Import neccessary packages

```
! pip install kaggle
     Looking in indexes: <a href="https://pypi.org/simple">https://us-python.pkg.dev/colab-wheels/pub</a>.
     Requirement already satisfied: kaggle in /usr/local/lib/python3.9/dist-packages (1.5.13)
     Requirement already satisfied: six>=1.10 in /usr/local/lib/python3.9/dist-packages (from
     Requirement already satisfied: certifi in /usr/local/lib/python3.9/dist-packages (from |
     Requirement already satisfied: urllib3 in /usr/local/lib/python3.9/dist-packages (from |
     Requirement already satisfied: python-slugify in /usr/local/lib/python3.9/dist-packages
     Requirement already satisfied: tqdm in /usr/local/lib/python3.9/dist-packages (from kags
     Requirement already satisfied: requests in /usr/local/lib/python3.9/dist-packages (from
     Requirement already satisfied: python-dateutil in /usr/local/lib/python3.9/dist-packages
     Requirement already satisfied: text-unidecode>=1.3 in /usr/local/lib/python3.9/dist-pack
     Requirement already satisfied: idna<4,>=2.5 in /usr/local/lib/python3.9/dist-packages (1
     Requirement already satisfied: charset-normalizer~=2.0.0 in /usr/local/lib/python3.9/dis
from google.colab import drive
drive.mount('/content/gdrive')
     Mounted at /content/gdrive
from google.colab import files
files.upload()
     Choose Files No file chosen
                                       Upload widget is only available when the cell has been
     executed in the current browser session. Please rerun this cell to enable.
     Saving kaggle.json to kaggle (1).json
     {'kaggle ison'.
!ls -lha kaggle.json
     -rw-r--r-- 1 root root 75 Apr 20 23:00 kaggle.json
!pip install -q kaggle
!mkdir -p ~/.kaggle
!cp kaggle.json ~/.kaggle/
!chmod 600 /root/.kaggle/kaggle.json
!pwd
```

/content

```
!kaggle datasets download -d emmarex/plantdisease
     plantdisease.zip: Skipping, found more recently modified local copy (use --force to force
! unzip plantdisease.zip
     Archive: plantdisease.zip
     replace PlantVillage/Pepper bell Bacterial spot/0022d6b7-d47c-4ee2-ae9a-392a53f48647
     replace PlantVillage/Pepper__bell___Bacterial_spot/006adb74-934f-448f-a14f-62181742127b_
from google.colab import drive
drive.mount('/content/drive')
     Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mour
import numpy as np
import pickle
import cv2
from os import listdir
from sklearn.preprocessing import LabelBinarizer
from keras.models import Sequential
from tensorflow.keras.layers import BatchNormalization
from keras.layers.convolutional import Conv2D
from keras.layers.convolutional import MaxPooling2D
from keras.layers.core import Activation, Flatten, Dropout, Dense
from keras import backend as K
from keras.preprocessing.image import ImageDataGenerator
from keras.optimizers import Adam
from keras.preprocessing import image
from tensorflow.keras.preprocessing.image import img to array
from sklearn.preprocessing import MultiLabelBinarizer
from sklearn.model_selection import train_test_split
import matplotlib.pyplot as plt
EPOCHS = 25
INIT LR = 1e-3
BS = 32
default image size = tuple((256, 256))
image size = 0
directory root = '/content/plantvillage'
width=256
height=256
depth=3
```

Function to convert images to array

```
def convert_image_to_array(image_dir):
    try:
        image = cv2.imread(image_dir)
        if image is not None :
            image = cv2.resize(image, default_image_size)
            return img_to_array(image)
        else :
            return np.array([])
    except Exception as e:
        print(f"Error : {e}")
        return None
```

