```
from google.colab import drive
drive.mount('/content/drive')
    Drive already mounted at /content/drive; to attempt to forcibly remount, call
!unzip /content/drive/MyDrive/Tomato
import tensorflow as tf
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.preprocessing import image
import numpy as np
import os
import random
from shutil import copyfile
import matplotlib.image as mpimg
import matplotlib.pyplot as plt
from glob import glob
from tensorflow.keras.models import Model, Sequential
from tensorflow.keras.layers import Conv2D, Activation, MaxPooling2D, Dropout, Dens
from tensorflow.keras.layers import GlobalAveragePooling2D, Flatten, BatchNormaliza
from tensorflow.keras.applications.inception v3 import InceptionV3
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.callbacks import ModelCheckpoint, ReduceLROnPlateau, EarlySto
from keras.utils.np_utils import to_categorical
to create = [
'leaf disease',
'leaf disease/training',
'leaf disease/testing',
'leaf disease/training/Tomato Bacterial spot',
'leaf_disease/training/Tomato Late blight',
'leaf disease/training/Tomato Early blight',
'leaf disease/training/Tomato healthy',
'leaf disease/training/Tomato Yellow Leaf Curl Virus',
'leaf disease/training/Tomato Leaf Mold',
'leaf disease/testing/Tomato Bacterial spot',
'leaf disease/testing/Tomato Late blight',
'leaf disease/testing/Tomato Early blight',
'leaf disease/testing/Tomato healthy',
'leaf disease/testing/Tomato Yellow Leaf Curl Virus',
'leaf disease/testing/Tomato Leaf Mold'
for directory in to create:
    os.mkdir(directory)
    print(directory, 'created')
 except:
    print(directory, 'failed')
```

```
leaf disease created
    leaf disease/training created
    leaf disease/testing created
    leaf disease/training/Tomato Bacterial spot created
    leaf disease/training/Tomato Late blight created
    leaf disease/training/Tomato Early blight created
    leaf disease/training/Tomato healthy created
    leaf disease/training/Tomato Yellow Leaf Curl Virus created
    leaf disease/training/Tomato Leaf Mold created
    leaf disease/testing/Tomato Bacterial spot created
    leaf disease/testing/Tomato Late blight created
    leaf disease/testing/Tomato Early blight created
    leaf disease/testing/Tomato healthy created
    leaf disease/testing/Tomato Yellow Leaf Curl Virus created
    leaf disease/testing/Tomato Leaf Mold created
def split data(SOURCE, TRAINING, TESTING, SPLIT SIZE):
    all files = []
    for file name in os.listdir(SOURCE):
        file path = SOURCE + file name
        if os.path.getsize(file path):
            all files.append(file name)
        else:
            print('{} is zero length, so ignoring'.format(file name))
    n files = len(all files)
    split point = int(n files * SPLIT SIZE)
    shuffled = random.sample(all files, n files)
    train set = shuffled[:split point]
    test set = shuffled[split point:]
    for file name in train set:
        copyfile(SOURCE + file name, TRAINING + file name)
    for file_name in test_set:
        copyfile(SOURCE + file name, TESTING + file name)
SOURCE_DIR = "Tomato/Tomato_Late blight/"
TRAINING DIR = "leaf disease/training/Tomato Late blight/"
TESTING DIR = "leaf disease/testing/Tomato Late blight/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
SOURCE_DIR = "Tomato/Tomato_Bacterial spot/"
TRAINING DIR = "leaf disease/training/Tomato Bacterial spot/"
TESTING DIR = "leaf disease/testing/Tomato Bacterial spot/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
```

https://colab.research.google.com/drive/1cEpGJeq1zNPw1mIW756Z3DAwxa3hYj8S#printMode=true

```
TRAINING DIR = "leaf disease/training/Tomato Early blight/"
TESTING DIR = "leaf disease/testing/Tomato Early blight/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
SOURCE DIR = "Tomato/Tomato healthy/"
TRAINING DIR = "leaf disease/training/Tomato healthy/"
TESTING DIR = "leaf disease/testing/Tomato healthy/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
SOURCE DIR = "Tomato/Tomato Yellow Leaf Curl Virus/"
TRAINING DIR = "leaf disease/training/Tomato Yellow Leaf Curl Virus/"
TESTING DIR = "leaf disease/testing/Tomato Yellow Leaf Curl Virus/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
SOURCE DIR = "Tomato/Tomato Leaf Mold/"
TRAINING DIR = "leaf disease/training/Tomato Leaf Mold/"
TESTING DIR = "leaf disease/testing/Tomato Leaf Mold/"
split size = .5
split data(SOURCE DIR, TRAINING DIR, TESTING DIR, split size)
#!unzip /content/drive/MyDrive/leaf disease
data dir = "leaf disease/training"
classes = os.listdir(data dir)
classes
     ['Tomato Bacterial spot',
      'Tomato Early blight',
      'Tomato Yellow Leaf Curl Virus',
      'Tomato Leaf Mold',
      'Tomato healthy',
      'Tomato Late blight']
train datagen = ImageDataGenerator(rescale=1./255,
                                   shear range=0.2,
                                   zoom range=0.2,
                                   width shift range=0.2,
                                   height shift range=0.2,
                                   fill mode='nearest')
valid datagen = ImageDataGenerator(rescale=1./255)
batch size = 128
training_dir = 'leaf_disease/training'
validation dir = 'leaf disease/testing'
training generator = train datagen.flow from directory(training dir,
                                                  target size=(256, 256),
                                                  batch size=batch size,
```

```
TransferLearning_VGG16.ipynb - Colaboratory
                                                  class mode='categorical')
valid generator = valid datagen.flow from directory(validation dir,
                                             target size=(256, 256),
                                             batch size=batch size,
                                             class mode='categorical',
                                             shuffle=False)
    Found 4491 images belonging to 6 classes.
