IMAGE PROCESSING FOR PLANTIX USING CONVOLUTIONAL NEURAL NETWORKS.

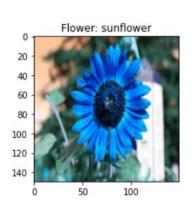
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WHAT IS A CONVOLUTIONAL NEURAL NETWORK?

- 1) Deep Learning algorithms
 - 1.1) Classify images based on multi class classification.
 - 1.2) Segmentation to find probability of each class possible.

Input features for Convolutional Neural Network

- Images extracted by cv2 library.
- Default mode is BGR.
- Images are resized to hard coded dimensions.
- Label images to get accuracy.



HOW DOES THE INPUT LOOK LIKE?

INPUT FOR CONVOLUTIONAL NEURAL NETWORKS

Input consists of images which are divided into the different classes.

Each class has multiple images based on which the training takes place.

```
import os
print(os.listdir('/content/flowers'))

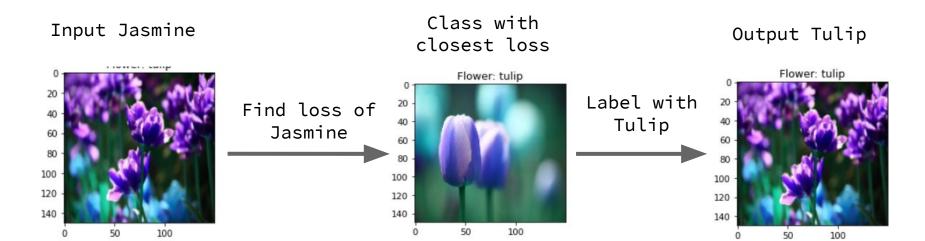
['sunflower', 'rose', 'tulip', 'dandelion', 'daisy']
```

- 5 classes
- Input trained based on only these classes.
- Output of image predicted is one of these classes.
- Images can not have any other class as output (like a jasmine or a iris)

WHAT HAPPENS TO IMAGES NOT BELONGING TO THE CLASS?

Predicts to closest possible class.

- 1) Finds the loss and validation of the input image.
- 2) Finds the class with the closest loss and validation.
- 3) Gives output with the closest class label.



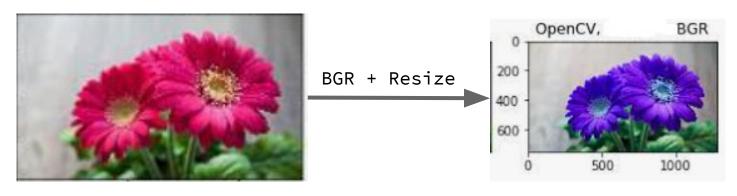
MODIFY INPUT FOR UNIFORM OUTCOMES

USING OPENCY LIBRARY

The most basic preprocessing library for image classification input.

OpenCV is generally used for preprocessing images and filtering out the input.

- 1) Convert colored images of RGB to BGR format. Also known as blue channel.
- 2) Resize images to uniform dimensions.



Size 1400 x 2800 Format RGB

Size 700 x 1400 Format BGR

CODE FOR INPUT

Import libraries like matplotlib(for plotting), cv2(for input modifications)

Filter out image

- 1) Image exists or not
- 2) Extension of jpg

Perform functions of

- 1) Image formatting to BGR
- 2) Resize image 150 x 150

Append the filtered input

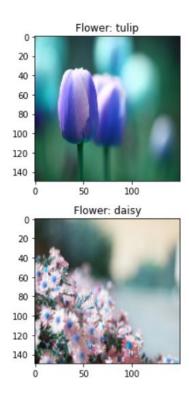
Filtered input divided into labels and images

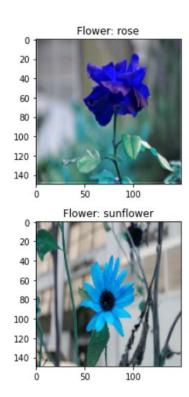
Label images with plots for dimensions

```
import matplotlib.pyplot as plt
import seaborn as sns
import cv2
features=[]
for i in categories:
    path=os.path.join(dire,i)
    num classes=categories.index(i)
    for img in os.listdir(path):
        if img.endswith('.jpg'):
            img array=cv2.imread(os.path.join(path,img),cv2.IMREAD COLOR)
            img array=cv2.resize(img array,(150,150))
            features.append([img array,num classes])
X=[]
y=[]
for i, j in features:
   X.append(i)
   y.append(j)
```

```
import matplotlib.pyplot as plt
import seaborn as sns
fig,ax=plt.subplots(5,2)
fig.set_size_inches(15,15)
for i in range(5):
    for j in range (2):
        l=np.random.randint(0,len(y))
        ax[i,j].imshow(X[1])
        ax[i,j].set_title('Flower: '+categories[y[1]])
plt.axis('off')
plt.tight_layout()
```

OUTPUT FOR THE CODE





PROCESSING THE INPUT

IMAGES TO ARRAYS

Images are readable only as matrices by the interpreter.

