plantdisease-train

April 30, 2024

1 Plant Disease Prediction

1.1 Importing Dataset

Dataset Link: https://www.kaggle.com/datasets/vipoooool/new-plant-diseases-dataset

1.2 Importing libraries

```
import tensorflow as tf
from tensorflow.keras import models, utils, optimizers
from tensorflow.keras.layers import Conv2D, MaxPool2D, Dropout, Flatten, Dense
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
import os
import json
from zipfile import ZipFile
from PIL import Image
from IPython.display import HTML
```

```
Downloading new-plant-diseases-dataset.zip to /content 100% 2.69G/2.70G [00:31<00:00, 89.2MB/s] 100% 2.70G/2.70G [00:31<00:00, 91.9MB/s]
```

1.3 Data Preprocessing

1.3.1 Training Image preprocessing

```
[3]: training_set = utils.image_dataset_from_directory(
         'train',
         labels="inferred",
         label_mode="categorical",
         class_names=None,
         color_mode="rgb",
         batch size=32,
         image_size=(128, 128),
         shuffle=True,
         seed=None,
         validation_split=None,
         subset=None,
         interpolation="bilinear",
         follow_links=False,
         crop_to_aspect_ratio=False
     )
```

Found 70295 files belonging to 38 classes.

1.3.2 Validation Image Preprocessing

```
[4]: validation_set = utils.image_dataset_from_directory(
         'valid'.
         labels="inferred",
         label_mode="categorical",
         class_names=None,
         color_mode="rgb",
         batch_size=32,
         image_size=(128, 128),
         shuffle=True,
         seed=None,
         validation_split=None,
         subset=None,
         interpolation="bilinear",
         follow_links=False,
         crop_to_aspect_ratio=False
     )
```

Found 17572 files belonging to 38 classes.

```
[5]: class_names = training_set.class_names class_names
```

```
[5]: ['Apple__Apple_scab',
      'Apple___Black_rot',
      'Apple___Cedar_apple_rust',
      'Apple__healthy',
      'Blueberry healthy',
      'Cherry_(including_sour)___Powdery_mildew',
      'Cherry_(including_sour)__healthy',
      'Corn_(maize)___Cercospora_leaf_spot Gray_leaf_spot',
      'Corn_(maize)___Common_rust_',
      'Corn_(maize)___Northern_Leaf_Blight',
      'Corn_(maize)__healthy',
      'Grape___Black_rot',
      'Grape___Esca_(Black_Measles)',
      'Grape___Leaf_blight_(Isariopsis_Leaf_Spot)',
      'Grape___healthy',
      'Orange___Haunglongbing_(Citrus_greening)',
      'Peach___Bacterial_spot',
      'Peach__healthy',
      'Pepper,_bell___Bacterial_spot',
      'Pepper,_bell__healthy',
      'Potato___Early_blight',
      'Potato___Late_blight',
      'Potato__healthy',
      'Raspberry__healthy',
      'Soybean___healthy',
      'Squash___Powdery_mildew',
      'Strawberry___Leaf_scorch',
      'Strawberry__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_Yellow_Leaf_Curl_Virus',
      'Tomato___Tomato_mosaic_virus',
      'Tomato___healthy']
```

1.4 Building Model

```
cnn.add(Conv2D(filters=64,kernel_size=3,padding='same',activation='relu'))
cnn.add(MaxPool2D(pool_size=2,strides=2))
cnn.add(Conv2D(filters=64,kernel_size=3,padding='same',activation='relu'))
cnn.add(MaxPool2D(pool_size=2,strides=2))
cnn.add(Dropout(0.25))
cnn.add(Conv2D(filters=64,kernel_size=3,padding='same',activation='relu'))
cnn.add(MaxPool2D(pool_size=2,strides=2))
cnn.add(Conv2D(filters=64,kernel_size=3,padding='same',activation='relu'))
cnn.add(MaxPool2D(pool_size=2,strides=2))
cnn.add(MaxPool2D(pool_size=2,strides=2))
cnn.add(Dropout(0.25))
cnn.add(Dropout(0.25))
cnn.add(Dropout(0.4))
cnn.add(Dense(units=38,activation='softmax'))
```

[8]: cnn.compile(optimizer=optimizers.Adam(
 learning_rate=0.0001),loss='categorical_crossentropy',metrics=['accuracy'])

[9]: cnn.summary()

