Leaf Disease Detection using Alexnet

```
import os
os.listdir("../input/plant-diseases-classification-using-alexnet")

['__results__.html',
    'best_weights_9.hdf5',
    'AlexNetModel.hdf5',
    '__results___files',
    '__output__.json',
    'custom.css']
```

Tensorflow is a foundation library that can be used to create deep learning models directly or by using wrapper libraries that simplify the process built on the top of tensorflow so for making deep learning model we are installing tensorflow. TensorFlow is mainly used for: Classification, Perception, Understanding, Discovering, Prediction and Creation.

```
# ! pip install tensorflow-gpu
```

from tensorflow we are importing device_lib that will provide us the full summary or list of information about local device like computational capability, memory limit, device type etc.

```
from tensorflow.python.client import device_lib
device_lib.list_local_devices()
     [name: "/device:CPU:0"
      device_type: "CPU"
      memory_limit: 268435456
      locality {
      }
      incarnation: 12930137146812964960, name: "/device:GPU:0"
      device_type: "GPU"
      memory_limit: 15900252570
      locality {
        bus_id: 1
        links {
        }
      }
      incarnation: 17279202340371761108
      physical_device_desc: "device: 0, name: Tesla P100-PCIE-16GB, pci bus id: 0000:00:04
```

Keras is a powerful and easy-to-use free open source Python library for developing and evaluating deep learning models. It wraps the efficient numerical computation libraries and allows you to define and train neural network models in just a few lines of code.

```
import os
import tensorflow as tf
from tensorflow import keras
```

!nvidia-smi

Building CNN Based On AlexNet Architecture

Here we are making the sequential model for that we are importing sequential from tensorflow. A Sequential model is appropriate for a plain stack of layers where each layer has exactly one input tensor and one output tensor. By the use of Convolution2D it will be summing up the results into a single output pixel. Maximum pooling, or max pooling, is a pooling operation that calculates the maximum, or largest, value in each patch of each feature map. Flatten can converts the pooled feature map to a single column that is passed to the fully connected layer. Dense adds the fully connected layer to the neural network. The Dropout layer randomly sets input units to 0 with a frequency of rate at each step during training time, which helps prevent overfitting. And hence Batch normalization accelerates training, in some cases by halving the epochs or better, and provides some regularization, reducing generalization error.

```
# Importing Keras libraries and packages
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Convolution2D
from tensorflow.keras.layers import MaxPooling2D
from tensorflow.keras.layers import Flatten
from tensorflow.keras.layers import Dense
from tensorflow.keras.layers import Dropout
from tensorflow.keras.layers import BatchNormalization
with tf.device('/device:GPU:0'):
    # Initializing the CNN
    classifier = Sequential()
    # Convolution Step 1
    classifier.add(Convolution2D(96, 11, strides = (4, 4), padding = 'valid', input_shape=
    # Max Pooling Step 1
    classifier.add(MaxPooling2D(pool_size = (2, 2), strides = (2, 2), padding = 'valid'))
    classifier.add(BatchNormalization())
    # Convolution Step 2
    classifier.add(Convolution2D(256, 11, strides = (1, 1), padding='valid', activation =
    # Max Pooling Step 2
    classifier.add(MaxPooling2D(pool size = (2. 2). strides = (2. 2). padding='valid'))
```

```
(-, -/, -----
classifier.add(BatchNormalization())
# Convolution Step 3
classifier.add(Convolution2D(384, 3, strides = (1, 1), padding='valid', activation = '
classifier.add(BatchNormalization())
# Convolution Step 4
classifier.add(Convolution2D(384, 3, strides = (1, 1), padding='valid', activation = '
classifier.add(BatchNormalization())
# Convolution Step 5
classifier.add(Convolution2D(256, 3, strides=(1,1), padding='valid', activation = 'rel
# Max Pooling Step 3
classifier.add(MaxPooling2D(pool_size = (2, 2), strides = (2, 2), padding = 'valid'))
classifier.add(BatchNormalization())
# Flattening Step
classifier.add(Flatten())
# Full Connection Step
classifier.add(Dense(units = 4096, activation = 'relu'))
classifier.add(Dropout(0.4))
classifier.add(BatchNormalization())
classifier.add(Dense(units = 4096, activation = 'relu'))
classifier.add(Dropout(0.4))
classifier.add(BatchNormalization())
classifier.add(Dense(units = 1000, activation = 'relu'))
classifier.add(Dropout(0.2))
classifier.add(BatchNormalization())
classifier.add(Dense(units = 38, activation = 'softmax'))
classifier.summary()
```

