



## Agriculture based plant leaf health assessment tool: A Deep Learning perspective

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### ABSTRACT

For the most part, mango leaves principally get influenced by three basic sicknesses they are Anthracnose, Bacterial canker, Powdery mildew. The previously mentioned infections influence the development of mango tree, decline the life expectancy and decrease the natural product creation. Thinking about this we were doing the task under the territory of leaf malady characterization. The fundamental point of our venture is to arrange the state of the leaf, needs to recognize the malady from which it is experiencing. Thus, it will be for the most part helpful for ranchers keeping from crop misfortune financially. They can kill the illness in the underlying state itself. We were utilizing the profound learning system to recognize the tainted leaves. We propose a CNN single-stream model to order the picture. Our informational collection comprises a sum of 800 pictures arranged into two kinds, the main sort comprises a preparation set and the subsequent kind comprises of the testing set. Preparing comprises of 150 pictures and testing comprises of 50 pictures in every envelope.

**Key words:** CNN, Max-pooling, Convolution, Soft-max, dense layers.

### 1. INTRODUCTION

Picture classification is an energetic region of research in picture comprehension and PC vision. Inexhaustible classifiers have been proposed in writing for various applications. In this paper, we propose to utilize a portion of these models of feature extraction and classification. Fungal illnesses are normal in plant leaves. The illnesses in the plants are caused by dropping the quality and the amount of the horticulture production. The plant infections influence the nature of the leaves, organic products, stem, vegetables, and their items. This vigorously impacts efficiency and reflects against the price. Food and Agricultural Organization evaluated that the total populace will reach nine billion in the upcoming years, in this manner requiring about seventy percent development in the nourishment generation for a relentless inventory. The illnesses are the biotic variables that are either brought about by the organisms, microbes or green growth though, disarranges are the abiotic components brought about by the temperature, precipitation, supplement insufficiency,

dampness and so forth. Consequently, in the field of agriculture, identification of infection in plants assumes a significant job. To distinguish a plant ailment in the extremely introductory stage, utilization of programmed sickness discovery strategy is valuable. For example, illnesses named are an Anthracnose ailment, powdery mildew, bacterial canker found in the mango tree. In such situations, early location could have been productive. Automatic detection of the diseases is cheaper in cost and just by seeing the diseased symptoms on the plant leaves, detection becomes simple. The proposed CNN strategy is programmed, computationally it is proficient. Along these lines, this work proposes a profound learning procedure named as CNN for the arrangement of leaves contaminated by bacterial canker, anthracnose, and powdery mildew. Deep Learning can be comprehended as a calculation that is made out of concealed layers of different neural systems. It takes a shot at solo information and is known to give precise outcomes than conventional calculations. As it goes further and more profound, it channels the intricate highlights and consolidates with those of the last layer, in this manner better outcome. Profound learning structures, for example, profound neural systems, profound conviction systems, repetitive neural systems and convolutional neural systems have been applied to fields including PC vision, discourse acknowledgment, normal language handling, sound acknowledgment, informal organization sifting, machine interpretation, bioinformatics, medicate plan, therapeutic picture examination, material review and table top game projects, where they have delivered results similar to and at times better than human specialists.

### 2. LITERATURE REVIEW

In [1] the deep learning approach that depends on improved convolutional neural systems for the constant discovery of apple leaf sicknesses. Advances utilized are information expansion and picture comment. Models utilized are GoogLe-Net Inception structure, Rainbow link, R-CNN, VGG-INCE, AlexNet, INAR-SSD are utilized. Generating the apple leaf malady image annotation and data augmentation then the detection by the single-shot multi-box detector. Structure of origin module, VGG-INCEP network model is for the most part utilized. In the exploratory assessment, a trial arrangement is done, the informational index of 26,377

pictures are gathered. Examination of pre-organize acknowledgment precision, disarray lattice, result correlation of different location calculations, information expansion examination tests, highlight representation process, exactness versus speed, identification representation, and disappointment investigation are prepared. Alternaria leaf spot, grey spot, brown spot, rust, mosaic is five normal sorts of apple leaf ailments that seriously influence apple yield. The outcome is broke down as the INAR-SSD model understands a location execution of 78.80% mAP on ALDD, with a high-detection speed of 23.13 FPS of the sickness contaminated to the apple leaf.

In [2] the field of farming data, automatic identification, and conclusion of maize leaf illnesses are finished. Models utilized are improved GoogLeNet, Cifar10 dependent on profound getting the hang of, pooling blends, dropout tasks, and rectified straight unit capacities are done. Dataset is gathered, the growth strategy is utilized, picture pre-handling and marking ought to be finished. Convolutional neural systems procedure will be finished. Steps of the stream are convolution, initiate the capacity, pooling, dropout, misfortune work, hyper parameters to be determined from the given informational index. Results are taken from the tested examination for the given dataset. Nine kinds of maize leaves are recognized by utilizing profound CNN, the exactness rate is acquired with 98.8%, 98.9% for the two models.