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 128, 128, 32)	896
<pre>max_pooling2d (MaxPooling2 D)</pre>	(None, 64, 64, 32)	0
dropout (Dropout)	(None, 64, 64, 32)	0
conv2d_1 (Conv2D)	(None, 64, 64, 64)	18496
<pre>max_pooling2d_1 (MaxPoolin g2D)</pre>	(None, 32, 32, 64)	0
conv2d_2 (Conv2D)	(None, 32, 32, 64)	36928
<pre>max_pooling2d_2 (MaxPoolin g2D)</pre>	(None, 16, 16, 64)	0

```
dropout_1 (Dropout)
                          (None, 16, 16, 64)
     conv2d_3 (Conv2D)
                           (None, 16, 16, 64)
                                               36928
    max_pooling2d_3 (MaxPoolin (None, 8, 8, 64)
                                               0
     g2D)
     conv2d_4 (Conv2D)
                           (None, 8, 8, 64)
                                               36928
    max_pooling2d_4 (MaxPoolin (None, 4, 4, 64)
                                               0
     g2D)
                                               0
     dropout_2 (Dropout)
                           (None, 4, 4, 64)
                           (None, 1024)
    flatten (Flatten)
     dense (Dense)
                           (None, 1500)
                                               1537500
     dropout_3 (Dropout)
                           (None, 1500)
                                               57038
     dense 1 (Dense)
                           (None, 38)
    ______
    Total params: 1724714 (6.58 MB)
    Trainable params: 1724714 (6.58 MB)
    Non-trainable params: 0 (0.00 Byte)
[10]: training_history = cnn.
     fit(x=training_set,validation_data=validation_set,epochs=10)
    Epoch 1/10
    accuracy: 0.2346 - val_loss: 1.9025 - val_accuracy: 0.4936
    Epoch 2/10
    2197/2197 [============= - - 98s 44ms/step - loss: 1.2859 -
    accuracy: 0.6128 - val_loss: 0.9892 - val_accuracy: 0.7251
    Epoch 3/10
    accuracy: 0.7539 - val_loss: 0.5564 - val_accuracy: 0.8449
    Epoch 4/10
    2197/2197 [============= ] - 87s 39ms/step - loss: 0.5710 -
    accuracy: 0.8209 - val_loss: 0.4645 - val_accuracy: 0.8573
    Epoch 5/10
    accuracy: 0.8610 - val_loss: 0.3923 - val_accuracy: 0.8786
    Epoch 6/10
    2197/2197 [============== ] - 83s 38ms/step - loss: 0.3476 -
```

```
Epoch 7/10
    accuracy: 0.9067 - val_loss: 0.2383 - val_accuracy: 0.9264
    Epoch 8/10
    2197/2197 [============ ] - 79s 36ms/step - loss: 0.2456 -
    accuracy: 0.9186 - val loss: 0.2320 - val accuracy: 0.9267
    Epoch 9/10
    2197/2197 [============= ] - 85s 39ms/step - loss: 0.2130 -
    accuracy: 0.9287 - val_loss: 0.1899 - val_accuracy: 0.9391
    Epoch 10/10
    accuracy: 0.9379 - val_loss: 0.1883 - val_accuracy: 0.9389
    1.5 Evaluating Model
[11]: # Training Accuracy
     train_loss, train_acc = cnn.evaluate(training_set)
     print('Training accuracy:', train_acc)
    accuracy: 0.9660
    Training accuracy: 0.9659719467163086
[12]: # Validation Accuracy
     val_loss, val_acc = cnn.evaluate(validation_set)
     print('Validation accuracy:', val_acc)
    550/550 [============ ] - 12s 22ms/step - loss: 0.1883 -
    accuracy: 0.9389
    Validation accuracy: 0.9389369487762451
    1.5.1 Saving Model
[13]: cnn.save('trained_plant_disease_model.h5')
    /usr/local/lib/python3.10/dist-packages/keras/src/engine/training.py:3103:
    UserWarning: You are saving your model as an HDF5 file via `model.save()`. This
    file format is considered legacy. We recommend using instead the native Keras
    format, e.g. `model.save('my_model.keras')`.
      saving_api.save_model(
[14]: training_history.history
[14]: {'loss': [3.062305212020874,
      1.28593111038208,
      0.7981175780296326,
      0.5709856152534485,
```