Fetch images from directory

```
import os
image list, label list = [], []
try:
   print("[INFO] Loading images ...")
   root_dir = os.listdir(directory_root)
   for directory in root dir :
        # remove .DS_Store from list
        if directory == ".DS Store" :
            root dir.remove(directory)
   for plant folder in root dir :
        plant_folder_path = os.path.join(directory_root, plant_folder)
        if os.path.isdir(plant folder path):
            plant_disease_folder_list = os.listdir(plant_folder_path)
            for disease folder in plant disease folder list :
                # remove .DS_Store from list
                if disease folder == ".DS Store" :
                    plant disease folder list.remove(disease folder)
            for plant disease folder in plant disease folder list:
                print(f"[INFO] Loading images: {plant_disease_folder} ...")
                plant disease folder path = os.path.join(plant folder path, plant disease fol
                if os.path.isdir(plant disease folder path):
                    print(f"[INFO] Processing {plant_disease_folder} ...")
                    plant disease image list = os.listdir(plant disease folder path)
                    for single plant disease image in plant disease image list :
                        if single plant disease image == ".DS Store" :
                            plant_disease_image_list.remove(single_plant_disease_image)
```

```
for image in plant disease image list[:200]:
                        image_directory = os.path.join(plant_disease_folder_path, image)
                        if image_directory.endswith(".jpg") or image_directory.endswith(".JPG
                            image list.append(convert image to array(image directory))
                            label list.append(plant disease folder)
   print("[INFO] Image loading completed")
except Exception as e:
   print(f"Error : {e}")
     [INFO] Loading images ...
     [INFO] Loading images: Tomato Spider mites Two spotted spider mite ...
     [INFO] Processing Tomato Spider mites Two spotted spider mite ...
     [INFO] Loading images: Potato healthy ...
     [INFO] Processing Potato healthy ...
     [INFO] Loading images: Tomato healthy ...
     [INFO] Processing Tomato healthy ...
     [INFO] Loading images: Potato Early blight ...
     [INFO] Processing Potato___Early_blight ...
     [INFO] Loading images: Potato Late blight ...
     [INFO] Processing Potato Late blight ...
     [INFO] Loading images: Tomato Late blight ...
     [INFO] Processing Tomato Late blight ...
     [INFO] Loading images: Tomato Early blight ...
     [INFO] Processing Tomato Early blight ...
     [INFO] Loading images: Tomato Bacterial spot ...
     [INFO] Processing Tomato Bacterial spot ...
     [INFO] Loading images: Pepper bell healthy ...
     [INFO] Processing Pepper bell healthy ...
     [INFO] Loading images: Tomato Septoria leaf spot ...
     [INFO] Processing Tomato Septoria leaf spot ...
     [INFO] Loading images: Tomato Tomato mosaic virus ...
     [INFO] Processing Tomato Tomato mosaic virus ...
     [INFO] Loading images: Tomato__Tomato_YellowLeaf__Curl_Virus ...
     [INFO] Processing Tomato Tomato YellowLeaf Curl Virus ...
     [INFO] Loading images: Tomato Leaf Mold ...
     [INFO] Processing Tomato_Leaf_Mold ...
     [INFO] Loading images: Tomato Target Spot ...
     [INFO] Processing Tomato Target Spot ...
     [INFO] Loading images: Pepper__bell___Bacterial_spot ...
     [INFO] Processing Pepper bell Bacterial spot ...
     [INFO] Image loading completed
Get Size of Processed Image
image size = len(image list)
```

```
image_size = len(image_list
image_size
```

