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Identification of Medicinal Plants using Deep Learning

R. Upendar Rao¹, M. Sai Lahari², K. Pavana Sri³, K. Yaminee Srujana⁴, D. Yaswanth⁵

¹Associate Professor, Dept of ECE, NRI Institute of Technology (Autonomous) (Affiliated to Jawaharlal Nehru Technological University Kakinada), Pothavarappadu, Andhra Pradesh

^{2, 3, 4, 5}Student Dept of ECE, NRI Institute of Technology (Autonomous), (Affiliated to Jawaharlal Nehru Technological University Kakinada), Pothavarappadu, Andhra Pradesh

Abstract: Identification of the correct medicinal plants that goes in to the preparation of a medicine is very important in ayurvedic, folk and herbal medicinal industry. The main features required to identify a medicinal plant is its leaf shape, color and texture. Color and texture from both sides of the leaf contain deterministic parameters to identify the species. In this project we explore feature vectors from both the front and back side of a green leaf along with morphological features to arrive at a unique optimum combination of features that maximizes the identification rate. A database of medicinal plant leaves is created from scanned images of front and back side of leaves of commonly used medicinal plants. The leaves are classified based on the shape and dimension combination. It is expected that for the automatic identification of medicinal plants this system will help the community people to develop their knowledge on medicinal plants, help taxonomists to develop more efficient species identification techniques and also participate significantly in the pharmaceutical drug manufacturing.

Keywords: Classification, feature extraction, morphological features, optimization, plant identification, texture features.

I. INTRODUCTION

A. Introduction

The world bears thousands of plant species, many of which have medicinal values, others are close to extinction, and still others that are harmful to man. Not only are plants an essential resource for human beings, but they form the base of all food chains. The medicinal plants are used mostly in herbal, ayurvedic and folk medicinal manufacturing.

Herbal plants are plants that can be used for alternatives to cure diseases naturally. About 80% of people in the world still depend on traditional medicine. Meanwhile, according to herbal plants are plants whose plant parts (leaves, stems, or roots) have properties that can be used as raw materials in making modern medicines or traditional medicines. These medicinal plants are often found in the forest. There are various types of herbal plants that we can know through the identification of these herbs, one of which is using identification through the leaves. and protect plant species, it is crucial to study and classify plants correctly. Combinations of a small subset amounting to 1500 of these plants are used in Herbal medicines of different systems of India. Specifically, commercial Ayurvedic preparations use 500 of these plants. Over 80% of plants used in ayurvedic formulations are collected from the forests and wastelands whereas the remaining are cultivated in agricultural lands. More than 8000 plants of Indian origin have been found to be of medicinal value.

B. Motivation

In the ancient past, the Ayurvedic physicians themselves picked the medicinal plants and prepared the medicines for their patients. Today only a few practitioners follow this practice. The manufacturing and marketing of Ayurvedic drugs has become a thriving industry whose turnover exceeds Rs. 4000 crores. The number of licensed Ayurvedic medicine manufacturers in India easily exceeds 8500. This commercialization of Ayurvedic sector has brought in to focus several questions regarding the quality of raw materials used for Ayurvedic medicines. Today the plants are collected by women and children from forest areas; those are not professionally trained in identifying correct medicinal plants. Manufacturing units often receive incorrect or substituted medicinal plants. Most of these units lack adequate quality control mechanisms to screen these plants. In addition to this, confusion due to variations in local name is also rampant. Some plants arrive in dried form and this make the manual identification task much more difficult. Incorrect use of medicinal plants makes the Ayurvedic medicine ineffective. It may produce unpredictable side effects also. In this situation, strict measures for quality control must be enforced on Ayurvedic medicines and raw materials used by the industry in order to sustain the present growth of industry by maintaining the efficacy and credibility of medicines.

A trained Botanist looks for all the available features of the plants such as leaves, flowers, seeds, root and stem to identify plants. Except for the leaf, all others are 3D objects and increase the complexity of analysis by computer. However, plant leaves are 2D objects and carry sufficient information to identify the plant. Leaves can be collected easily and image acquisition may be carried out using inexpensive digital cameras, mobile phones or document scanners. It is available at any time of the year in contrast to flowers and seeds. Leaves acquire a specific colour, texture and shape when it grows and these changes are relatively insignificant. Plant recognition based on leaves depends on finding exact descriptors and extracting the feature vectors from it. Then the feature vectors of the training samples are compared with the feature vectors of the test sample to find the degree of similarity using an appropriate classifier.

C. Problem Definition

Deep learning is one of the major subfields of machine learning framework. Machine learning is the study of design of algorithms, inspired from the model of human brain. Deep learning is becoming more popular in data science fields like robotics, artificial intelligence (AI), audio & video recognition and image recognition. Artificial neural network is the core of deep learning methodologies. Deep learning is supported by various libraries such as Theano, TensorFlow, Caffe, Mx net etc., Keras is one of the most powerful and easy to use python library, which is built on top of popular deep learning libraries like TensorFlow, Theano, etc., for creating deep learning models. Detection of correct medicinal leaves can help botanists, taxonomists and drug manufacturers to make quality drug and can reduce the side effects caused by the wrong drug delivery. To identify the leaves of the plants, a type of artificial neural network called convolutional neural network (CNN) is used. The architecture we used here is Densenet121, which is a convolutional neural network that is a powerful model capable of achieving high accuracies on challenging datasets.

II. SYSTEM ANALYSIS

A. Existing System

To overcome the problems of conventional classification algorithms in recognizing medicinal plants, Kan et al [2017] suggested an automated classification approach depending on leaf images of medicinal herbs. The technique will initially preprocess medicinal plant leaf images, then calculate 10 shape features and 5 texture features, and then identify medicinal plant leaves through a support vector machine (SVM) classification. The classification model was used to identify twelve various medicinal plant leaf images, with an average rate of success of 93.3 %. The findings demonstrate that utilizing multi-feature extortion of leaf pictures in connection with SVM, it is possible to automatically categorize medicinal herbs. The report presents a useful conceptual framework for medicinal herbs categorization model research and development.

Alimboyong et al. [2018] suggested a computer vision method for recognizing ayurvedic medicinal plant species present in India's Western Ghats. A mixture of Surface plot and HOG attributes derived from leaf images, as well as a categorization utilizing a k-NN classifier, are used in the suggested approach. Studies have yielded findings that appear to be adequate for developing apps for real-world use. Prasad and Singh [2017] developed an information exchange from object recognition to plant genetic analysis using deep attributes to describe the original plant leaf image. Such deep attributes have been shown to outperform the current in plant species identification in experiments. The study demonstrated a novel and effective leaf collection method. The image is then translated into a device-independent lab color space, which is then utilized to construct the VGG-16 feature map. To boost the effectiveness of species identification, this feature map is re-projected to PCA subspace. The study utilizes two kinds of plant leaf collections to demonstrate the sturdiness.

Turkoglu and Hanbay [2019] employed image processing methods like color, vein attributes, Fourier Descriptors, and Gray-Level Co-occurrence Matrix (GLCM) approaches to retrieve features from images. Rather than obtaining attributes for the entire leaf, research advises using features collected from leaves separated into 2 or four pieces. Extreme Learning Machines (ELM) classifier calculates the separate and aggregate performances of every attribute extortion approach. The Flavia leaf database was used to test the proposed method. The suggested technique's efficiency was evaluated using 10-fold cross-validation, that was then contrasted and tabulated with approaches from other studies. Also on Fluvial leaf database, the suggested method's results were evaluated to be 99.10 %

Nuril Aslina, Nursuriati Jamil et al. used Scale Invariant Feature Transform (SIFT) as a shape descriptor and colour moments. The image is decomposed in to HSV planes and each plane is divided into 9 grids. Colour moments are calculated for each grid of every plane and used as feature vector. Least Euclidean distance between test and training sets are used for identification. Database is created by the authors by acquiring 40 leaf images of Malaysian herbs from natural habitat in natural light. An accuracy of 87.5% is obtained independent of scaling and rotation of images. SIFT is computationally intensive when used to extract key point features.