    Found 4495 images belonging to 6 classes.
class dict = training generator.class indices
print(class dict)
    {'Tomato Bacterial spot': 0, 'Tomato Early blight': 1, 'Tomato Late blight': 2
target names = list(class dict.keys())
print(target names)
    ['Tomato Bacterial spot', 'Tomato Early blight', 'Tomato Late blight', 'Tomato
train_num = training_generator.samples
valid num = valid generator.samples
from tensorflow.keras.applications.vgg16 import VGG16, preprocess input
pre trained model = VGG16(input shape=(256, 256, 3), include top=False, weights="im
    Downloading data from <a href="https://storage.googleapis.com/tensorflow/keras-applicat">https://storage.googleapis.com/tensorflow/keras-applicat</a>
    for layer in pre trained model.layers:
   print(layer.name)
   layer.trainable = False
print(len(pre trained model.layers))
    input 1
    block1 conv1
    block1 conv2
    block1 pool
    block2_conv1
    block2 conv2
    block2 pool
    block3 conv1
    block3 conv2
    block3 conv3
    block3 pool
    block4 conv1
    block4 conv2
    block4 conv3
```

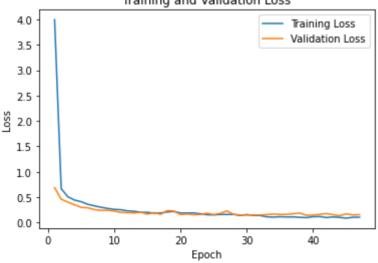
block4 pool block5 conv1

```
block5 conv2
   block5 conv3
   block5 pool
   19
last layer = pre trained model.get layer('block5 pool')
print('last layer output shape:', last layer.output shape)
last output = last layer.output
   last layer output shape: (None, 8, 8, 512)
from tensorflow.keras import layers
from tensorflow.keras import Model
from tensorflow.keras.applications.vgg16 import VGG16, preprocess input
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.callbacks import ReduceLROnPlateau, EarlyStopping
# Flatten the output layer to 1 dimension
x=layers.Flatten()(last output)
#x = layers.GlobalMaxPooling2D()(last output)
# Add a fully connected layer with 512 hidden units and ReLU activation
x = layers.Dense(512, activation='relu')(x)
# Add a dropout rate of 0.5
x = layers.Dropout(0.2)(x)
# Add a final sigmoid layer for classification
x = layers.Dense(6, activation='softmax')(x)
# Configure and compile the model
model = Model(pre trained model.input, x)
optimizer = Adam(lr=0.001, beta_1=0.9, beta_2=0.999, epsilon=None, decay=0.0, amsgr
model.compile(loss='categorical crossentropy',
           optimizer=optimizer,
           metrics=['accuracy'])
history = model.fit(training generator,
                epochs = 47, validation data = valid generator,
                verbose = 1, steps per epoch=(train num // batch size),
                validation steps=(valid num// batch size))
  Epoch 7/47
   Epoch 8/47
   35/35 [============ ] - 121s 3s/step - loss: 0.3090 - accurac
   Epoch 9/47
   Epoch 10/47
   Epoch 11/47
   35/35 [============== ] - 121s 3s/step - loss: 0.2582 - accurac
   Epoch 12/47
   35/35 [============== ] - 121s 3s/step - loss: 0.2301 - accurac
   Epoch 13/47
```

```
Epoch 14/47
  35/35 [============== ] - 123s 4s/step - loss: 0.1967 - accurac
  Epoch 15/47
  Epoch 16/47
  Epoch 17/47
  Epoch 18/47
  Epoch 19/47
  35/35 [============== ] - 121s 3s/step - loss: 0.2226 - accurac
  Epoch 20/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1991 - accurac
  Epoch 21/47
  Epoch 22/47
  Epoch 23/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1720 - accurac
  Epoch 24/47
  Epoch 25/47
  Epoch 26/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1421 - accurac
  Epoch 27/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1580 - accurac
  Epoch 28/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1734 - accurac
  Epoch 29/47
  Epoch 30/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1564 - accurac
  Epoch 31/47
  Epoch 32/47
  Epoch 33/47
  Epoch 34/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1085 - accurac
  Epoch 35/47
  35/35 [============== ] - 122s 3s/step - loss: 0.1132 - accurac
import pickle
filename = 'VGG16 model.pkl'
pickle.dump(history.history, open(filename, 'wb'))
tf.keras.models.save model(model, 'transferlearning VGG16.hdf5')
acc = history.history['accuracy']
val acc = history.history['val accuracy']
loss = history.history['loss']
val loss = history.history['val loss']
epochs = range(1, len(loss) + 1)
```

```
#accuracy proc
plt.plot(epochs, acc, label='Training Accuracy')
plt.plot(epochs, val acc, label='Validation Accuracy')
plt.title('Training and Validation Accuracy')
plt.ylabel('Accuracy')
plt.xlabel('Epoch')
plt.savefig(fname='Training Validation Accuracy VGG16')
plt.legend()
plt.figure()
#loss plot
plt.plot(epochs, loss, label='Training Loss')
plt.plot(epochs, val loss, label='Validation Loss')
plt.title('Training and Validation Loss')
plt.xlabel('Epoch')
plt.ylabel('Loss')
plt.legend()
plt.savefig(fname='Training Validation Loss VGG16')
plt.show()
```





```
from sklearn.metrics import confusion_matrix from sklearn.metrics import classification_report from sklearn.metrics import accuracy_score from sklearn.metrics import roc_curve, auc from sklearn.metrics import roc auc score https://colab.research.google.com/drive/1cEpGJeq1zNPw1mIW756Z3DAwxa3hYj8S#printMode=true
```

```
from sklearn import preprocessing
Y pred = model.predict(valid generator, valid num // batch size+1)