accuracy: 0.8865 - val_loss: 0.2675 - val_accuracy: 0.9187

```
0.4321916103363037,
        0.34760773181915283,
        0.28534531593322754,
        0.24562574923038483,
        0.21301718056201935,
        0.18715494871139526],
       'accuracy': [0.2346397340297699,
        0.6127747297286987,
        0.7538800835609436,
        0.8209260702133179,
        0.8609573841094971,
        0.8865352869033813,
        0.906735897064209,
        0.9186002016067505,
        0.9287146925926208,
        0.9379187822341919],
       'val_loss': [1.9025198221206665,
        0.9891754388809204,
        0.556390106678009,
        0.4644668698310852,
        0.39225131273269653,
        0.2675146162509918,
        0.23834316432476044,
        0.2319595366716385,
        0.18992629647254944,
        0.18827539682388306],
       'val_accuracy': [0.4935693144798279,
        0.7250739932060242,
        0.844923734664917,
        0.8572729229927063,
        0.8785567879676819,
        0.9187343716621399,
        0.9263601303100586,
        0.926701545715332,
        0.9390507340431213,
        0.9389369487762451]}
[15]: import json
      with open('training_hist.json','w') as f:
        json.dump(training_history.history,f)
[16]: print(training_history.history.keys())
     dict_keys(['loss', 'accuracy', 'val_loss', 'val_accuracy'])
```

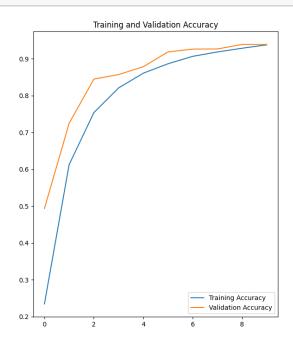
1.6 Visualising results

plt.show()

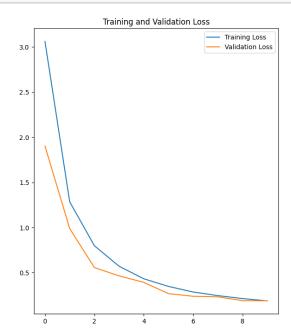
```
[17]: acc = training_history.history['accuracy']
    val_acc = training_history.history['val_accuracy']
    loss = training_history.history['loss']
    val_loss = training_history.history['val_loss']

[18]: plt.figure(figsize=(15, 8))
    plt.subplot(1, 2, 1)
    plt.plot(range(10), acc, label='Training Accuracy')
    plt.plot(range(10), val_acc, label='Validation Accuracy')
    plt.legend(loc='lower right')
    plt.title('Training and Validation Accuracy')

plt.subplot(1, 2, 2)
    plt.plot(range(10), loss, label='Training Loss')
    plt.plot(range(10), val_loss, label='Validation Loss')
    plt.legend(loc='upper right')
```



plt.title('Training and Validation Loss')



1.7 Model evaluation

```
[21]: class_name = validation_set.class_names
[22]: len(class_name)
```

```
[22]: 38
[28]: test_set = utils.image_dataset_from_directory(
          'test',
          labels="inferred",
          label_mode="categorical",
          class names=None,
          color_mode="rgb",
          batch_size=32,
          image_size=(128, 128),
          shuffle=False,
          seed=None,
          validation_split=None,
          subset=None,
          interpolation="bilinear",
          follow_links=False,
          crop_to_aspect_ratio=False
      )
     Found 17572 files belonging to 38 classes.
[30]: # Predicted
      y_pred = cnn.predict(test_set)
      predicted_categories = tf.argmax(y_pred, axis=1)
      predicted_categories
[30]: <tf.Tensor: shape=(17572,), dtype=int64, numpy=array([ 0,  0,  0,  ..., 37, 37,
      37])>
[32]: # Actual
      true_categories = tf.concat([y for x, y in test_set], axis=0)
      Y_true = tf.argmax(true_categories, axis=1)
      Y_true
[32]: <tf.Tensor: shape=(17572,), dtype=int64, numpy=array([ 0, 0, ..., 37, 37,
      37])>
[26]: len(Y_true)
[26]: 33
[27]: len(predicted_categories)
```