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 54, 54, 96)	34944
max_pooling2d (MaxPooling2D)	(None, 27, 27, 96)	0
batch_normalization (BatchNo	(None, 27, 27, 96)	384
conv2d_1 (Conv2D)	(None, 17, 17, 256)	2973952
max_pooling2d_1 (MaxPooling2	(None, 8, 8, 256)	0
batch_normalization_1 (Batch	(None, 8, 8, 256)	1024
conv2d_2 (Conv2D)	(None, 6, 6, 384)	885120
batch_normalization_2 (Batch	(None, 6, 6, 384)	1536
conv2d_3 (Conv2D)	(None, 4, 4, 384)	1327488
batch_normalization_3 (Batch	(None, 4, 4, 384)	1536
conv2d_4 (Conv2D)	(None, 2, 2, 256)	884992

max_pooling2d_2 (MaxPooling2	(None,	1, 1, 256)	0
batch_normalization_4 (Batch	(None,	1, 1, 256)	1024
flatten (Flatten)	(None,	256)	0
dense (Dense)	(None,	4096)	1052672
dropout (Dropout)	(None,	4096)	0
batch_normalization_5 (Batch	(None,	4096)	16384
dense_1 (Dense)	(None,	4096)	16781312
dropout_1 (Dropout)	(None,	4096)	0
batch_normalization_6 (Batch	(None,	4096)	16384
dense_2 (Dense)	(None,	1000)	4097000
dropout_2 (Dropout)	(None,	1000)	0
batch_normalization_7 (Batch	(None,	1000)	4000
dense_3 (Dense)	(None,	38)	38038
Total nanams: 29 117 700		·	

Total params: 28,117,790 Trainable params: 28,096,654 Non-trainable params: 21,136

Loading Weights To The Model

classifier.load_weights('../input/plant-diseases-classification-using-alexnet/best_weights

Fine Tuning By Freezing Some Layers Of Our Model

let's visualize layer names and layer indices to see how many layers we should freeze:

```
from tensorflow.keras import layers
for i, layer in enumerate(classifier.layers):
    print(i, layer.name)

    0 conv2d
    1 max_pooling2d
    2 batch_normalization
    3 conv2d_1
    4 max_pooling2d_1
    5 batch_normalization_1
    6 conv2d_2
```

- 7 batch_normalization_2
- 8 conv2d_3
- 9 batch_normalization_3
- 10 conv2d_4

```
11 max_pooling2d_2
12 batch_normalization_4
13 flatten
14 dense
15 dropout
16 batch_normalization_5
17 dense_1
18 dropout_1
19 batch_normalization_6
20 dense_2
21 dropout_2
22 batch_normalization_7
23 dense_3
```

we chose to train the top 2 conv blocks, i.e. we will freeze the first 8 layers and unfreeze the rest:

```
print("Freezed layers:")
for i, layer in enumerate(classifier.layers[:20]):
   print(i, layer.name)
   layer.trainable = False
     Freezed layers:
     0 conv2d
     1 max_pooling2d
     2 batch normalization
     3 conv2d 1
     4 max pooling2d 1
     5 batch_normalization_1
     6 conv2d 2
     7 batch_normalization_2
     8 conv2d 3
     9 batch_normalization_3
     10 conv2d 4
     11 max_pooling2d_2
     12 batch_normalization_4
     13 flatten
     14 dense
     15 dropout
     16 batch_normalization_5
     17 dense 1
     18 dropout 1
     19 batch_normalization_6
```

Model Summary After Freezing

trainable parameters decrease after freezing some bottom layers.

classifier.summary()