y pred = np.argmax(Y pred, axis=1)
print('Confusion Matrix')
print(confusion matrix(valid generator.classes, y pred))
print('Classification Report')
target names = ['Tomato Bacterial spot', 'Tomato Early blight',
                'Tomato Late blight', 'Tomato Leaf Mold',
                'Tomato Yellow Leaf Curl Virus', 'Tomato healthy']
print(classification report(valid generator.classes, y pred,
                            target names=target names))
    Confusion Matrix
    [[1014
            18
                18
                       0
                             7
                                  71
         6 427
                 42
                      12
                             1
                                 121
         1 41 903
                       6
                            0
                                  4 ]
         1
             9
                16 440
                            0
                                101
     Γ
                        6 672
             16
                   4
                                  21
     [
             1
                   0
                        1
                            0
                               79411
         ()
    Classification Report
                                                                   support
                                   precision
                                               recall f1-score
            Tomato Bacterial spot
                                        0.99
                                                  0.95
                                                            0.97
                                                                      1064
              Tomato Early blight
                                                  0.85
                                                            0.84
                                                                       500
                                        0.83
               Tomato Late blight
                                        0.92
                                                  0.95
                                                            0.93
                                                                       955
                 Tomato Leaf Mold
                                        0.95
                                                  0.92
                                                            0.94
                                                                       476
                                                            0.97
    Tomato Yellow Leaf Curl Virus
                                        0.99
                                                  0.95
                                                                       704
                   Tomato healthy
                                        0.96
                                                  1.00
                                                            0.98
                                                                       796
                                                            0.95
                                                                      4495
                         accuracy
                                        0.94
                                                  0.94
                                                            0.94
                                                                      4495
                        macro avq
                                        0.95
                                                  0.95
                                                            0.95
                                                                      4495
                     weighted avg
all labels = ['Tomato Bacterial spot', 'Tomato Early blight', 'Tomato Late blight',
       'Tomato_Leaf_Mold', 'Tomato_Yellow_Leaf_Curl_Virus', 'Tomato_healthy']
fig, c ax = plt.subplots(1,1, figsize = (12, 8))
def multiclass roc auc score(y test, y pred, average="macro"):
   lb = preprocessing.LabelBinarizer()
   #lb = LabelBinarizer()
   lb.fit(y test)
   y test = lb.transform(y test)
   y pred = lb.transform(y pred)
   for (idx, c label) in enumerate(all labels): # all labels: no of the labels
        fpr, tpr, thresholds = roc_curve(y_test[:,idx].astype(int), y_pred[:,idx])
       plt.xlabel('False Positive Rate')
       plt.ylabel('True Positive Rate')
       plt.legend(loc = 'lower right')
       plt.title('Receiver Operating Characteristic')
        plt.plot(fpr, tpr, label = '%s (AUC:%0.2f)' % (c label, auc(fpr, tpr)))
   plt.plot(fpr, fpr, 'b-', label = 'Random Guessing')
   plt.savefig(fname='roc auc VGG16')
```

```
return roc_auc_score(y_test, y_pred, average=average)
```

```
# calling
valid_generator.reset() # resetting generator
y_pred = model.predict(valid_generator, verbose = True)
y_pred = np.argmax(y_pred, axis=1)
multiclass_roc_auc_score(valid_generator.classes, y_pred)
```

36/36 [==========] - 39s 1s/step No handles with labels found to put in legend. 0.9636495147919181

