```
import numpy as np
import pickle
import cv2
import os
import matplotlib.pyplot as plt
from os import listdir
from sklearn.preprocessing import LabelBinarizer
from keras.models import Sequential
from keras.layers import BatchNormalization, Conv2D, MaxPooling2D, Flatten, Dense, Dropout
from keras.optimizers import Adam
from \ keras.preprocessing.image \ import \ ImageDataGenerator, \ img\_to\_array
from sklearn.model_selection import train_test_split
import kagglehub
# Download latest version
path = kagglehub.dataset download("emmarex/plantdisease")
print("Path to dataset files:", path)
🕁 Warning: Looks like you're using an outdated `kagglehub` version, please consider updating (latest version: 0.3.4)
     Path to dataset files: /root/.cache/kagglehub/datasets/emmarex/plantdisease/versions/1
# Dimension of resized image
DEFAULT_IMAGE_SIZE = tuple((256, 256))
# Number of images used to train the model
N_IMAGES = 100
# Path to the dataset folder
train_dir = os.path.join(path, 'train')
val_dir = os.path.join(path, 'val')
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)
           return np.array([])
    except Exception as e:
       print(f"Error : {e}")
        return None
image_list, label_list = [], []
try:
    print("[INFO] Loading images ...")
    plant_disease_folder_list = listdir(train_dir)
    for plant_disease_folder in plant_disease_folder_list:
        print(f"[INFO] Processing {plant_disease_folder} ...")
        plant_disease_image_list = listdir(f"{train_dir}/{plant_disease_folder}/")
        for image in plant_disease_image_list[:N_IMAGES]:
            image_directory = f"{train_dir}/{plant_disease_folder}/{image}"
            if image_directory.endswith(".jpg")==True or image_directory.endswith(".JPG")==True:
                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}")
# Transform the loaded training image data into numpy array
np_image_list = np.array(image_list, dtype=np.float16) / 225.0
print()
# Check the number of images loaded for training
image_len = len(image_list)
print(f"Total number of images: {image_len}")
→ [INFO] Loading images ...
     [INFO] Processing Soybean___healthy ...
     [INFO] Processing Cherry_(including_sour)___Powdery_mildew ...
     [INFO] Processing Corn_(maize)___Northern_Leaf_Blight ...
     [INFO] Processing Tomato__healthy ...
[INFO] Processing Tomato__Target_Spot ...
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[INFO] Processing Potato___Early_blight ..
     [INFO] Processing Pepper,_bell___Bacterial_spot ...
     [INFO] Processing Peach___healthy ...
     [INFO] Processing Corn_(maize)___Cercospora_leaf_spot Gray_leaf_spot ...
     [INFO] Processing Corn_(maize)___healthy ..
     [INFO] Processing Cherry_(including_sour)___healthy ...
     [INFO] Processing Tomato__Leaf_Mold ...
[INFO] Processing Apple__Cedar_apple_rust ...
     [INFO] Processing Potato__healthy \dots
     [INFO] Processing Squash___Powdery_mildew ...
     [INFO] Processing Strawberry___healthy ..
     [INFO] Processing Orange__Haunglongbing_(Citrus_greening) \dots
     [INFO] Processing Apple__healthy ...
     [INFO] Processing Grape___Leaf_blight_(Isariopsis_Leaf_Spot) ...
     [INFO] Processing Strawberry__Leaf_scorch ...