### 3. METHODOLOGY

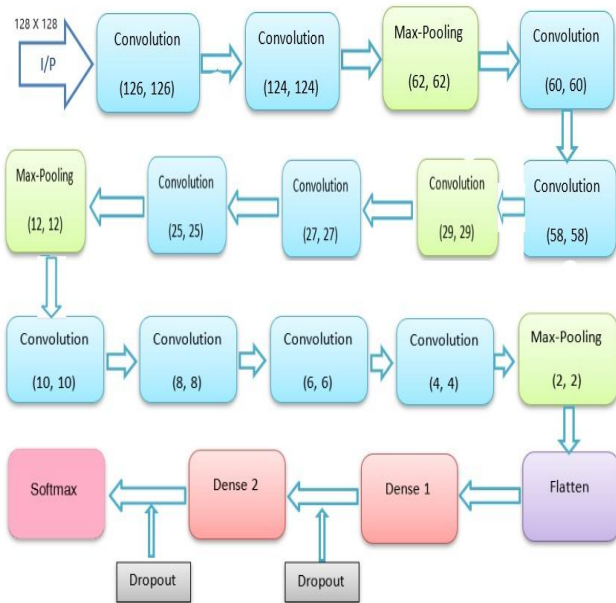


Figure 1: Block diagram

Fig 1: This block diagram explains the layers associated with the project. The information is given as a picture of size 128\*128 after the convolution layer the yield will be 126\*126. The convolution layer will extract in the form of a matrix. Again convolution layer happens and the size of the picture

gets 124\*124. Convolutional layers apply convolution operation to the information, passing the outcome to the following layer. The convolution copies the reaction of an individual neuron to visual boosts. In the maximum pooling layer, the most well-known sort of pooling is a pooling layer of filter size (2,2) utilizing the MAX activity. It would take the limit of each 4\*4 grid of the first picture. Again convolution is applied to the came about sign then the yield gets 60\*60, max-pooling is applied the yield gets 12\*12. Again 4 layers of convolution are applied and the yield gets 4\*4. At that point, max-pooling is applied yield gets 2\*2. It is levelled and goes through the dense layers and the dropout layer is included for each thick layer. In conclusion, it is given to the soft-max layer. The soft-max work computes the probabilities of each target class's overall conceivable objective classes. Later the determined probabilities will be useful for determining the target class for the given source inputs. The principle bit of leeway of utilizing Soft-max is the output probabilities range. The range will be from 0 to 1, and the whole of the considerable number of probabilities will be equivalent to one.

### DATASET OF MANGO LEAVES COLLECTED IN OUR UNIVERSITY



Figure 2: Healthy dataset collected in KL (Deemed to be university)



**Figure 3:** Powdery Mildew dataset collected in KL (Deemed to be university)

Figures 2,3,4,5: The training and testing dataset of 800 pictures of mango leaves were collected in our university. In these pictures we can see the infected leaves and healthy leaves.

**Table 1:** Layer information and parameters of CNN

Layer(type)	Function	Output Shape
conv2d_1 (Conv2D)	Convolution	8X126X126
activation_1 (Activation)	Activation	8X126X126
conv2d_2 (Conv2D)	Convolution	8X124X124
max_pooling2d_1 (MaxPooling2)	MaxPooling	8X62X62
conv2d_3 (Conv2D)	Convolution	16X60X60
conv2d_4 (Conv2D)	Convolution	16X58X58
max_pooling2d_2 (MaxPooling2)	MaxPooling	16X29X29
conv2d_5 (Conv2D)	Convolution	32X27X27
conv2d_6 (Conv2D)	Convolution	32X25X25
max_pooling2d_3 (MaxPooling2)	MaxPooling	32X12X12
conv2d_7 (Conv2D)	Convolution	64X10X10
conv2d_8 (Conv2D)	Convolution	64X8X8
conv2d_9 (Conv2D)	Convolution	64X6X6
conv2d_10 (Conv2D)	Convolution	64X4X4
max_pooling2d_4 (MaxPooling2)	MaxPooling	64X2X2

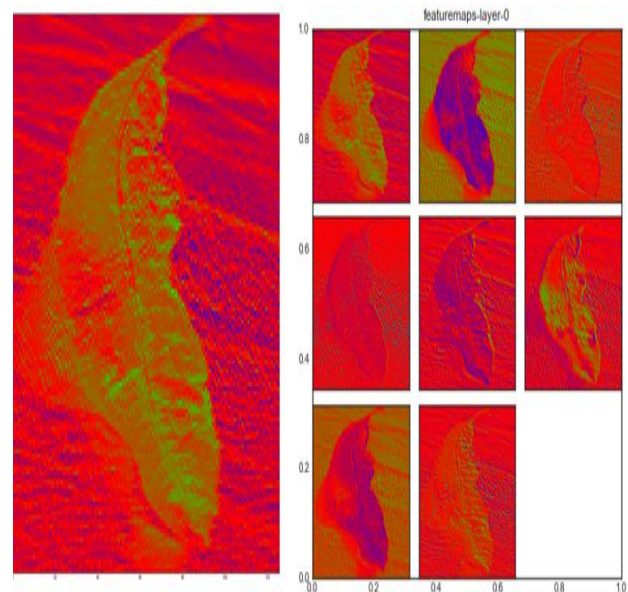
Table 1 clarifies the number of channels we have utilized for each layer processing and how the image layers are differing by separating every one of the highlights of the picture. Numerically we have completed ten times of convolution which is two dimensional, max-pooling of four times and an activation function. The order of processing an image is convolution, activation function which includes Re-Lu function followed by max-pooling.

**Table 2:** Table for recognition rates

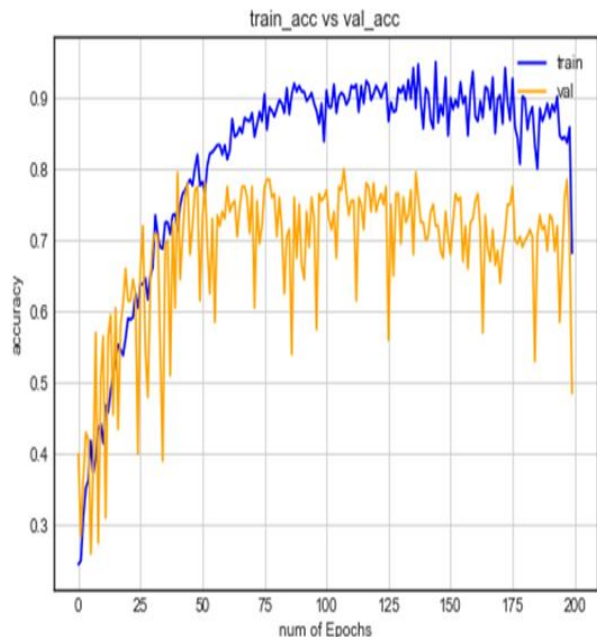
Classifier	Recognition Rates (%)					
	Batch-I Training		Batch-II Training		Batch-III Training	
	Testing with		Testing with		Testing with	
	Same data	Different data	Same data	Different data	Same data	Different data
ANN	76.39	74.26	78.47	75.29	79.13	77.42
Deep ANN	84.32	82.16	86.92	83.49	88.6	84.58
Our Proposed CNN(Single stride)	96.72	91.09	93.24	90.62	97.35	93.22

Table 2 clarifies the order of three classes i.e. ANN, deep ANN, our proposed structure of CNN. For finding out the recognition rates percentage of the diseased leaves, we had performed three batches of the data for three different diseases involved in mango leaves. We tested the information with our preparation data and different data. The accuracies obtained are mentioned in the table accordingly. We had got the correctness's of ANN and profound ANN from the online sources.

**4. RESULTS**



**Figure 4:** a) Feature maps



**Figure 4 :** b) Graphical representation of training accuracy vs epochs

Figure 4 a) The feature maps extricate the picture data; the feature map is the yield of one channel applied to the past layer. A given channel is drawn over the whole past layer, moved each pixel in turn. Each position brings about the enactment of the neuron and the yield is gathered in the element map. b) We had drawn the graph for the number of epoch's vs accuracy for training and testing of the data.



**Figure 5:** Accuracy obtained for the infected leaves

Figure 5s shows the exactness acquired for each leaf we had tested a portion of the leaves got 100% precision, some got 90% precision. Anthracnose, powdery mildew, bacterial canker are the recognized sicknesses their precision normal is 96.72%. The mismatch classification has happened for just a couple of leaves. The acquired after effect of our own proposed CNN model experiment for the leaves is appeared in the figure above.

### 5.CONCLUSION

This paper proposes the detection of diseases for the infected mango leaves. We selected the deep learning approach of our proposed design CNN which gives the average accuracy of 96.72% for batch1 and 97.35% for batch3 data. This deep learning technique extracts all features of the images and identifies the diseases of bacterial canker, anthracnose, and powdery mildew. The model used in CNN is single stride. A total of 800 pictures were taken and they were trained and tested 3 times like batch1, batch2, batch3. The results of recognition rates are mentioned in the table. 150 pictures are pre-trained for the system, 50 are tested to know the accuracy. In our proposed model we used Re-Lu as the activation function, convolution and max-pooling to extract the information regarding input image, dropout layers and dense layers are used to determine the output through the softmax layer. Our proposed model mainly helps the farmers to identify the disease that occurred to the leaves of the mango and can eradicate them in the initial state itself. So, the loss of harvesting crops for farmers can be reduced. Our project is completely a real-time approach mainly for the benefit of the farmers.

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