- 1) *GLCM*: Developed plant identification method using texture features alone. 10 texture features are extracted using GLCM of leaf image. These leaves are used for classification using dissimilarity method. The model is trained with 63 samples and tested with 33 leaves. As it was tested only on limited samples, the accuracy was low and it was not fully invariant to rotation of leaf.
- 2) *Root Mean Square Error*: In this method the leaf shape features and edges are extracted using predefined structural elements. Root mean square error method is used between captured image feature vectors and image in database that is trained. 10 different species of plants are considered with 40 leaf samples in the dataset. The identification rate is comparatively low to be in practical use.
- 3) *SIFT*: SIFT is used as a shape descriptor and color moments. Image is divided into HSV planes and each plane into 9 parts. Color moments are used as feature vectors. Least Euclidean distance is used between training and testing data set. 40 leaf species are considered.
- 4) *Geometric Features*: In this method five geometric features are used and a boundary descriptor named Directional fragment histogram for identification. 3070 scanned images are used and 897 scan like images. Another method used four geometrical features, hue invariant methods and polar fourier transform coefficients as shape descriptor for identification. The colour descriptor used is mean, standard deviation, skewness and kurtosis of RGB color planes. 100 samples are considered out of which 50 images are used for testing. This method is not fully invariant to rotation.

B. Proposed System

A novel method for identification of medicinal plants from images of different angles of both front and backside of the leaves have been proposed. The work is based on a database of leaf images of medicinal plants. Unique features of texture and shape combinations of morphological features have been identified, that maximizes the identification rate of green leaves. By using this method, when the image of any plant leaf is given to the system it gives whether the leaf belongs to the medicinal plant or not, local name, scientific name and properties of the leaf or the disease it cures along with the image of the leaf are displayed. In this method Dense Net type of Convolutional Neural Network(CNN) is used because of its several compelling advantages like it strengthens the feature propagation and also encourages feature reuse, which in turn increases the efficiency and decrease the loss of valuation. Here Keras is used for training the data to the model.

- 1) *Data*: The leaf samples of medicinal plants were collected. At least 30 leaves of 50 different medicinal plant species were collected. Perform a basic manual sampling and remove severely damaged leaves. Select 30 leaves of each species for scanning and 60 image samples per medicinal plant species are generated. The top and bottom faces of the leaves are captured. The below table is the list of leaves collected.

| S.No | Local name | Scientific name |
|------|------------------------|---------------------------|
| 1 | Asthma plant | Euphorbia hirta |
| 2 | Avaram | Senna auriculata |
| 3 | Balloon vine | Cardiospermum halicacabum |
| 4 | Bellyache bush (Green) | Jatropha gossypifolia |
| 5 | Benghal day flower | Commelia benghalensis |
| 6 | Big caltrops | Gokru bada |
| 7 | Black honey shrub | Phyllanthus reticulatus |
| 8 | Bristly wild grape | Cyphostemma setosum |
| 9 | Butterfly pea | Clitoria ternatea |
| 10 | Cape gooseberry | Physalis peruviana |
| 11 | Coat buttons | Tridax procumbens |
| 12 | Common wireweed | Sida acuta |
| 13 | Country mallow | Sida cortifolia |
| 14 | Crown flower | Caltropis gigantea |
| 15 | Green chireta | Andrographis paniculate |
| 16 | Heart leaved moon seed | Tinospora cardifolia |
| 17 | Holy basil | Ocimum tenuiflorum |

| | | |
|----|-------------------------|---------------------------|
| 18 | Indian copperleaf | Acalypha indica |
| 19 | Indian jujube | Ziziphus maritiana |
| 20 | Indian sarsaparilla | Hemidesmus indicus |
| 21 | Indian stinging nettle | Urtica dioica |
| 22 | Indian thorn apple | Datura inoxia |
| 23 | Indian worm weed | Artemisia absinthium |
| 24 | Ivy gourd | Coccinia grandis |
| 25 | Kokilaksha | Hygrophila auriculata |
| 26 | Land caltrops | Tribulus terrestris |
| 27 | Madagascar periwinkle | Catharanthus roseus |
| 28 | Madras pea pumpkin | Cucumis maderaspatnus |
| 29 | Malabar catmint | Anisomeles indica |
| 30 | Mexican mint | Coleus amboinicus |
| 31 | Mexican prickly poppy | Argemone Mexicana |
| 32 | Mountain knotgrass | Aerva lanata |
| 33 | Nalta jute | Corchorus olitorius |
| 34 | Night blooming cereus | Selenicereus grandiflorus |
| 35 | Panicled foldwing | Dicleptera paniculate |
| 36 | Prickly chaff flower | Achyranthus aspera |
| 37 | Punarnava | Boerhavia diffusa |
| 38 | Purple fruited eggplant | Solanum toruvum |
| 39 | Purple tephrosia | Tephrosia purpurea |
| 40 | Rosary pea | Abrus precatorious |
| 41 | Shaggy button weed | Spermacoe hispida linn |
| 42 | Small water clover | Marsilea minuta |
| 43 | Spider wisp | Cleome gynandra |
| 44 | Square stalked wine | Cissus quadrangularise |
| 45 | Stinking passion flower | Passiflora foetida |
| 46 | Sweet basil | Ocimum basillium |
| 47 | Sweet flag | Acorus calamus |
| 48 | Tinnevelly senna | Aleexandrian senna |
| 49 | Trellis vine | Liana |
| 50 | Velvet bean | Mucunna prureins |

Table 2.1 Leaf dataset used for training

III. ARCHITECTURE

A. Algorithm

- 1) *Step 1:* The input is taken in two ways- Using camera and the images.
- 2) *Step 2:* The front and back side of the leaves are scanned using a scanner that has the maximum possible resolution. These images are stored in leaf image dataset.
- 3) *Step 3:* These images are pre-processed. The dimensions of the images in the dataset are set to the required size.
- 4) *Step 4:* Now the pre-processed dataset is divided into testing and training dataset.
- 5) *Step 5:* Training data set is now driven as input to the Convolutional neural network.
- 6) *Step 6:* The output of the CNN layer along with the testing dataset is provided as input for the performance assessment. In this step the accuracy and loss of the model and validation set is considered, the accuracy and loss graphs are plotted accordingly using confusion matrix.
- 7) *Step 7:* The image i.e., the result of the output layer of convolutional neural network is displayed.