     [INFO] Processing Tomato__Spider_mites Two-spotted_spider_mite ...
[INFO] Processing Tomato__Tomato_Yellow_Leaf_Curl_Virus ...
     [INFO] Processing Pepper,_bell__healthy ...
     [INFO] Processing Potato___Late_blight ...
     [INFO] Processing background ...
     [INFO] Processing Raspberry___healthy ...
     [INFO] Processing Corn_(maize)___Common_rust_ ...
     [INFO] Processing Apple___Black_rot ...
     [INFO] Processing Peach___Bacterial_spot ...
     [INFO] Processing Grape__healthy ...
[INFO] Processing Tomato__Septoria_leaf_spot ...
     [INFO] Processing Apple__Apple_scab ...
     [INFO] Processing Blueberry__healthy ...
     [INFO] Processing Grape__Black_rot ...
[INFO] Processing Tomato__Bacterial_spot ...
     [INFO] Processing Grape___Esca_(Black_Measles) \dots
     [INFO] Processing Tomato___Late_blight ...
     [INFO] Processing Tomato___Tomato_mosaic_virus ...
[INFO] Processing Tomato___Early_blight ...
     [INFO] Image loading completed
     Total number of images: 3900
label_binarizer = LabelBinarizer()
image_labels = label_binarizer.fit_transform(label_list)
pickle.dump(label_binarizer,open('plant_disease_label_transform.pkl', 'wb'))
n_classes = len(label_binarizer.classes_)
print("Total number of classes: ", n_classes)
→ Total number of classes: 39
augment = 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")
print("[INFO] Splitting data to train and test...")
x_train, x_test, y_train, y_test = train_test_split(np_image_list, image_labels, test_size=0.2, random_state = 42)

→ [INFO] Splitting data to train and test...
EPOCHS = 25
STEPS = 100
LR = 1e-3
BATCH_SIZE = 32
WIDTH = 256
HEIGHT = 256
DEPTH = 3
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))
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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"))
{\tt 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: "sequential\_1"

Model: Sequential_1		
Layer (type)	Output Shape	Param #
conv2d_1 (Conv2D)	(None, 256, 256, 32)	896
activation_1 (Activation)	(None, 256, 256, 32)	0
batch_normalization_1 (Batch	(None, 256, 256, 32)	128
max_pooling2d_1 (MaxPooling2	(None, 85, 85, 32)	0
dropout_1 (Dropout)	(None, 85, 85, 32)	0
conv2d_2 (Conv2D)	(None, 85, 85, 64)	18496
activation_2 (Activation)	(None, 85, 85, 64)	0
batch_normalization_2 (Batch	(None, 85, 85, 64)	256
conv2d_3 (Conv2D)	(None, 85, 85, 64)	36928
activation_3 (Activation)	(None, 85, 85, 64)	0
batch_normalization_3 (Batch	(None, 85, 85, 64)	256
max_pooling2d_2 (MaxPooling2	(None, 42, 42, 64)	0
dropout_2 (Dropout)	(None, 42, 42, 64)	0
conv2d_4 (Conv2D)	(None, 42, 42, 128)	73856
activation_4 (Activation)	(None, 42, 42, 128)	0
batch_normalization_4 (Batch	(None, 42, 42, 128)	512
conv2d_5 (Conv2D)	(None, 42, 42, 128)	147584
activation_5 (Activation)	(None, 42, 42, 128)	0
batch_normalization_5 (Batch	(None, 42, 42, 128)	512
max_pooling2d_3 (MaxPooling2	(None, 21, 21, 128)	0
dropout_3 (Dropout)	(None, 21, 21, 128)	0
flatten_1 (Flatten)	(None, 56448)	0
dense_1 (Dense)	(None, 1024)	57803776
activation_6 (Activation)	(None, 1024)	0
batch_normalization_6 (Batch	(None, 1024)	4096
dropout_4 (Dropout)	(None, 1024)	0
dense_2 (Dense)	(None, 39)	39975

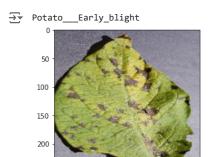
```
# Initialize optimizer
opt = Adam(lr=LR, decay=LR / EPOCHS)