- 1) Use numpy to convert images to arrays.
- 2) Final resizing of images

The resizing is from $n \times m$ to inverted(a x a x 3)

An image is basically divided into its three channels, inverted from RGB to BGR and has equal dimensions of width and height after resizing.

X=np.array(X).reshape(-1,150,150,3)/255.0

Scaled from
[0, 255] to
[0, 1]. -1 is
for inverting
and 3 is for
number of
channels(BGR)
and 150 X 150
is the new
dimensions.

USING KERAS AND SKLEARN

1) Convert class matrix or numpy array to binary class matrix. It is to convert a row/column vector to matrix.

```
from tensorflow.keras.utils import to_categorical
y=to_categorical(y)
```

2) Sklearn to split the input dataset into training and testing. Ratios are used to define test size.

```
from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test=train_test_split(X,y,test_size=0.2,random_state=12)
```

Test Size images: 20 % of total images.

Training Size images: 80 % of total images.

Random State is set to any value to get uniform results according to test size. If set to None, it will give random results in test size inputs.

... MORE ABOUT KERAS

Models: Sequential - Stacking up layers in network

Layers: Dense - implement the following neural network equation

output = activation(dot(input, kernel) + bias)

For convolutional neural network:

Layers: Dropout - Prevent overfitting by setting random inputs as zero

Layers: Flatten - Convert matrix to 1 dimensional vector for input to next layer

Layers: Activation - Used as switch to on or

off a layer in the network

Layers: Conv2D - Number of filters for learning

Layers: MaxPooling2D - Find max in 2D dataset.

Layers: BatchNormalization - Helps in

coordinating after each output to uniformise.

```
from keras import backend as K
from keras.models import Sequential
from keras.layers import Dense
from keras.optimizers import Adam,SGD,Adagrad,Adadelta,RMSprop
```

specifically for cnn

from keras.layers import Dropout, Flatten,Activation
from keras.layers import Conv2D, MaxPooling2D, BatchNormalization

import tensorflow as tf
from keras.preprocessing.image import ImageDataGenerator

MODEL STRUCTURE AND SEQUENCE

SEQUENTIAL MODEL AND STRUCTURE

1) Define a sequential model

```
model = Sequential()
```

2) Add layers based on keras imports

model.add

3) Remove layers based on keras imports

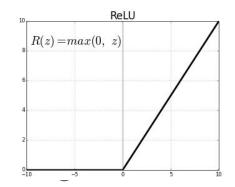
4) 4 layers used

model.pop()

LAYER 1

```
model.add(Conv2D(32, (3, 3), input_shape=(150,150,3)))
model.add(Activation("relu"))
model.add(MaxPooling2D(2, 2, padding="same"))
model.add(Dropout(0.2))
```

- 1) Conv2D layer
 - 1.1) Input of 150 \times 150 \times 3 dimensions
 - 1.2) Binarize using 32 (3 x 3) filters
- 2) Activation used relu function
- 3) MaxPooling2D
 - 5.1) Find maximum among 2 x 2 squares in 🖟



- 4) Dropout 0.2
 - 6.1) If accuracy is not high, 20 % of nodes are randomized to zero or dropped

OUTPUT OF LAYER 1

1) Conv2D:

Number of filters: f

```
conv2d (Conv2D) (None, 148, 148, 32) 896

activation (Activation) (None, 148, 148, 32) 0

max_pooling2d (MaxPooling2D) (None, 74, 74, 32) 0

dropout (Dropout) (None, 74, 74, 32) 0
```

Input - Image: m x n and Filter: a x b

Output - Image: $(m - a + 1) \times (n - b + 1) \times f$

 $= (150 - 3 + 1) \times (150 - 3 + 1) \times 32 = 148 \times 148 \times 32 =$ Input for next

2) MaxPooling2D:

Number of filers: f

Input - Filter: c x d

Output - Image: $(m / c) \times (n / d) \times f$

 $= (148 / 2) \times (148 / 2) \times 32 = 74 \times 74 \times 32$

LAYER 2 AND LAYER 3

Layer 2 and Layer 3: increased number of filters

The formulas used in Conv2D:

$$m \times n \times f$$
 $\xrightarrow{a \times b}$ $(m - a + 1) \times (n - b + 1) \times f$

The formulas used in MaxPool2D:

