import pandas as pd from sklearn.metrics import classification_report import seaborn as sns

```
y_prob = model.predict(X_test_scaled)
y_pred = np.argmax(y_prob, axis=1)
report = classification_report(y_test, y_pred, output_dict=True)
```

418/418 [===========] - 9s 3ms/step

create a dataframe from the classification report
df = pd.DataFrame(report).transpose()

df.tail()

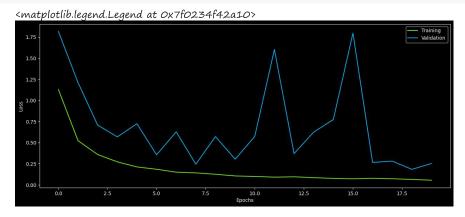
	precision	recall	f1-score	support	
37	•	0.879121		91.000000	
38	0.994830	0.986091	0.990441	1366.000000	
accuracy	0.945588	0.945588	0.945588	0.945588	
macro avg	0.928437	0.926737	0.925231	13361.000000	
weighted avg	0.948465	0.945588	0.944935	13361.000000	

Graph

```
loss = history.history['loss']
val_loss = history.history['val_loss']

epochs = range(len(loss))

fig = plt.figure(figsize=(15,6))
plt.plot(epochs,loss,c="lawngreen",label="Training")
plt.plot(epochs,val_loss,c="deepskyblue",label="Validation")
plt.xlabel("Epochs")
plt.ylabel("Loss")
plt.legend()
```



```
acc = history.history['accuracy']
val_acc = history.history['val_accuracy']
epochs = range(len(acc))
fig = plt.figure(figsize=(15,6))
plt.plot(epochs,acc,c="deeppink",label="Training")
plt.plot(epochs,val_acc,c="yellow",label="Validation")
plt.xlabel("Epochs")
```

plt.ylabel("Accuracy") plt.legemdtplotlib.legend.Legend at 0x7f023508bfd0>