B. Flowchart

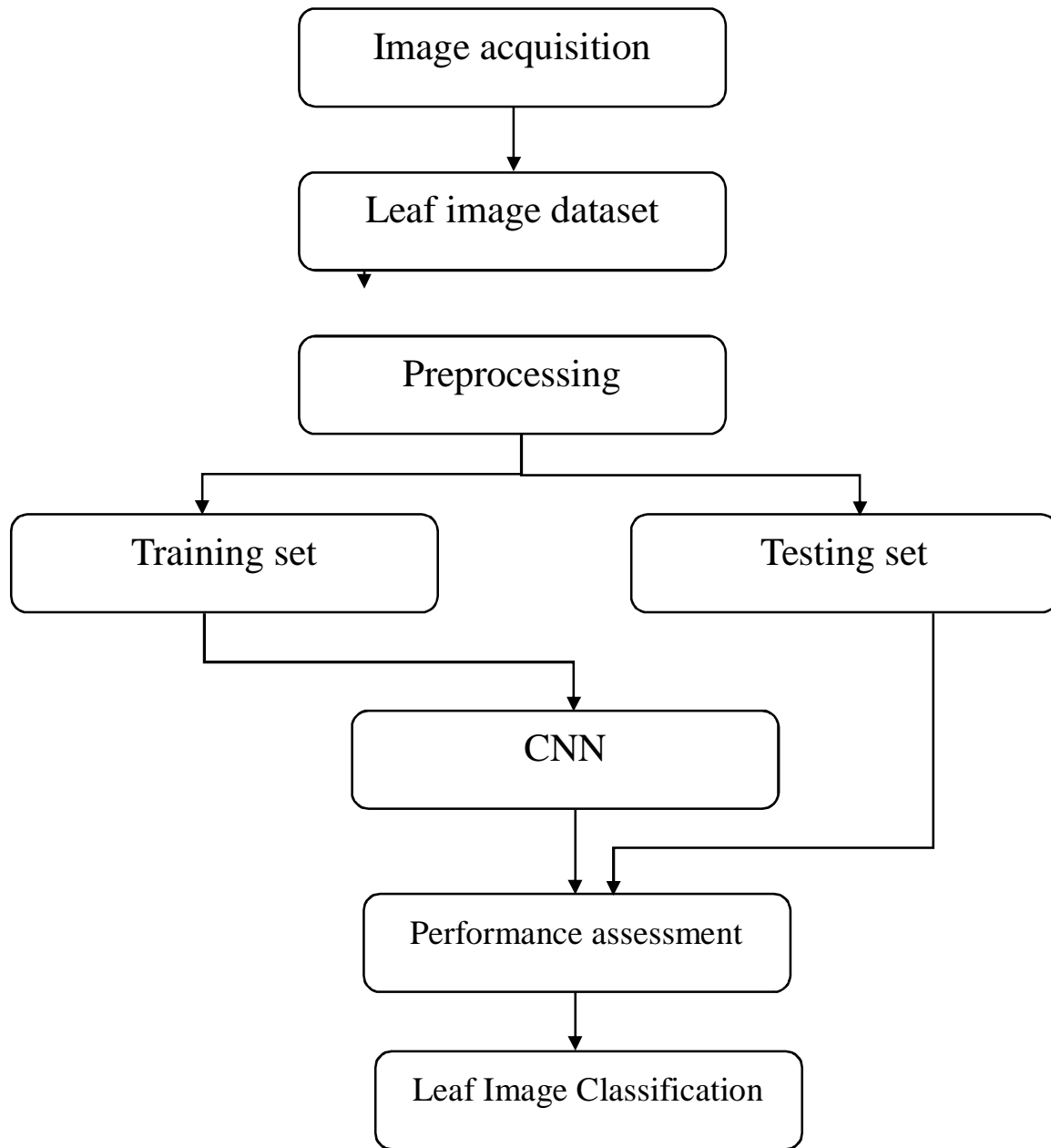


Fig 3.1 Flowchart of proposed system

IV. SOFTWARE

Python 3.9 version is used as software and the IDE used is Jupyter Notebook. Keras is used to train the model. It is a high-level neural network library which trains the deep learning model by using epochs and back propagation. Epochs means considering the data into batches and training them through iterations, while training it checks for minimum loss and maximum accuracy. DenseNet is the type of CNN used and the library used for the numerical calculations in DenseNet is Tensorflow. It is an open source software library which performs computations using dataflow graphs and provides multiple application interfaces. The input activation function used in first layer of cnn is ReLU and the output activation function used in last layer of cnn is Softmax. ReLU is a piecewise linear function that will output the input directly if it is positive, otherwise it will output zero. Softmax scales the input values between 0 and 1 i.e., it is used to normalize the output. The optimizer type used is Adam and the learning rate is 0.001. Adam optimizer is a stochastic gradient descent method that is based on the estimation of first order and second order moments.

V. OUTPUT

A. Outputs For Image Dataset

1) *Training the Data:* The considered dataset of 3777 images are trained using epochs and the accuracy and loss are calculated for each epoch. Epochs is the number of times a learning algorithm sees the complete dataset. One Epoch is when an entire dataset is passed forward and backward through the neural network only once. Loss is the summation of errors in the model and accuracy is the ratio of correct predictions to the total predictions. This process takes place through back propagation.

```
Epoch 1/30
71/71 [=====] - 61s 811ns/step - loss: 3.3095 - acc: 0.1695 - val_loss: 2.0830 - val_acc: 0.4950
Epoch 2/30
71/71 [=====] - 56s 798ns/step - loss: 2.0930 - acc: 0.4285 - val_loss: 1.3174 - val_acc: 0.6413
Epoch 3/30
71/71 [=====] - 56s 794ns/step - loss: 1.4913 - acc: 0.5808 - val_loss: 1.0691 - val_acc: 0.7042
Epoch 4/30
71/71 [=====] - 57s 801ns/step - loss: 1.2509 - acc: 0.6368 - val_loss: 0.8991 - val_acc: 0.7492
Epoch 5/30
71/71 [=====] - 57s 800ns/step - loss: 1.0414 - acc: 0.6929 - val_loss: 0.9038 - val_acc: 0.7359
Epoch 6/30
71/71 [=====] - 56s 795ns/step - loss: 0.9701 - acc: 0.7158 - val_loss: 0.7236 - val_acc: 0.7915
Epoch 7/30
71/71 [=====] - 56s 787ns/step - loss: 0.8277 - acc: 0.7485 - val_loss: 0.7134 - val_acc: 0.8008
Epoch 8/30
71/71 [=====] - 55s 782ns/step - loss: 0.7638 - acc: 0.7590 - val_loss: 0.6528 - val_acc: 0.8233
Epoch 9/30
71/71 [=====] - 702s 10s/step - loss: 0.6953 - acc: 0.7970 - val_loss: 0.6302 - val_acc: 0.8180
Epoch 10/30
71/71 [=====] - 61s 859ns/step - loss: 0.6450 - acc: 0.8032 - val_loss: 0.6109 - val_acc: 0.8206
Epoch 11/30
71/71 [=====] - 65s 917ns/step - loss: 0.5601 - acc: 0.8310 - val_loss: 0.6418 - val_acc: 0.8233
Epoch 12/30
71/71 [=====] - 57s 801ns/step - loss: 0.5803 - acc: 0.8217 - val_loss: 0.6197 - val_acc: 0.8160
Epoch 13/30
71/71 [=====] - 55s 780ns/step - loss: 0.5550 - acc: 0.8301 - val_loss: 0.5867 - val_acc: 0.8273
Epoch 14/30
71/71 [=====] - 57s 805ns/step - loss: 0.5409 - acc: 0.8301 - val_loss: 0.6598 - val_acc: 0.8120
Epoch 15/30
71/71 [=====] - 68s 962ns/step - loss: 0.4816 - acc: 0.8535 - val_loss: 0.5593 - val_acc: 0.8438
Epoch 16/30
71/71 [=====] - 70s 987ns/step - loss: 0.4398 - acc: 0.8588 - val_loss: 0.5865 - val_acc: 0.8312
Epoch 17/30
71/71 [=====] - 57s 799ns/step - loss: 0.4261 - acc: 0.8636 - val_loss: 0.5468 - val_acc: 0.8445
Epoch 18/30
71/71 [=====] - 55s 778ns/step - loss: 0.3998 - acc: 0.8707 - val_loss: 0.5734 - val_acc: 0.8478
Epoch 19/30
71/71 [=====] - 55s 777ns/step - loss: 0.4066 - acc: 0.8676 - val_loss: 0.6029 - val_acc: 0.8312
Epoch 20/30
71/71 [=====] - 55s 781ns/step - loss: 0.3817 - acc: 0.8848 - val_loss: 0.5790 - val_acc: 0.8425
Epoch 21/30
71/71 [=====] - 55s 780ns/step - loss: 0.3499 - acc: 0.8928 - val_loss: 0.5364 - val_acc: 0.8425
Epoch 22/30
71/71 [=====] - 53s 756ns/step - loss: 0.3851 - acc: 0.8773 - val_loss: 0.5810 - val_acc: 0.8359
Epoch 23/30
71/71 [=====] - 53s 748ns/step - loss: 0.3323 - acc: 0.8972 - val_loss: 0.5452 - val_acc: 0.8379
Epoch 24/30
71/71 [=====] - 53s 746ns/step - loss: 0.3007 - acc: 0.8981 - val_loss: 0.5606 - val_acc: 0.8518
Epoch 25/30
71/71 [=====] - 53s 757ns/step - loss: 0.3282 - acc: 0.8914 - val_loss: 0.5578 - val_acc: 0.8557
Epoch 26/30
71/71 [=====] - 53s 747ns/step - loss: 0.2946 - acc: 0.9117 - val_loss: 0.5623 - val_acc: 0.8524
Epoch 27/30
71/71 [=====] - 53s 744ns/step - loss: 0.2674 - acc: 0.9166 - val_loss: 0.5927 - val_acc: 0.8398
Epoch 28/30
71/71 [=====] - 53s 753ns/step - loss: 0.2728 - acc: 0.9113 - val_loss: 0.5818 - val_acc: 0.8418
Epoch 29/30
71/71 [=====] - 53s 747ns/step - loss: 0.2663 - acc: 0.9078 - val_loss: 0.5436 - val_acc: 0.8484
Epoch 30/30
71/71 [=====] - 53s 749ns/step - loss: 0.2689 - acc: 0.9184 - val_loss: 0.5944 - val_acc: 0.8445
```

Fig 5.1 Training process through back propagation

2) *Displaying the Training Images:* During the training process, some of the images are displayed that are used for training.