LAYER 4

```
model.add(Flatten())
model.add(Dense(512, activation="relu"))
model.add(Dropout(0.2))
model.add(Dense(128, activation="relu"))
model.add(Dropout(0.2))
model.add(Dense(5, activation="softmax"))
```

1) Flatten matrix to vector

$$a \times b \times f \xrightarrow{\text{Flatten}} (a \times b \times f) \times 1 \times 1$$

2) Dense to hardcode reshaping of matrix to converge

(a x b x f) x 1 x 1
$$\xrightarrow{\text{Dense by h}}$$
 h x 1 x 1 dimensions

3) Apply dropout if necessary

Final output is given by softmax for

Final output is given by softmax function

$$\sigma(ec{z})_i = rac{e^{z_i}}{\sum_{j=1}^K e^{z_j}}$$

OUTPUT OF LAYER 4

Since there are 5 classes of flowers, there will be a dense function with 5 as the required output dimensions for class predictions.

flatten (Flatten)	(None,	36992)	0
dense (Dense)	(None, S	512)	18940416
dropout_3 (Dropout)	(None, S	512)	0
dense_1 (Dense)	(None, 1	128)	65664
dropout_4 (Dropout)	(None, 1	128)	0
dense_2 (Dense)	(None,	5)	645

DATA GENERATOR

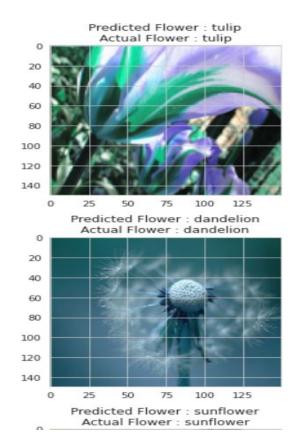
The functions on the models can be slightly modified with the following for more accuracy.

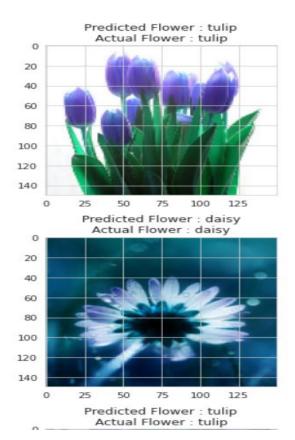
APPLY GENERATOR

BATCH SIZE = 128

NUMBER OF STEPS = 3328 / 128 = 26 per EPOCH

OUTPUT OF THE CONVOLUTIONAL NEURAL NETWORK





PROJECT WORK

INPUT

39 classes of leaves
Total number of images 3900

```
[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 ...
[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 ...
[INFO] Processing Squash Powdery mildew ...
[INFO] Processing Strawberry healthy ...
[INFO] Processing Orange Haunglongbing (Citrus greening) ...
[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) ...
[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

MODEL USED

7 Layers
Used Conv2D, MaxPool2D,
ReLu, Softmax, etc.
functions

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
activation 7 (Activation)	(None,	39)	0

(None, 256, 256, 32)

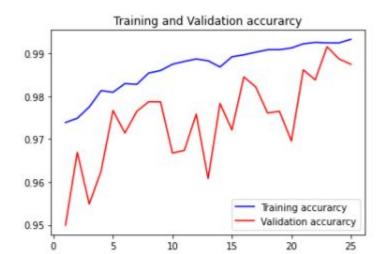
896

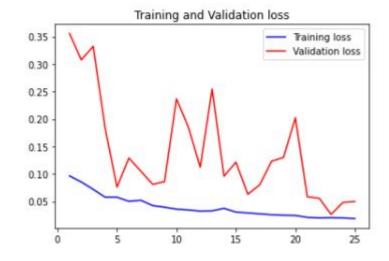
conv2d_1 (Conv2D)

OUTPUT

As the number of EPOCHS increased accuracy increased and loss decreased
Final accuracy = 98.75 %

Test Accuracy: 98.74754548072815



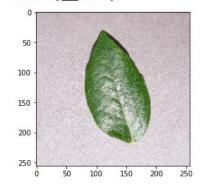


TESTING

Used images to test

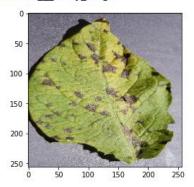
predict_disease('/content/PlantVillage/val/Blueberry__healthy/008c85d0-a954-4127-bd26-861dc8a1e6ff__RS_HL 2431.JPG')

Blueberry healthy



predict_disease('/content/PlantVillage/val/Potato___Early_blight/03b0d3c1-b5b0-48f4-98aa-f8904670290f___RS_Early.B 7051.JPG')

Potato___Early_blight



RESOURCES

- 1. IEEE papers provided by Bhavin Sir.
- 2. My own github account having all the source code for the images used in the presentation:

https://github.com/mehul14062001/Practice-School-1