# Compile model
model.compile(loss="binary_crossentropy", optimizer=opt, metrics=["accuracy"])
```

```
# Train model
print("[INFO] Training network...")
history = model.fit_generator(augment.flow(x_train, y_train, batch_size=BATCH_SIZE),
                 validation_data=(x_test, y_test),
                 steps_per_epoch=len(x_train) // BATCH_SIZE,
                 epochs=EPOCHS,
                 verbose=1)
→ [INFO] Training network...
   Epoch 1/25
   97/97 [===:
            Epoch 2/25
   97/97 [======
            Epoch 3/25
   97/97 [=====
          =============================  - 793s 8s/step - loss: 0.0718 - accuracy: 0.9775 - val_loss: 0.3322 - val_accuracy: 0.9549
   Epoch 4/25
   97/97 [=====
            Epoch 5/25
   97/97 [=========] - 791s 8s/step - loss: 0.0767 - accuracy: 0.9809 - val loss: 0.0760 - val accuracy: 0.9767
   Epoch 6/25
   Epoch 7/25
   97/97 [============= - 795s 8s/step - loss: 0.0520 - accuracy: 0.9828 - val loss: 0.1049 - val accuracy: 0.9765
   Epoch 8/25
   97/97 [=====
            ============================== - 802s 8s/step - loss: 0.0422 - accuracy: 0.9854 - val_loss: 0.0811 - val_accuracy: 0.9787
   Epoch 9/25
   97/97 [============ - 796s 8s/step - loss: 0.0395 - accuracy: 0.9860 - val loss: 0.0859 - val accuracy: 0.9787
   Epoch 10/25
   97/97 [=====
                :========] - 804s 8s/step - loss: 0.0359 - accuracy: 0.9875 - val_loss: 0.2368 - val_accuracy: 0.9668
   Epoch 11/25
   Epoch 12/25
   Epoch 13/25
   97/97 [==========] - 806s 8s/step - loss: 0.0328 - accuracy: 0.9883 - val loss: 0.2544 - val accuracy: 0.9608
   Epoch 14/25
   97/97 [============ - 804s 8s/step - loss: 0.0373 - accuracy: 0.9868 - val loss: 0.0958 - val accuracy: 0.9783
   Epoch 15/25
   97/97 [====
               :========] - 810s 8s/step - loss: 0.0305 - accuracy: 0.9892 - val_loss: 0.1213 - val_accuracy: 0.9722
   Epoch 16/25
   Epoch 17/25
   97/97 [=====
             Epoch 18/25
   Epoch 19/25
   Epoch 20/25
   Epoch 21/25
   97/97 [=====
              Epoch 22/25
   97/97 [====
            Epoch 23/25
   97/97 [==========] - 825s 9s/step - loss: 0.0203 - accuracy: 0.9925 - val loss: 0.0264 - val accuracy: 0.9916
   Enoch 24/25
   97/97 [=====
           Epoch 25/25
   97/97 [=========] - 813s 8s/step - loss: 0.0187 - accuracy: 0.9933 - val loss: 0.0497 - val accuracy: 0.9875
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()

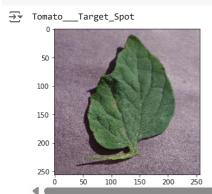
 $\overline{\Rightarrow}$ Training and Validation accurarcy 0.99 0.98 0.97 0.96 Training accurarcy Validation accurarcy 0.95 20 Training and Validation loss 0.35 Training loss Validation loss 0.30 0.25 0.20 0.15 0.10 0.05 print("[INFO] Calculating model accuracy") scores = model.evaluate(x\_test, y\_test) print(f"Test Accuracy: {scores[1]\*100}") → [INFO] Calculating model accuracy 780/780 [========== ] - 52s 66ms/step Test Accuracy: 98.74754548072815 # Dump pickle file of the model print("[INFO] Saving model...") pickle.dump(model,open('plant\_disease\_classification\_model.pkl', 'wb'))  $\rightarrow$  [INFO] Saving model... # Dump pickle file of the labels print("[INFO] Saving label transform...") filename = 'plant\_disease\_label\_transform.pkl' image\_labels = pickle.load(open(filename, 'rb')) → [INFO] Saving label transform... def predict\_disease(image\_path): image\_array = convert\_image\_to\_array(image\_path) np\_image = np.array(image\_array, dtype=np.float16) / 225.0 np\_image = np.expand\_dims(np\_image,0) plt.imshow(plt.imread(image\_path)) result = model.predict\_classes(np\_image) print((image\_labels.classes\_[result][0]))  $predict\_disease('/content/PlantVillage/val/Blueberry\_\_healthy/008c85d0-a954-4127-bd26-861dc8a1e6ff\_\_RS\_HL\ 2431.JPG')$ → Blueberry healthy 50 100 200 250 100

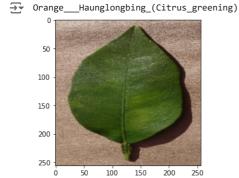


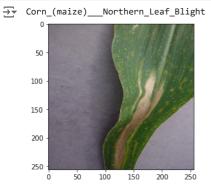
100

250

 $predict\_disease('/content/PlantVillage/val/Tomato\_\_Target\_Spot/1006b3dd-22d8-41b8-b83d-08bf189fcdaa\_\_Com.G\_TgS\_FL~8118.JPG')$ 







Start coding or  $\underline{\text{generate}}$  with AI.