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 54, 54, 96)	34944

plant_diseases_classification_using_alexnet_(2) (1).ipynb - Colaborate				
<pre>max_pooling2d (MaxPooling2D)</pre>	(None,	27, 27, 96)	0	
batch_normalization (BatchNo	(None,	27, 27, 96)	384	
conv2d_1 (Conv2D)	(None,	17, 17, 256)	2973952	
max_pooling2d_1 (MaxPooling2	(None,	8, 8, 256)	0	
batch_normalization_1 (Batch	(None,	8, 8, 256)	1024	
conv2d_2 (Conv2D)	(None,	6, 6, 384)	885120	
batch_normalization_2 (Batch	(None,	6, 6, 384)	1536	
conv2d_3 (Conv2D)	(None,	4, 4, 384)	1327488	
batch_normalization_3 (Batch	(None,	4, 4, 384)	1536	
conv2d_4 (Conv2D)	(None,	2, 2, 256)	884992	
max_pooling2d_2 (MaxPooling2	(None,	1, 1, 256)	0	
batch_normalization_4 (Batch	(None,	1, 1, 256)	1024	
flatten (Flatten)	(None,	256)	0	
dense (Dense)	(None,	4096)	1052672	
dropout (Dropout)	(None,	4096)	0	
batch_normalization_5 (Batch	(None,	4096)	16384	
dense_1 (Dense)	(None,	4096)	16781312	
dropout_1 (Dropout)	(None,	4096)	0	
batch_normalization_6 (Batch	(None,	4096)	16384	
dense_2 (Dense)	(None,	1000)	4097000	
dropout_2 (Dropout)	(None,	1000)	0	
batch_normalization_7 (Batch	(None,	1000)	4000	
dense_3 (Dense)	(None,	38)	38038	

Total params: 28,117,790
Trainable params: 4,137,038
Non-trainable params: 23,080,75

Non-trainable params: 23,980,752

Compiling the Model

Image Preprocessing

```
# image preprocessing
from tensorflow.keras.preprocessing.image import ImageDataGenerator
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
base dir = "../input/new-plant-diseases-dataset/new plant diseases dataset(augmented)/New
training_set = train_datagen.flow_from_directory(base_dir+'/train',
                                                                                                                                         target_size=(224, 224),
                                                                                                                                         batch size=batch size,
                                                                                                                                         class mode='categorical')
valid_set = valid_datagen.flow_from_directory(base_dir+'/valid',
                                                                                                                           target_size=(224, 224),
                                                                                                                           batch_size=batch_size,
                                                                                                                           class mode='categorical')
              Found 70295 images belonging to 38 classes.
              Found 17572 images belonging to 38 classes.
 Printing all the classes of training data set with index value.
class_dict = training_set.class_indices
print(class dict)
              {'Apple__Apple_scab': 0, 'Apple__Black_rot': 1, 'Apple__Cedar_apple_rust': 2, 'Apple_Cedar_apple_rust': 2, 'Apple_rust': 2, 'Apple_
listing all the classes without index
li = list(class_dict.keys())
print(li)
              ['Apple Apple scab', 'Apple Black rot', 'Apple Cedar apple rust', 'Apple heal
```

Assigning all the training set and valid set samples to the variable

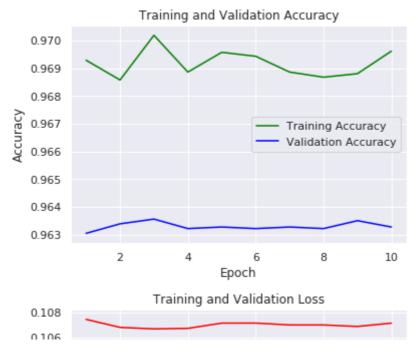
```
train_num = training_set.samples
valid num = valid set.samples
```

Importing checkpoint for checking the accuracy of each epoch to avoid the overfitting problem.

```
# checkpoint
from tensorflow.keras.callbacks import ModelCheckpoint
weightpath = "./best weights 9.hdf5"
checkpoint = ModelCheckpoint(weightpath, monitor='val_acc', verbose=1, save_best_only=True
callbacks_list = [checkpoint]
#fitting images to CNN
history = classifier.fit_generator(training_set,
                steps_per_epoch=train_num//batch_size,
                validation_data=valid_set,
                epochs=10,
                validation_steps=valid_num//batch_size,
                callbacks=callbacks list)
#saving model
filepath="AlexNetModel.hdf5"
classifier.save(filepath)
   Epoch 1/10
   Epoch 00001: val_acc improved from -inf to 0.96305, saving model to ./best_weights_9
   Epoch 2/10
   Epoch 00002: val_acc improved from 0.96305 to 0.96339, saving model to ./best_weights
   549/549 [================== ] - 821s 1s/step - loss: 0.0933 - acc: 0.9685
   Epoch 3/10
   Epoch 00003: val acc improved from 0.96339 to 0.96356, saving model to ./best weights
   Epoch 4/10
   Epoch 00004: val_acc did not improve from 0.96356
   549/549 [============= ] - 832s 2s/step - loss: 0.0930 - acc: 0.9687
   Epoch 5/10
   Epoch 00005: val acc did not improve from 0.96356
   549/549 [=============] - 832s 2s/step - loss: 0.0916 - acc: 0.9696
   Epoch 6/10
   Epoch 00006: val acc did not improve from 0.96356
   549/549 [============= ] - 831s 2s/step - loss: 0.0918 - acc: 0.9693
   Epoch 7/10
   Epoch 00007: val_acc did not improve from 0.96356
   549/549 [================ ] - 804s 1s/step - loss: 0.0938 - acc: 0.9688
   Epoch 8/10
   Epoch 00008: val acc did not improve from 0.96356
   549/549 [============== ] - 851s 2s/step - loss: 0.0932 - acc: 0.9686
   Epoch 9/10
```

Visualising Training Progress

```
#plotting training values
import matplotlib.pyplot as plt
import seaborn as sns
sns.set()
acc = history.history['acc']
val_acc = history.history['val_acc']
loss = history.history['loss']
val_loss = history.history['val_loss']
epochs = range(1, len(loss) + 1)
#accuracy plot
plt.plot(epochs, acc, color='green', label='Training Accuracy')
plt.plot(epochs, val_acc, color='blue', label='Validation Accuracy')
plt.title('Training and Validation Accuracy')
plt.ylabel('Accuracy')
plt.xlabel('Epoch')
plt.legend()
plt.figure()
#loss plot
plt.plot(epochs, loss, color='pink', label='Training Loss')
plt.plot(epochs, val_loss, color='red', label='Validation Loss')
plt.title('Training and Validation Loss')
plt.xlabel('Epoch')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



Predicting New Test Image(s)

```
0.102
# predicting an image
from tensorflow.keras.preprocessing import image
import numpy as np
image_path = "../input/new-plant-diseases-dataset/test/test/TomatoEarlyBlight1.JPG"
new img = image.load img(image path, target size=(224, 224))
img = image.img_to_array(new_img)
img = np.expand_dims(img, axis=0)
img = img/255
print("Following is our prediction:")
prediction = classifier.predict(img)
# decode the results into a list of tuples (class, description, probability)
# (one such list for each sample in the batch)
d = prediction.flatten()
j = d.max()
for index,item in enumerate(d):
    if item == j:
        class name = li[index]
##Another way
# img_class = classifier.predict_classes(img)
# img prob = classifier.predict proba(img)
# print(img_class ,img_prob )
#ploting image with predicted class name
plt.figure(figsize = (4,4))
plt.imshow(new img)
plt.axis('off')
plt.title(class_name)
plt.show()
```

Following is our prediction:

Tomato___Early_blight



new_model = tf.keras.models.load_model('./AlexNetModel.hdf5')

```
# predicting an image
from keras.preprocessing import image
import numpy as np
image_path = "../input/new-plant-diseases-dataset/test/test/AppleCedarRust2.JPG"
new_img = image.load_img(image_path, target_size=(224, 224))
img = image.img_to_array(new_img)
img = np.expand_dims(img, axis=0)
img = img/255
print("Following is our prediction:")
prediction = new_model.predict(img)
# decode the results into a list of tuples (class, description, probability)
# (one such list for each sample in the batch)
d = prediction.flatten()
j = d.max()
for index,item in enumerate(d):
    if item == j:
        class name = li[index]
##Another way
# img class = classifier.predict classes(img)
# img prob = classifier.predict proba(img)
# print(img class ,img prob )
#ploting image with predicted class name
plt.figure(figsize = (4,4))
plt.imshow(new img)
plt.axis('off')
plt.title(class_name)
plt.show()
```

Using TensorFlow backend. Following is our prediction:

Apple__Cedar_apple_rust