[27]: 33

1.8 Precision Recall Fscore

[36]: from sklearn.metrics import confusion_matrix,classification_report

f1-score	support	precision	recall
II SCOLE	support		
	AppleApple_sca	b 0.97	0.91
0.94	504		
0.06	AppleBlack_ro	t 0.94	0.99
0.96	497 AppleCedar_apple_rus	t 0.89	0.99
0.94	440	0.00	0.00
	Applehealth	y 0.91	0.96
0.93	502		
0.05	Blueberryhealth	y 0.96	0.94
0.95	454 Cherry_(including_sour)Powdery_milde	w 1.00	0.96
0.98	421	w 1.00	0.50
	Cherry_(including_sour)health	y 0.95	0.89
0.92	456		
	ze)Cercospora_leaf_spot Gray_leaf_spo	t 0.89	0.91
0.90	410 Corn_(maize)Common_rust	0.99	0.00
0.99	477	_ 0.99	0.99
	Corn_(maize)Northern_Leaf_Bligh	t 0.93	0.96
0.94	477		
	Corn_(maize)health	y 0.97	1.00
0.99	465		0.04
0.94	GrapeBlack_ro	t 0.93	0.94
0.31	GrapeEsca_(Black_Measles) 0.96	0.97
0.97	480		
	rapeLeaf_blight_(Isariopsis_Leaf_Spot	0.97	0.97
0.97	430	0.04	0.00
0.95	Grapehealth	y 0.91	0.99
0.95	OrangeHaunglongbing_(Citrus_greening	0.93	0.98
0.95	503	,	
	PeachBacterial_spo	t 0.91	0.89
0.90	459	2 2-	0.00
0.97	Peachhealth	y 0.95	0.99
0.31	Pepper,_bellBacterial_spo	t 0.94	0.93
		- 0.01	3.20

0.94	478			
0.89	497	Pepper,_bellhealthy	0.96	0.84
0.96	485	PotatoEarly_blight	0.93	0.98
0.90	485	PotatoLate_blight	0.97	0.84
		Potatohealthy	0.95	0.95
0.95	456	Raspberryhealthy	0.94	0.97
0.95	445	Soybeanhealthy	0.84	0.99
0.91	505	SquashPowdery_mildew	0.98	0.97
0.98	434	StrawberryLeaf_scorch	0.96	0.95
0.96	444	·		
0.99	456	Strawberryhealthy	1.00	0.98
0.90	425	TomatoBacterial_spot	0.85	0.95
0.84	480	TomatoEarly_blight	0.85	0.82
0.87	463	TomatoLate_blight	0.93	0.82
		TomatoLeaf_Mold	0.98	0.93
0.95	470	TomatoSeptoria_leaf_spot	0.93	0.76
0.84	436 TomatoSpi	der_mites Two-spotted_spider_mite	0.93	0.95
0.94	435	TomatoTarget_Spot	0.91	0.86
0.88	457 Tomat	coTomato_Yellow_Leaf_Curl_Virus	0.97	0.98
0.97	490	TomatoTomato_mosaic_virus	0.99	0.96
0.97	448			
0.97	481	Tomatohealthy	0.95	1.00
		accuracy		
0.94	17572	macro avg	0.94	0.94
0.94	17572	weighted avg	0.94	0.94
0.94	17572	weighted avg	0.94	0.94

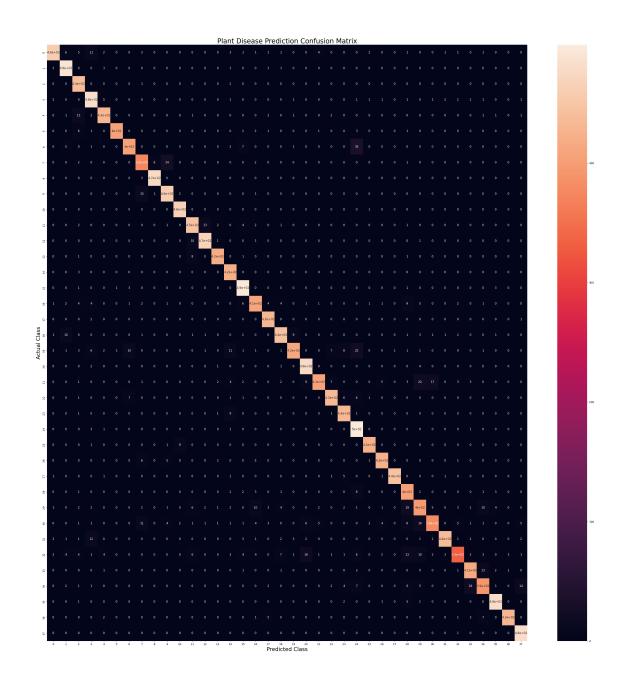
[34]: <IPython.core.display.HTML object>

1.8.1 Confusion Matrix Visualization

```
[38]: cm = confusion_matrix(Y_true, predicted_categories)

[42]: plt.figure(figsize=(40, 40))
    sns.heatmap(cm,annot=True,annot_kws={"size": 11})

    plt.xlabel('Predicted Class',fontsize = 20)
    plt.ylabel('Actual Class',fontsize = 20)
    plt.title('Plant Disease Prediction Confusion Matrix',fontsize = 25)
    plt.show()
```



[29]: