

# Tomato Leaf Diseases Identification Using CNN

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## ABSTRACT

Crop diseases are a key danger for food security, but their speedy identification still difficult in many portions of the world because of the lack of the essential infrastructure. The mixture of increasing worldwide smartphone dispersion and current advances in computer vision made conceivable by deep learning has cemented the way for web-based application for disease identification.

Using a public dataset of 20000 images of infected and healthy Tomato leaves collected under controlled conditions, we trained a deep convolutional neural network to identify 9 diseases and 1 healthy.

The trained model achieved an accuracy of 88.34% on a held-out test set, demonstrating the feasibility of this approach. Overall, the approach of training deep learning models on increasingly large and publicly available image datasets presents a clear path toward web application for crop disease diagnosis on a massive global scale.

**Keywords:** Deep Learning, Tomato Leaves, Inception v3, Detection, CNN

## 1. INTRODUCTION

Since the dawn of time, humans were depending on edible plants to survive, our ancestors would travel for long distances searching for food, no surprise that the first human civilizations began after the invention of agriculture, without crops will be impossible for humanity to survive. Modern technologies have given human society the ability to produce enough food to meet the demand of more than 7 billion people. However, food security still threatened by many factors including plant diseases, see (Strange and Scott, 2005)

Plant diseases are major threats for smallholder farmers, whose depend on healthy crops to survive, and about 80% of the agricultural production in the developing world is generated by them. see (UNEP, 2013), Identifying a disease correctly when it first appears is a crucial step for efficient disease management, traditional approaches to identify diseases is done by visiting local plant clinics.

But recent development in smartphones and computer vision would make their advanced HD cameras very interesting tool to identify diseases. It is widely estimated that there will be between 5 and 6 billion smartphones and also systems on the globe by 2020. At the end of 2015, already 69% of the world's population had access to mobile broadband coverage.

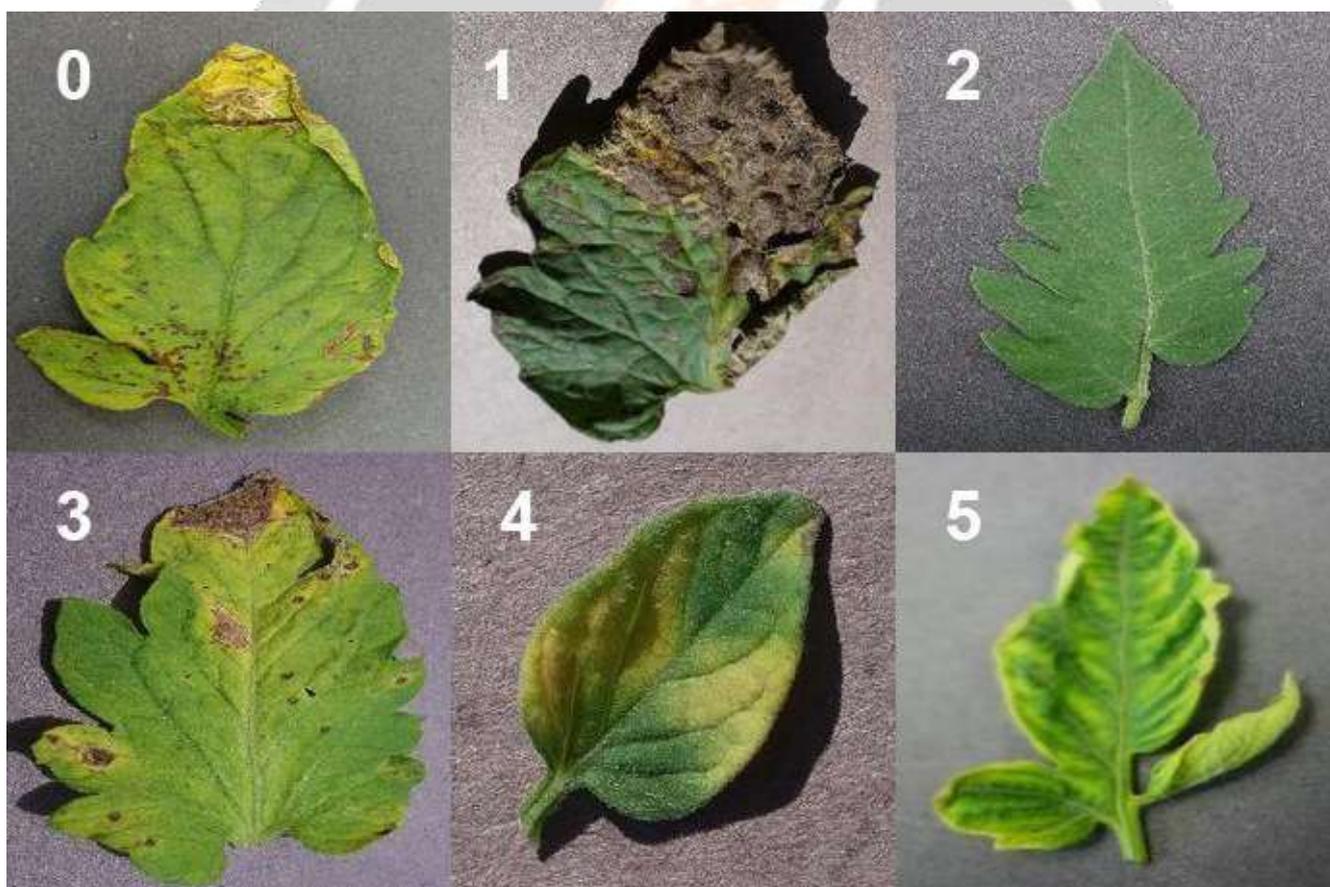
Significant impacts in image recognition were felt from 2011 to 2012. Although CNNs trained by backpropagation had been around for decades, and GPU implementations of NNs for years, including CNNs, fast implementations of CNNs with max-pooling on GPUs in the style of Ciresan and colleagues were needed to progress on computer vision. In 2011, this approach achieved for the first time superhuman performance in a visual pattern recognition contest. Also in 2011, it won the ICDAR Chinese handwriting

contest, and in May 2012, it won the ISBI image segmentation contest. Until 2011, CNNs did not play a major role at computer vision conferences, but in June 2012, a paper by Ciresan et al. at the leading conference CVPR showed how max-pooling CNNs on GPU can dramatically improve many vision benchmark records. In October 2012, a similar system by Krizhevsky et al. won the large-scale ImageNet competition by a significant margin over shallow machine learning methods. In November 2012, Ciresan et al.'s system also won the ICPR contest on analysis of large medical images for cancer detection, and in the following year also the MICCAI Grand Challenge on the same topic. In 2013 and 2014, the error rate on the ImageNet task using deep learning was further reduced, following a similar trend in large-scale speech recognition. The Wolfram Image Identification project publicized these improvements. Some researchers assess that the October 2012 ImageNet victory anchored the start of a "deep learning revolution" that has transformed the AI industry.

So here, using state of the art deep learning techniques, we demonstrated the feasibility of our approach by using a public dataset of 10000 images for healthy and infected Tomato leaves, to produce a model that can be used in smartphones applications to identify 9 types of Tomato leaf diseases, with an accuracy of 88.34% on a held-out test set.

## 2. STUDY OBJECTIVES

- 1- Demonstrating the feasibility of using deep convolutional neural networks to classify plant diseases.
- 2- Developing a model that can be used by developer to create web application to detect plant diseases.



**Figure 1:** Dataset Samples

## 3.DATASET SAMPLES

We extracted our dataset from the well-known Plant village dataset, which contains nearly 50,000 images of 14 crop species and 26 diseases. We choose to work with 20,000 images on Tomato leaves; our dataset contains samples for 9 types of Tomato diseases in addition to healthy leaves, 10 classes in total as follow[16]:

class (0): Bacterial Spot.

class (1): Early Blight.

class (2): Healthy.

class (3): Septorial Leaf Spot.

class (4): Leaf mold.

class (5): Yellow Leaf Curl Virus.

The images were resized into 224×224 for faster computations but without compromising the quality of the data.

## **4.NETWORKS: AN INTRODUCTION**

### **4.1.THE ARTIFICIAL CONVOLUTIONAL NEURAL NETWORK**

In machine learning, a Convolutional Neural Network (CNN, or ConvNet) is a class of deep, feed-forward artificial neural networks, most commonly applied to analyzing visual imagery.

CNNs use a variation of multilayer perceptrons designed to require minimal preprocessing. They are also known as shift invariant or Space Invariant Artificial Neural Networks (SIANN), based on their shared-weights architecture and translation invariance characteristics.

Convolutional networks were inspired by biological processes in that the connectivity pattern between neurons resembles the organization of the animal visual cortex. Individual cortical neurons respond to stimuli only in a restricted region of the visual field known as the receptive field. The receptive fields of different neurons partially overlap such that they cover the entire visual field.

CNNs use relatively little pre-processing compared to other image classification algorithms. This means that the network learns the filters that in traditional algorithms were hand-engineered. This independence from prior knowledge and human effort in feature design is a major advantage.

They have applications in image and video recognition, recommender systems and natural language processing.

### **4.2 Design**

A CNN consists of an input and an output layer, as well as multiple hidden layers. The hidden layers of a CNN typically consist of convolutional layers, pooling layers, fully connected layers and normalization layers [1-5].

Description of the process as a convolution in neural networks is by convention. Mathematically it is a cross-correlation rather than a convolution. This only has significance for the indices in the matrix, and thus which weights are placed at which index.

### **4.3 Convolutional**

Convolutional layers apply a convolution operation to the input, passing the result to the next layer. The convolution emulates the response of an individual neuron to visual stimuli.

Each convolutional neuron processes data only for its receptive field.

Although fully connected feedforward neural networks can be used to learn features as well as classify data, it is not practical to apply this architecture to images. A very high number of neurons would be necessary, even in shallow (opposite of deep) architecture, due to the very large input sizes associated with images, where each pixel is a relevant variable. For instance, a fully connected layer for a (small) image of size 100 x 100 has 10000 weights for each neuron in the second layer. The convolution operation brings a solution to this problem as it reduces the number of free parameters, allowing the network to be deeper with fewer parameters. For instance, regardless of image size, tiling regions of size 5 x 5, each with the same shared weights, requires only 25 learnable parameters. In this way, it resolves the vanishing or exploding gradients problem in training traditional multi-layer neural networks with many layers by using backpropagation.

### **4.4 Pooling**

Convolutional networks may include local or global pooling layers [8,9,10], which combine the outputs of neuron clusters at one layer into a single neuron in the next layer. For example, max pooling uses the maximum value from each of a cluster of neurons at the prior layer. Another example is average pooling, which uses the average value from each of a cluster of neurons at the prior layer[11-15].

#### 4.5 Fully Connected

Fully connected layers connect every neuron in one layer to every neuron in another layer. It is in principle the same as the traditional multi-layer perceptron neural network (MLP)

#### 4.6 Receptive Field

In neural networks, each neuron receives input from some number of locations in the previous layer. In a fully connected layer, each neuron receives input from every element of the previous layer. In a convolutional layer, neurons receive input from only a restricted subarea of the previous layer. Typically the subarea is of a square shape (e.g., size 5 by 5). The input area of a neuron is called its receptive field. So, in a fully connected layer, the receptive field is the entire previous layer. In a convolutional layer, the receptive area is smaller than the entire previous layer.

#### 4.7 Weights

Each neuron in a neural network computes an output value by applying some function to the input values coming from the receptive field in the previous layer. The function that is applied to the input values is specified by a vector of weights and a bias (typically real numbers). Learning in a neural network progresses by making incremental adjustments to the biases and weights. The vector of weights and the bias are called a filter and represents some feature of the input (e.g., a particular shape). A distinguishing feature of CNNs is that many neurons share the same filter. This reduces memory footprint because a single bias and a single vector of weights is used across all receptive fields sharing that filter, rather than each receptive field having its own bias and vector of weights.

### 5.METHODS

We experimented with two types of images to see how the model work and what exactly it learns, first we take the image as it is infected, and then we experimented with the another leaf which is healthy as an input. We have used compile method to configure the model, fit method to train the model with the trained dataset, plot

Layer (type)	Output Shape	Param #
conv2d_1 (Conv2D)	(None, 148, 148, 32)	896
max_pooling2d_1 (MaxPooling2)	(None, 74, 74, 32)	0
conv2d_2 (Conv2D)	(None, 72, 72, 64)	18496
max_pooling2d_2 (MaxPooling2)	(None, 36, 36, 64)	0
conv2d_3 (Conv2D)	(None, 34, 34, 128)	73856
max_pooling2d_3 (MaxPooling2)	(None, 17, 17, 128)	0
conv2d_4 (Conv2D)	(None, 15, 15, 256)	295168
max_pooling2d_4 (MaxPooling2)	(None, 7, 7, 256)	0
Activation (activation)	(None, 5,5,192)	0
Mixed (Concatenate)	(None, 5,5,2048)	0
Flatten (Flatten)	(None, 51200)	0
dense_2 (Dense)	(None, 10)	51210

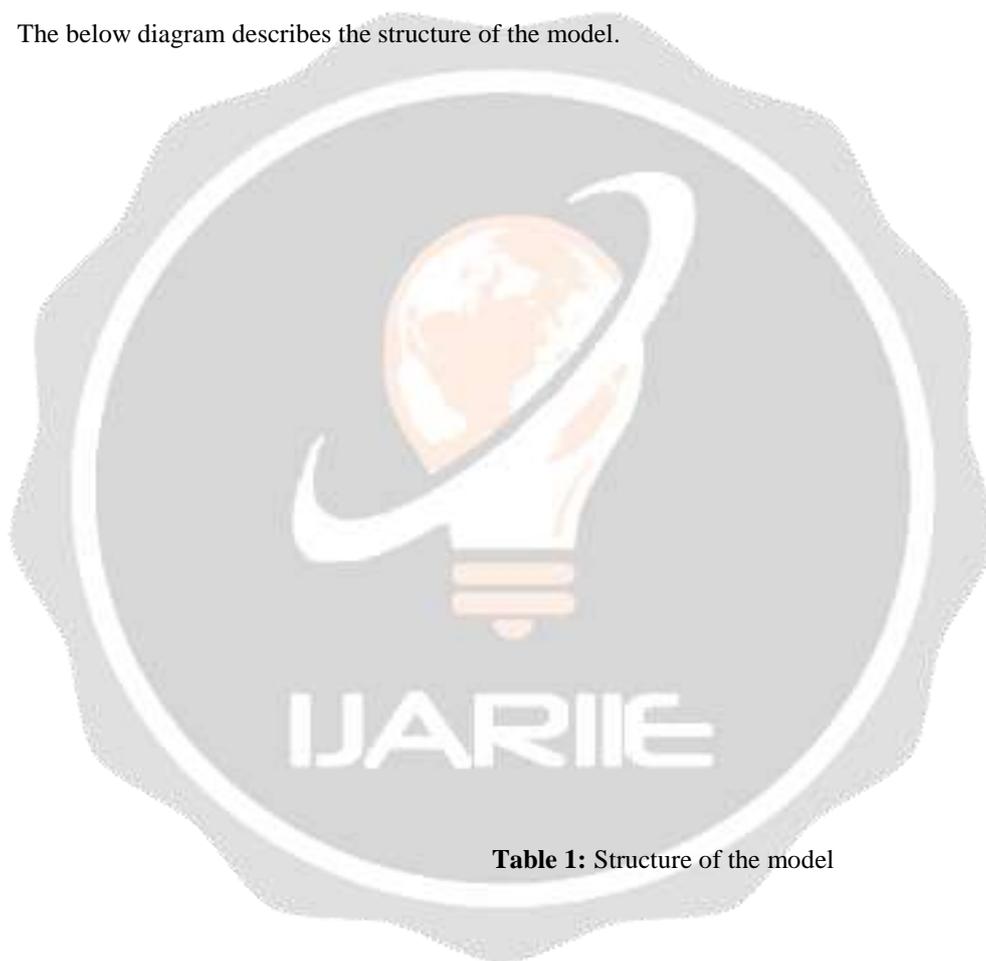
method to plot the graphs of accuracy and loss of training and testing, flatten method to remove the last layer

and adding the dense layer at last. In order to import the model, we are using inception method by passing the various arguments.