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Transform Image Labels uisng Scikit Learn's LabelBinarizer

```
label binarizer = LabelBinarizer()
image labels = label binarizer.fit transform(label list)
pickle.dump(label_binarizer,open('label_transform.pkl', 'wb'))
n classes = len(label binarizer.classes )
Print the classes
print(label binarizer.classes )
     ['Pepper bell Bacterial spot' 'Pepper bell healthy'
      'Potato___Early_blight' 'Potato___Late_blight' 'Potato___healthy'
      'Tomato_Bacterial_spot' 'Tomato_Early_blight' 'Tomato_Late_blight'
      'Tomato_Leaf_Mold' 'Tomato_Septoria_leaf_spot'
      'Tomato Spider mites Two spotted spider mite' 'Tomato Target Spot'
      'Tomato__Tomato_YellowLeaf__Curl_Virus' 'Tomato__Tomato_mosaic_virus'
      'Tomato healthy']
np_image_list = np.array(image_list, dtype=np.float16) / 225.0
print("[INFO] Spliting data to train, test")
x train, x test, y train, y test = train test split(np image list, image labels, test size=0.
     [INFO] Spliting data to train, test
aug = ImageDataGenerator(
   rotation range=25, width shift range=0.1,
   height shift range=0.1, shear range=0.2,
   zoom range=0.2,horizontal flip=True,
    fill mode="nearest")
model = Sequential()
inputShape = (height, width, depth)
chanDim = -1
if K.image data format() == "channels first":
    inputShape = (depth, height, width)
   chanDim = 1
model.add(Conv2D(32, (3, 3), padding="same",input_shape=inputShape))
model.add(Activation("relu"))
model.add(BatchNormalization(axis=chanDim))
model.add(MaxPooling2D(pool size=(3, 3)))
model.add(Dropout(0.25))
model.add(Conv2D(64, (3, 3), padding="same"))
model.add(Activation("relu"))
model.add(BatchNormalization(axis=chanDim))
```

```
model.add(Conv2D(64, (3, 3), padding="same"))
model.add(Activation("relu"))
model.add(BatchNormalization(axis=chanDim))
model.add(MaxPooling2D(pool size=(2, 2)))
model.add(Dropout(0.25))
model.add(Conv2D(128, (3, 3), padding="same"))
model.add(Activation("relu"))
model.add(BatchNormalization(axis=chanDim))
model.add(Conv2D(128, (3, 3), padding="same"))
model.add(Activation("relu"))
model.add(BatchNormalization(axis=chanDim))
model.add(MaxPooling2D(pool_size=(2, 2)))
model.add(Dropout(0.25))
model.add(Flatten())
model.add(Dense(1024))
model.add(Activation("relu"))
model.add(BatchNormalization())
model.add(Dropout(0.5))
model.add(Dense(n classes))
model.add(Activation("softmax"))
```

Model Summary

model.summary()

```
dropout (Dropout)
                            (None, 85, 85, 32)
                                                       0
conv2d_1 (Conv2D)
                            (None, 85, 85, 64)
                                                       18496
activation_1 (Activation)
                            (None, 85, 85, 64)
                                                       0
batch normalization 1 (Batc (None, 85, 85, 64)
                                                       256
hNormalization)
conv2d 2 (Conv2D)
                            (None, 85, 85, 64)
                                                       36928
activation 2 (Activation)
                            (None, 85, 85, 64)
                                                       0
```

```
(None, 42, 42, 128)
     conv2d 4 (Conv2D)
                                                      147584
     activation 4 (Activation)
                              (None, 42, 42, 128)
     batch normalization 4 (Batc (None, 42, 42, 128)
                                                      512
     hNormalization)
     max pooling2d 2 (MaxPooling (None, 21, 21, 128)
                                                      0
     2D)
     dropout 2 (Dropout)
                              (None, 21, 21, 128)
                                                      0
     flatten (Flatten)
                              (None, 56448)
     dense (Dense)
                              (None, 1024)
                                                      57803776
                              (None, 1024)
     activation 5 (Activation)
     batch normalization 5 (Batc (None, 1024)
                                                      4096
     hNormalization)
                              (None, 1024)
     dropout 3 (Dropout)
     dense 1 (Dense)
                              (None, 15)
                                                      15375
     activation 6 (Activation)
                              (None, 15)
    ______
    Total params: 58,102,671
    Trainable params: 58,099,791
opt = Adam(lr=INIT LR, decay=INIT LR / EPOCHS)
# distribution
model.compile(loss="binary crossentropy", optimizer=opt,metrics=["accuracy"])
# train the network
print("[INFO] training network...")
    [INFO] training network...
    /usr/local/lib/python3.9/dist-packages/keras/optimizers/legacy/adam.py:117: UserWarning
      super(). init (name, **kwargs)
history = model.fit generator(
   aug.flow(x_train, y_train, batch_size=BS),
   validation_data=(x_test, y_test),
   steps per epoch=len(x train) // BS,
   epochs=EPOCHS, verbose=1
   )
    <ipython-input-27-f5a41846f0f2>:1: UserWarning: `Model.fit_generator` is deprecated and
      history = model.fit generator(
    Epoch 1/25
```

```
Epoch 2/25
73/73 [============= ] - 39s 526ms/step - loss: 0.2858 - accuracy: 0.296
Epoch 3/25
73/73 [============= ] - 40s 550ms/step - loss: 0.1873 - accuracy: 0.516
Epoch 4/25
Epoch 5/25
Epoch 6/25
73/73 [============= ] - 39s 530ms/step - loss: 0.1171 - accuracy: 0.699
Epoch 7/25
73/73 [============= ] - 39s 524ms/step - loss: 0.1121 - accuracy: 0.718
Epoch 8/25
Epoch 9/25
73/73 [============ - 40s 547ms/step - loss: 0.1201 - accuracy: 0.702
Epoch 10/25
73/73 [============ ] - 40s 550ms/step - loss: 0.0965 - accuracy: 0.754
Epoch 11/25
73/73 [============= ] - 39s 527ms/step - loss: 0.0872 - accuracy: 0.781
Epoch 12/25
73/73 [============= ] - 38s 515ms/step - loss: 0.0970 - accuracy: 0.756
Epoch 13/25
73/73 [============ ] - 39s 530ms/step - loss: 0.0909 - accuracy: 0.77!
Epoch 14/25
73/73 [============ ] - 39s 540ms/step - loss: 0.0815 - accuracy: 0.799
Epoch 15/25
73/73 [============ ] - 39s 536ms/step - loss: 0.0771 - accuracy: 0.806
Epoch 16/25
Epoch 17/25
Epoch 18/25
73/73 [============= ] - 39s 533ms/step - loss: 0.0757 - accuracy: 0.816
Epoch 19/25
73/73 [============= ] - 39s 536ms/step - loss: 0.0655 - accuracy: 0.843
Epoch 20/25
73/73 [=========== ] - 38s 519ms/step - loss: 0.0637 - accuracy: 0.856
Epoch 21/25
73/73 [============= ] - 39s 534ms/step - loss: 0.0591 - accuracy: 0.86
Epoch 22/25
73/73 [============ ] - 39s 539ms/step - loss: 0.0566 - accuracy: 0.869
Epoch 23/25
73/73 [============= ] - 39s 536ms/step - loss: 0.0536 - accuracy: 0.87!
Epoch 24/25
Epoch 25/25
73/73 [============= ] - 37s 512ms/step - loss: 0.0489 - accuracy: 0.889
```

Plot the train and val curve

```
acc = history.history['accuracy']
val_acc = history.history['val_accuracy']
```

```
loss = history.history['loss']
val_loss = history.history['val_loss']
epochs = range(1, len(acc) + 1)
#Train and validation accuracy
plt.plot(epochs, acc, 'b', label='Training accurarcy')
plt.plot(epochs, val_acc, 'r', label='Validation accurarcy')
plt.title('Training and Validation accurarcy')
plt.legend()

plt.figure()
#Train and validation loss
plt.plot(epochs, loss, 'b', label='Training loss')
plt.plot(epochs, val_loss, 'r', label='Validation loss')
plt.title('Training and Validation loss')
plt.legend()
plt.show()
```

Training and Validation accurarcy 0.9 Training accurarcy Validation accurarcy 0.8 0.7 0.6 0.5 Model Accuracy I print("[INFO] Calculating model accuracy") scores = model.evaluate(x_test, y_test) print(f"Test Accuracy: {scores[1]*100}") [INFO] Calculating model accuracy 19/19 [==========] - 1s 36ms/step - loss: 0.4226 - accuracy: 0.7394 Test Accuracy: 73.94247055053711 Save model using Pickle iraining ioss print("[INFO] Saving model...") #file name='cnn model.pkl' pickle.dump(model,open('cnn_model.pkl', 'wb')) model.save('C:\\Users\\en0216\\Downloads\\cnn model.pkl') print("saved") [INFO] Saving model... WARNING:absl:Found untraced functions such as jit compiled convolution op, jit compile saved ١ image dir="/content/PlantVillage/Potato Early blight/0a8a68ee-f587-4dea-beec-79d02e7d3fa4 #image dir="C:\\Users\\SHARATHVEER REDDY K\\Desktop\archive\\PlantVillage\\PlantVillage\\plan im=convert_image_to_array(image_dir) #print(im) np_image_li = np.array(im, dtype=np.float16) / 225.0 npp image = np.expand dims(np image li, axis=0)

X