Fig 5.2 Displaying some of the training images

3) *Input and Output:* The path of the image from the test data is given in the path and the program is run. The output image is displayed with it's local name, scientific name and the propoerties of the leaf or the disease it cures is displayed along with it's image.

```

return np.expand_dims(x, axis=0)
path=r"D:\Lahari\project\major\Identification of medicinal plants\ayurvedic_leaf_classification\test_data\img1.jpg"
result = model.predict([prepare(path)])
d=image.load_img(path)
plt.imshow(d)
x=np.argmax(result,axis=1)
print (leaf_name[int(x)])

```

Balloon vine-Cardiospermum halicacabum-treats nervous diseases and stiffness in limbs

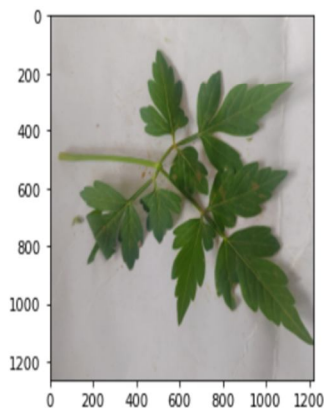


Fig 5.3 Displaying the output

4) *Output 1*

Local name – Butterfly pea

Scientific name – *Clitoria ternatea*

Uses – rich in antioxidants

Butterfly Pea-*Clitoria ternatea*-rich in antioxidants

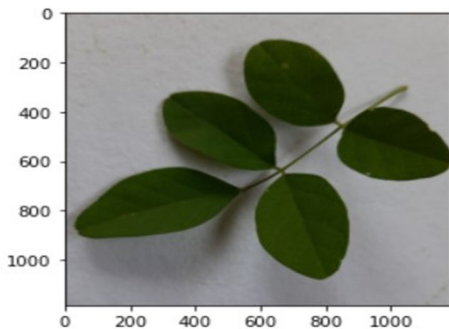


Fig 5.4 Butterfly pea

5) *Output 2*

Local name – Balloon vine

Scientific name – *Cardiospermum halicacabum*

Uses – treats nervous disorders and stiffness in limbs

Balloon vine-*Cardiospermum halicacabum*-treats nervous diseases and stiffness in limbs

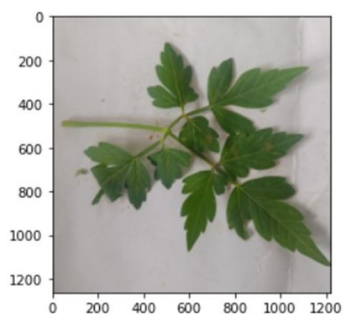


Fig 5.5 Balloon vine

6) *Output 3*

Local name – Bellyache bush

Scientific name – *Jatropha gossypifolia*

Uses – contains antimicrobial, antidiabetic properties

Bellyache bush (Green)-*Jatropha gossypifolia*-antimicrobial,antidiabetic

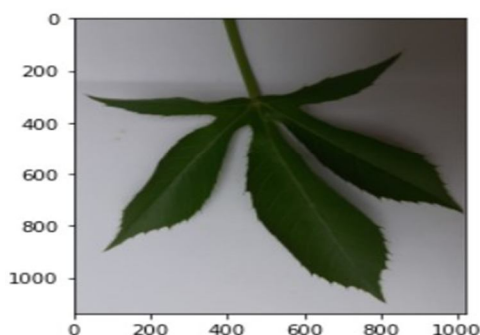


Fig 5.6 Bellyache bush

7) *Output 4*

Local name – Purple fruited pea eggplant

Scientific name – *Solanum torurum*

Uses – rich in antioxidants, used in herbal teas and cosmetics

Purple Fruited Pea Eggplant-*Solanum torurum*-tones nervous tissue and ejects phelgm from lungs

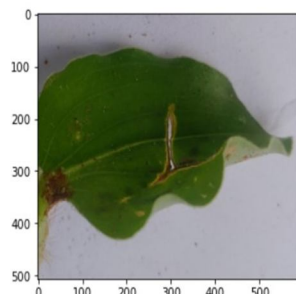


Fig 5.7 Purple fruited pea eggplant

8) *Output 5*

Local name – Asthma plant

Scientific name – *Euphoria hirta*

Uses – cures breathing disorders

Asthma Plant-*Euphoria hirta*-cures breathing disorders

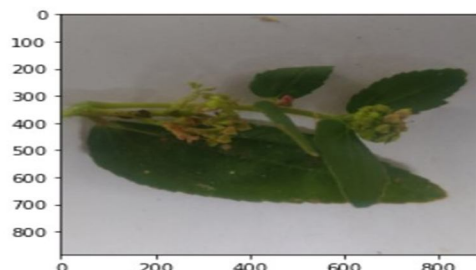


Fig 5.8 Asthma plant

9) *Output 6*

Local name – Big caltrops

Scientific name – *Gokru bada*

Uses – Used in treatment of asthma, cough, edema

Big Caltrops-*Gokru bada*-treatment of asthma, cough, edema

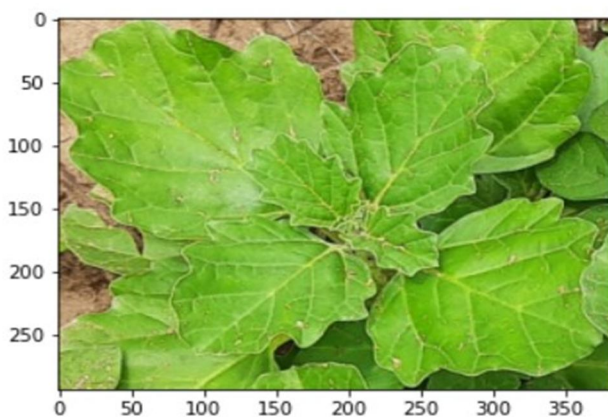


Fig 5.9 Big caltrops

B. Output For Camera Capturing

- 1) *Input-Capturing the leaf Using Camera:* Initially when the code is run, the capturing window opens to capture the leaf as shown in the figure below.

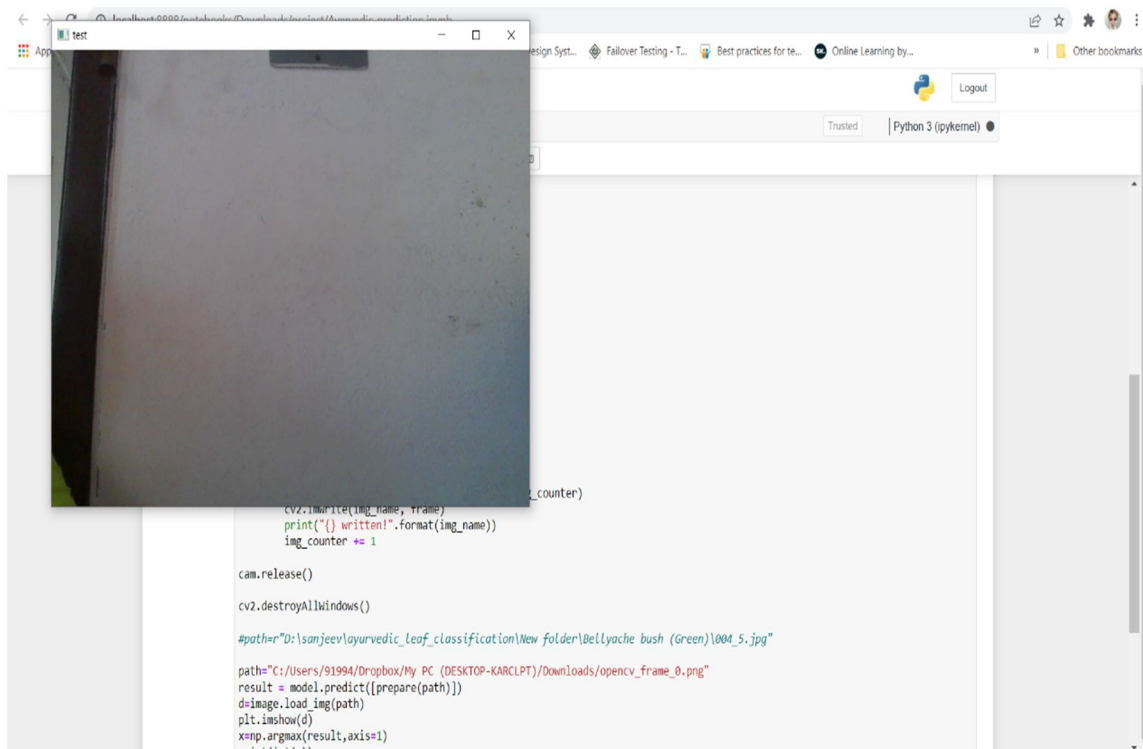


Fig 5.10 Opening of camera window

- 2) *Capturing the Frames:* The leaf is placed in front of the camera and the image is written into the model on pressing the spacebar. When the image is written into the model it is displayed as “opencv_frame_0.png written!”. Maximum of three frames can be written into the model.

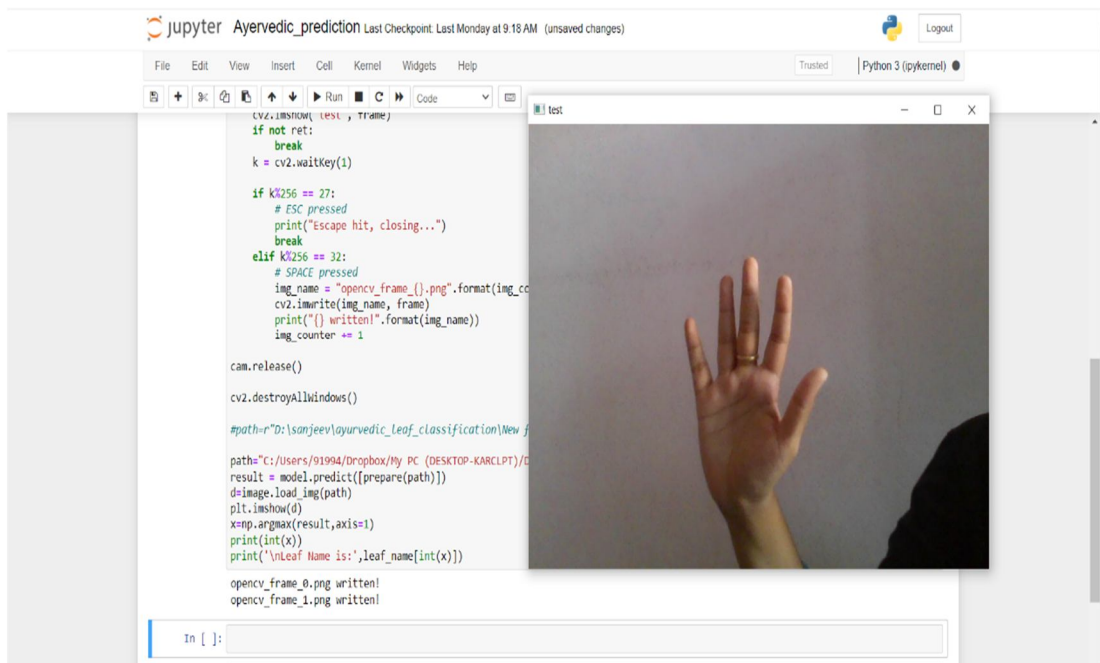


Fig 5.11 Capturing the frames

3) *Displaying the Output:* After capturing and reading into the model, escape button is pressed to close the camera window. The leaf name is displayed immediately after the camera window is closed. The closure of the camera window is also displayed as “escape hit, closing...”. Also, the number of frames captured is also displayed along with the leaf name.

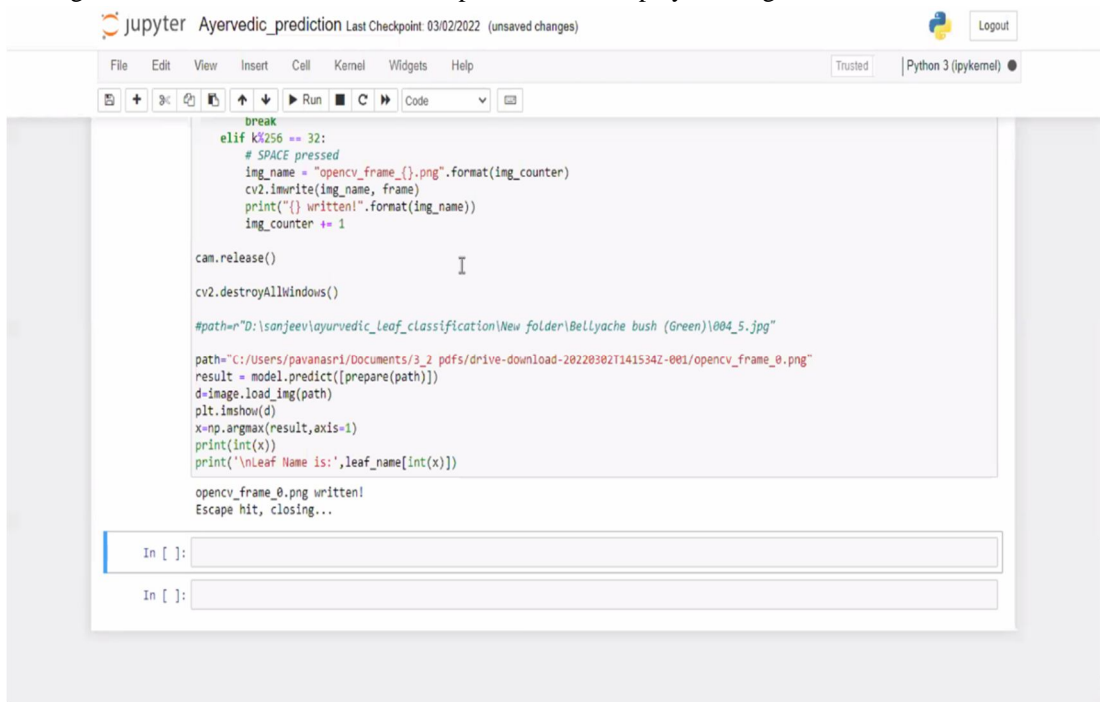


Fig 5.12 After hitting escape the camera window closes

4) *Output-1*

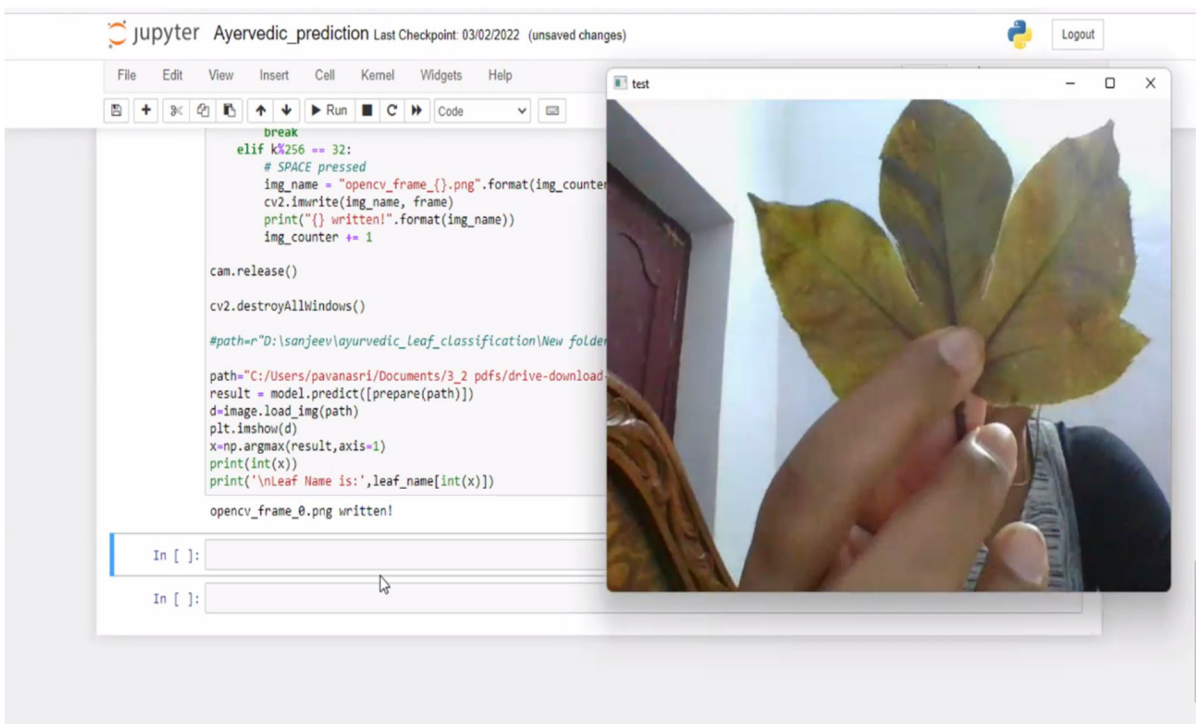


Fig 5.13 Capturing Bellyache bush(Green) leaf

Bellyache Bush(Green) leaf is considered here and is captured using camera. The camera displays the output as shown in the figure.

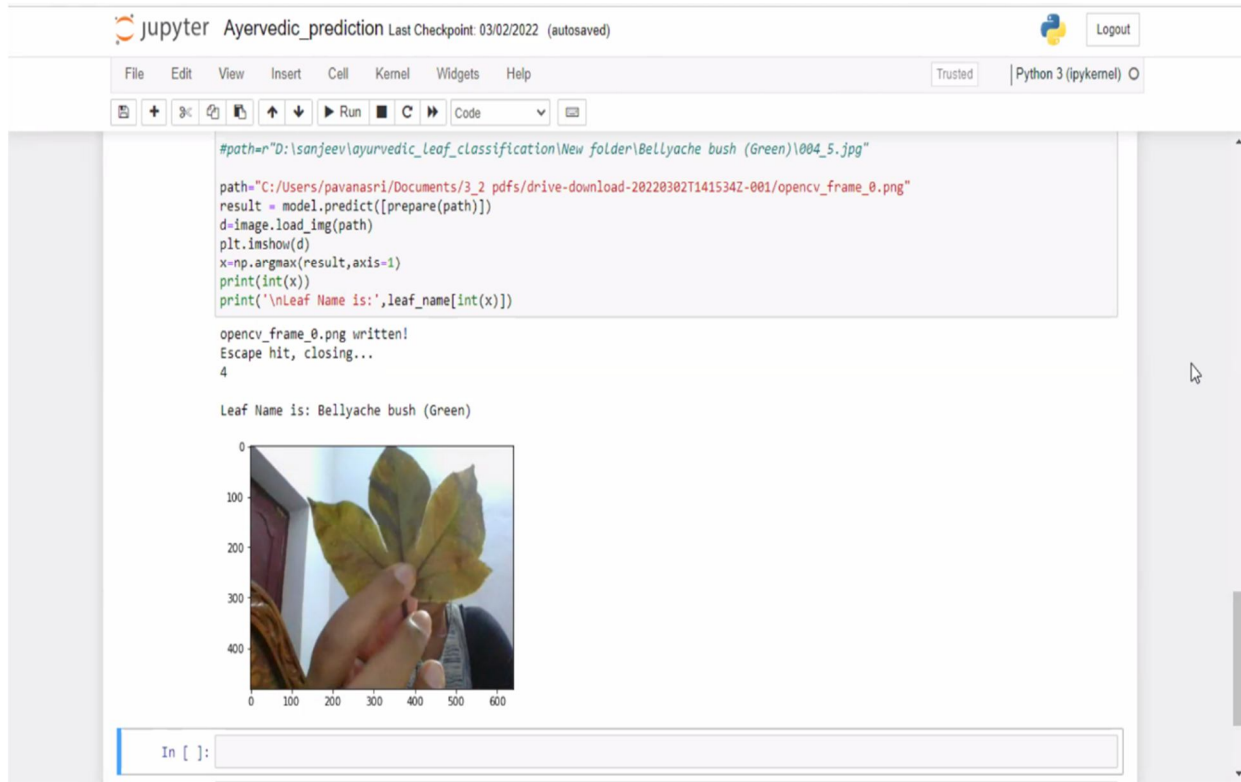


Fig 5.14 Displaying the captured leaf name (Bellyache Bush (Green))

5) Output-2

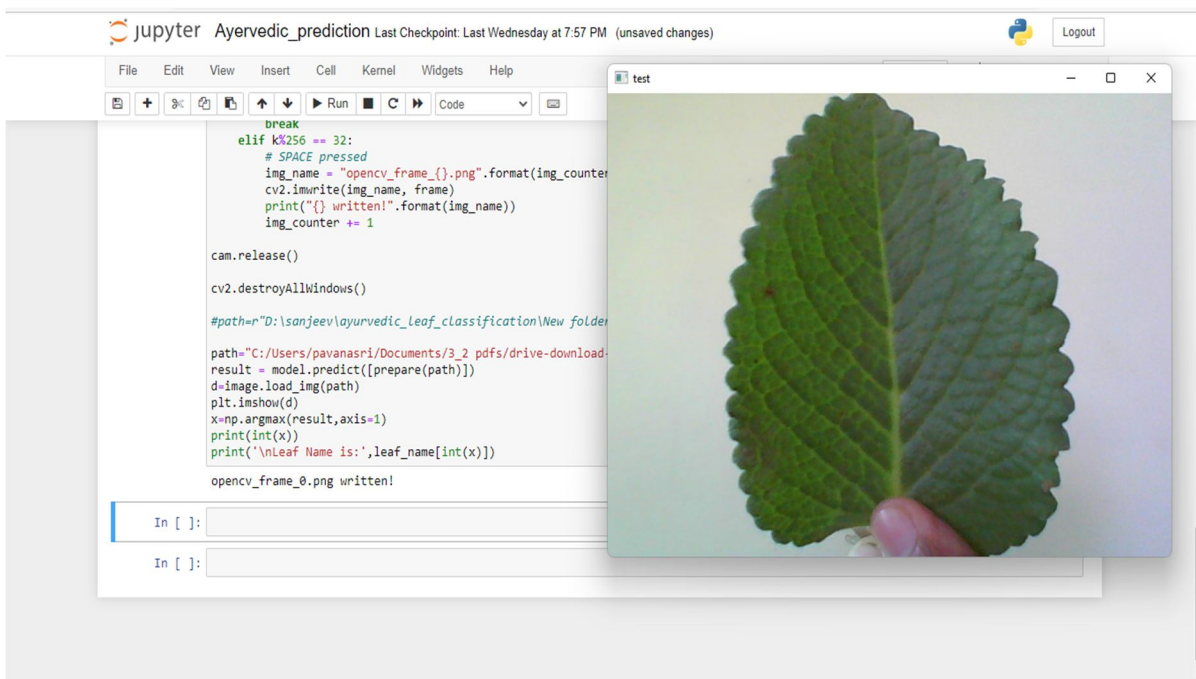


Fig 5.15 Capturing the Mexican mint leaf

Mexican mint leaf is considered here and is captured using camera. The camera displays the output as shown in the figure.

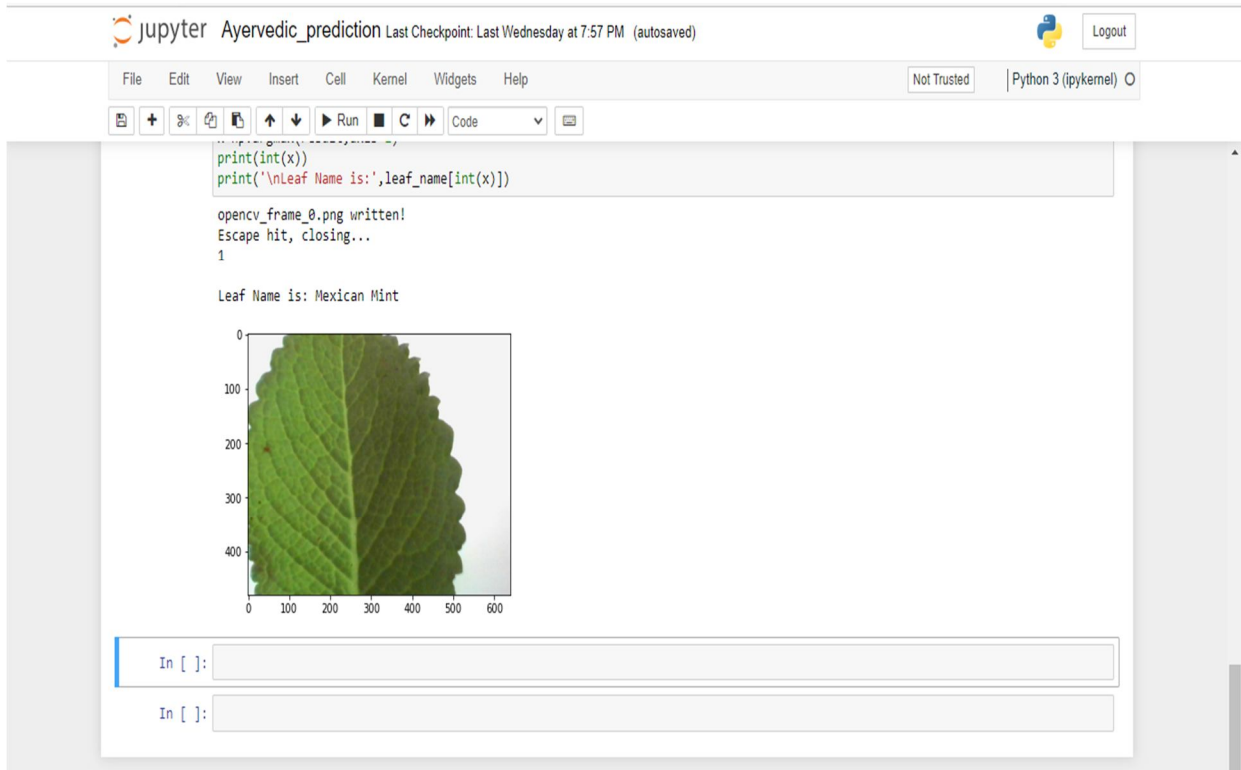


Fig 5.16 Displaying the captured leaf name (Mexican Mint)

6) Output-3

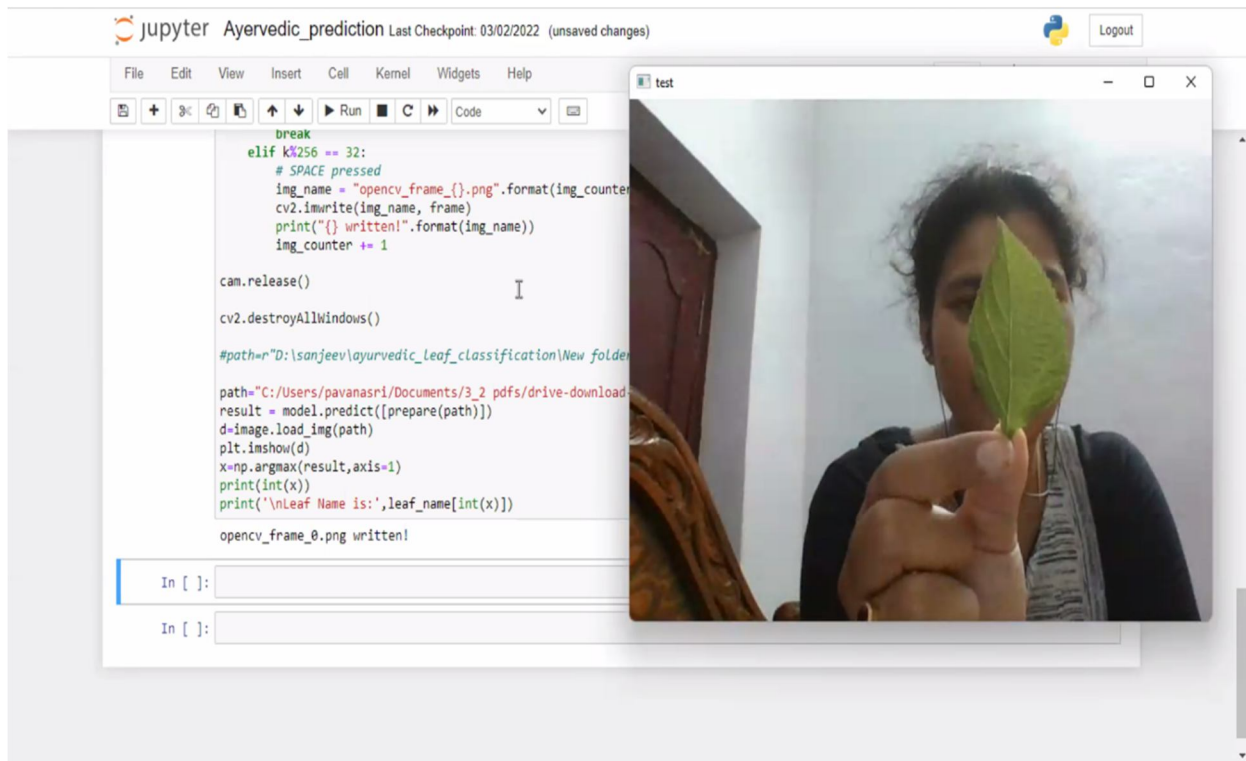


Fig 5.17 capturing of Nalta jute leaf

Nalta jute leaf is considered here and is captured using camera. The camera displays the output as shown in the figure.

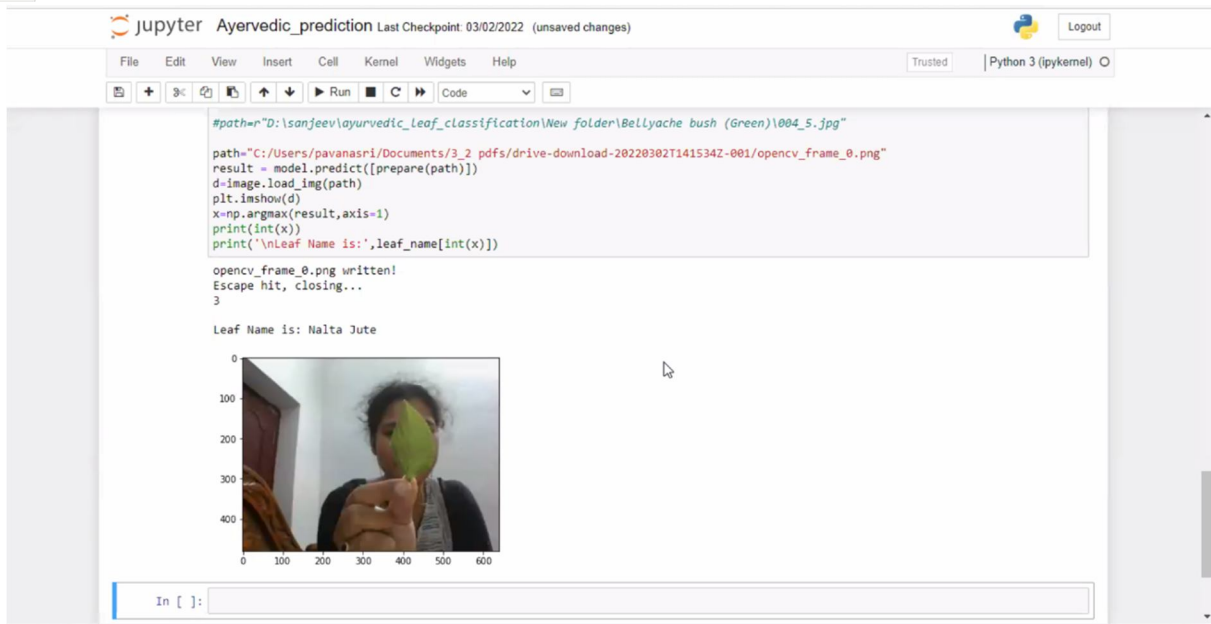


Fig 5.18 Displaying the captured leaf name (Nalta Jute)

C. Training And Validation Accuracy Graph

After the process of training, the graph between training accuracy, validation accuracy and 30 epochs is plotted. Here the blue line represents validation accuracy and the red line represents training accuracy. Accuracy is calculated using the below formula.

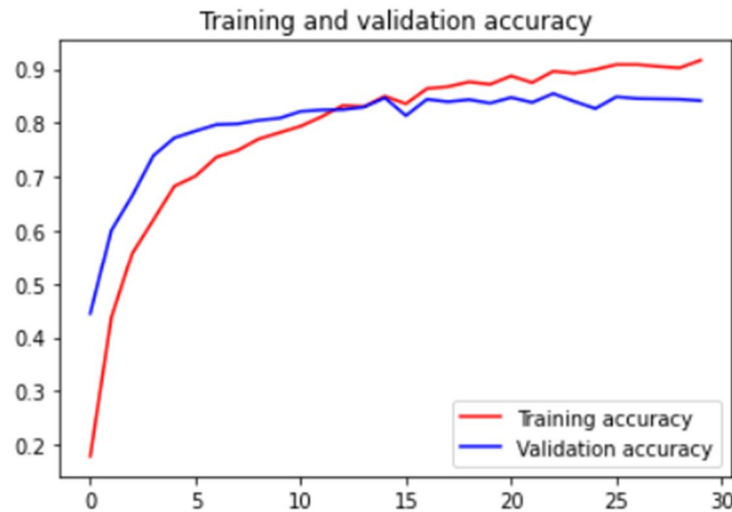


Fig 5.19 epoch vs accuracy graph

Formula to calculate accuracy:

$$Accuracy = \frac{\text{Number of correct predictions}}{\text{Total number of predictions}}$$

Number of correct predictions – positive predictions

Total numbers of predictions – sum of positive and negative predictions

Here positive predictions are the correctly predicted values according to the desired input and negative predictions are the error values or the incorrectly predicted values.

Here the values of the training accuracy and validation accuracy with respect to the epochs are mentioned in the below table.

| | Model accuracy | Validation accuracy |
|----|----------------|---------------------|
| 1 | 0.178 | 0.444 |
| 2 | 0.436 | 0.600 |
| 3 | 0.557 | 0.665 |
| 4 | 0.620 | 0.739 |
| 5 | 0.682 | 0.772 |
| 6 | 0.701 | 0.784 |
| 7 | 0.736 | 0.796 |
| 8 | 0.748 | 0.798 |
| 9 | 0.770 | 0.804 |
| 10 | 0.782 | 0.808 |
| 11 | 0.793 | 0.821 |
| 12 | 0.811 | 0.8240 |
| 13 | 0.832 | 0.8246 |
| 14 | 0.830 | 0.829 |
| 15 | 0.849 | 0.846 |
| 16 | 0.835 | 0.813 |
| 17 | 0.863 | 0.843 |
| 18 | 0.867 | 0.839 |
| 19 | 0.876 | 0.843 |
| 20 | 0.871 | 0.836 |
| 21 | 0.887 | 0.847 |
| 22 | 0.874 | 0.837 |
| 23 | 0.895 | 0.854 |
| 24 | 0.891 | 0.839 |
| 25 | 0.898 | 0.826 |
| 26 | 0.908 | 0.848 |
| 27 | 0.908 | 0.845 |
| 28 | 0.904 | 0.844 |
| 29 | 0.902 | 0.843 |
| 30 | 0.916 | 0.841 |

Table 5.1 Epoch vs model accuracy and validation accuracy

D. Training And Validation Loss Graph

After the process of training, the graph between training accuracy, validation accuracy and 30 epochs is plotted. Here the blue line represents validation accuracy and the red line represents training accuracy. Accuracy is calculated using the below formula.



Fig 5.20 epoch vs loss graph

Formula to calculate loss:

$$\text{Loss} = \frac{1}{n} \sum_{i=1}^n (Y_i - \hat{Y}_i)^2$$

Y_i – The actual output

\hat{Y}_i – The predicted output

n – number of inputs

i – Iteration

Here the values of the training accuracy and validation accuracy with respect to the epochs are mentioned in the below table.

| Epochs | Model loss | Validation loss |
|--------|------------|-----------------|
| 1 | 3.32 | 2.23 |
| 2 | 2.09 | 1.43 |
| 3 | 1.56 | 1.10 |
| 4 | 1.31 | 0.92 |
| 5 | 1.11 | 0.76 |
| 6 | 0.98 | 0.72 |
| 7 | 0.88 | 0.71 |
| 8 | 0.82 | 0.667 |
| 9 | 0.73 | 0.662 |
| 10 | 0.70 | 0.63 |
| 11 | 0.67 | 0.60 |
| 12 | 0.62 | 0.58 |
| 13 | 0.54 | 0.56 |
| 14 | 0.54 | 0.57 |
| 15 | 0.47 | 0.54 |
| 16 | 0.50 | 0.60 |
| 17 | 0.45 | 0.53 |
| 18 | 0.42 | 0.51 |
| 19 | 0.40 | 0.53 |
| 20 | 0.40 | 0.54 |
| 21 | 0.37 | 0.52 |
| 22 | 0.37 | 0.53 |
| 23 | 0.34 | 0.50 |
| 24 | 0.35 | 0.53 |
| 25 | 0.31 | 0.64 |
| 26 | 0.31 | 0.57 |
| 27 | 0.29 | 0.55 |
| 28 | 0.31 | 0.52 |
| 29 | 0.31 | 0.56 |
| 30 | 0.26 | 0.57 |

Table 5.2 Epochs vs model loss and validation loss

VI. CONCLUSION

Plants are necessary for human survival. Herbs, particularly, are employed by indigenous populations as folk medicines from old period. Herbs are typically recognized by clinicians based on decades of intimate sensory or olfactory experience. Recent improvements in analytical technology have made it much easier to identify herbs depending on scientific evidence. This helps a lot of individuals, particularly those are not used to recognising herbs. additionally for time-consuming methods, laboratory-based analysis necessitates expertise in sample healing and data explanation. As a result, a simple and reliable method for identifying herbs is required. Herbal identification anticipated to benefit from the combination of computation and statistical examination. This non-destructive technique will be the preferred approach for quickly identifying herbs, especially for individuals who cannot able to use expensive analytical equipment. This work reviews about different methods for plants recognition and also reviews their advantages and disadvantages.

VII. FUTURE SCOPE

The proposed methods are not suitable for tiny leaves or plants without a proper leaf. Efforts may be made to develop methods to identify these types of plants. The algorithms may be implemented on a standalone single board computer connected to a scanner. A portable system may be developed for field use. In future research in the area of plants identification, improved machine learning classifier with some pre-processing and feature selection models will be used to solve the accuracy related issues and enhance the performance.

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45.98



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