## 6.MODEL

We have used inception v3 which was a pretrained model for this experiment. Our model takes raw images as an input, so we used Convolutional Neural Networks (CNNs) to extract features.

- The model (features extraction), consist of Convolutional layers with ReLu activation function and batch normalization layer which is used for avoiding over fitting, each followed by Max Pooling layer.
- the flatten layer contains the dense layer to our approach, it has both softmax activation function and 10 outputs representing 10 classes.
- The below diagram describes the structure of the model.



**Table 1:** Structure of the model

## 7.DATA VISUALISATION

To see how the model works and what exactly learns we choose to visualize intermediate activations that consists of displaying the feature maps that are output by various convolution and pooling layers in a network, given a certain input (the output of a layer is often called its activation, the output of the activation function).

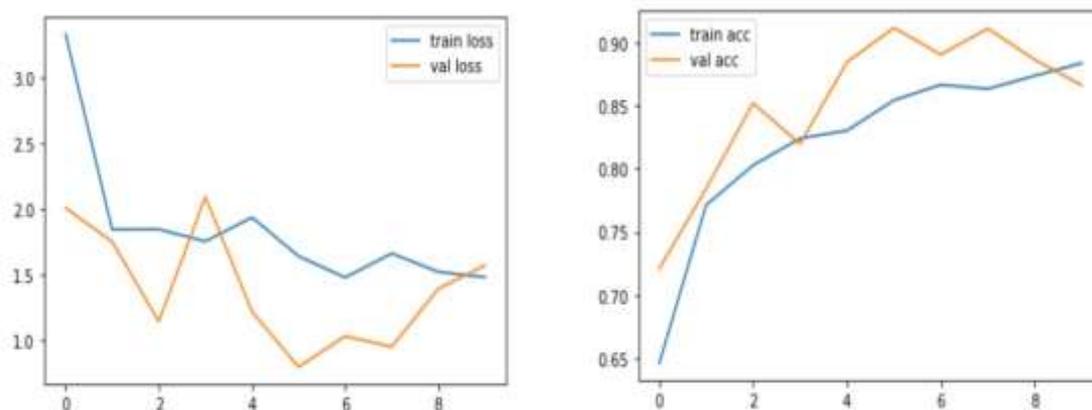
This gives a view into how an input is decomposed into the different filters learned by the network .

The image is taken as input image. The output of this is the type of the disease the leaf is.



**Figure 7.1: Input Image**

The graphs of the Accuracy and loss for training and testing dataset are shown below.

**Graph 1: Loss Graph****Graph 2: Accuracy Graph**

## 8. Conclusion

In this paper, a classification method of tomato leaf diseases based on deep learning was introduced to identify and classify ten common tomato leaf diseases: Spot blight, Late blight and Yellow leaf curl etc. Based on the inception v3, leaf pictures (after data augmentation) of three different diseases were selected and prepared for batch learning and training. After about 10 epochs, the recognition accuracy of the ten diseases reached 88.3% in training set and 86.0% in test set, respectively.

The pictures were sorted out in advance and named uniformly in one folder, instead of making labels for each one by hand. At the same time, a small number of pictures are easy to obtain and handle in real practice, making training process more flexible and efficient [13]. It has certain practical value for the development of modern ecological agriculture. In addition, this method can be applied to many situations. For other crops with analogous disease features, it can also achieve good results with a certain amount of pictures collected and processed.

Of course, this paper may make some improvements.

the segmentation and localization of the disease area can be combined to make the effect more reliable and abundant. Finally, due to the lack of a relatively complete and high quality crop disease database, the classification of some rare diseases or species is still not ideal. How to establish a more comprehensive and high quality disease data system is still the most important part of crop disease prevention

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