Defect Identification and Classification of Tomato Leaf Using Convolutional Neural Network

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Abstract - Tomatoes are the most commonly grown crop globally, and they are used in almost every kitchen. India holds second place in the production of tomatoes. Due to the various kinds of diseases, the quantity and quality of tomato crop go down. Identifying the diseases in the earlier stage is very important and will help the farmers save the crop. The first initial step is pre-processing, for the Canny edge detection method is used for detecting the edges in the tomato leaves. The classification of tomato leaves is to be carried out by extracting the features like color, shape, and texture. Extracted features from segmented images are fed into classification. The convolutional neural network algorithm will be used, which will give a better accuracy to classify the diseases in the tomato leaves.

Keywords: Canny Edge Detection, Classification, Convolutional Neural Network

I. INTRODUCTION

India is a country where the agriculture sector employs the majority of the population. Tomatoes are the most widely consumed crop in India. Tomatoes contain the three most essential antioxidants: vitamin E, vitamin C, and beta-carotene. They are also high in potassium, a mineral that's important for good health. (Solanum Lycopersicum) can fill important for the plant's vascular framework, but the leaves can stay green. A contaminated plant stem seems brown in cross-area with yellowish material emerging from it. We proposed a clever strategy for recognizing sickness in tomato crops by breaking down leaves' photos in the examination paper. Ranchers would have the option to tackle their plant sickness discovery issues without pursuing down plant researchers. Hence, it will help them in relieving the plant's sickness instantly, in this way expanding the quality and amount of food crops delivered and helping them.

The tomato issues might be partitioned into microorganisms or organisms or helplessness development propensities, causing 16 infections, while creepy crawlies were causing five different sorts of illnesses. Ralstonia Solanacearum microorganisms cause a genuine type of Bacterial shrink. This bacterium can get by in the soil for a critical stretch and enter roots through typical wounds made during helper roots advancement or fake during creating or migrating or even bugs. High sogginess and high-temperature favor contamination headway.

By quickly duplicating inside the plant's water-directing tissue, the microbes fill it with ooze, which influences the plant's vascular framework, however the leaves can stay green. A contaminated plant stem seems brown in cross-area with yellowish material emerging from it. We proposed a clever strategy for recognizing sickness in tomato crops by breaking down leaves' photos in the examination paper. Ranchers would have the option to tackle their plant sickness discovery issues without pursuing down plant researchers. Hence, it will help them in relieving the plant's sickness instantly, in this way expanding the quality and amount of food crops delivered and helping them.

The methodology alludes to the most well-known tomato infections, for example, Bacterial leaf spot, Septoria leaf spot, and Yellow Leaf Curl. Any leaf picture might be ordered into one of the infection still up in the air to be steady. The evaluation data set is a subset of Plant Village, a vault with 54,306 photographs of 14 yields invaded with the vermin. Around 18160 photographs of tomato leaf
sicknesses are remembered for this subset. Information assortment, pre-preparing, and arrangement are the three fundamental stages in the proposed approach. As referenced before, the photographs used to apply the proposed procedure were gotten from a freely accessible dataset called Plant Village. We built a Convolution Neural Network model to classify the images after downloading the dataset. For a pre-trained model, the model's performance was evaluated using different parameters such as training accuracy, validation accuracy, testing accuracy, and the number of trainable and trainable parameters.

II. LITERATURE SURVEY

Machine Learning are applied in various fields, anyway incorporate planning remaining parts the standard issue. With the advancement of a significant neural association, promising results are available for plant pathology without troublesome segment planning. Significant neural associations through and through increase picture gathering accuracy. This part gives distinctive significant learning techniques used by experts in plant ailment ID. Mohanty et al., [10] used AlexNet to get ready to arrange plant diseases that were not seen already. Model precision was liberally diminished while testing picture conditions are not equivalent to planning pictures. To a great extent, the disease appears on the leaves' upper sides, sometimes, the leaves' lower sides. Rangarajan et al., [14] arranged both AlexNet and VGG16net with least bundle size, weight, and tendency learning rate as hyper-limits. Accuracy is antagonistically related with least bunch size by virtue of VGG16net. Convolution and pooling layers together stacked in a module and applied to GoogleNet plan as Inception V4 for estimation decline. Too et al., [19] applied burdens pre-arranged on ImageNet to this designing typical pooling layer of 8 × 8 for changing. DenseNets obtains a test accuracy score of 99.75% for the 30th age, beating the models' rest. DenseNets is, thusly, a good designing for the task of plant picture based disease identification. All past layers' segment maps are used as commitments to each layer, and its component maps are used as commitments to each resulting layer. DenseNets have various convincing benefits, including the end of the disappearing inclination issue, further developed element proliferation, highlight reuse, and a critical decrease in the quantity of parameters. On four exceptionally cutthroat article acknowledgment benchmark undertakings, the writer tried his proposed engineering (CIFAR-10, CIFAR-100, SVHN, and ImageNet). On the majority of them, DenseNets achieve extensive updates over the bleeding edge while requiring less estimation to achieve high capability. Caffe outfits media specialists and specialists with an unblemished and adaptable framework for state of the art significant learning estimations and an arrangement of reference models. The construction is a BSD-approved C++ library with Python and MATLAB ties for getting ready and passing on comprehensively valuable convolutional neural associations and other significant models gainfully on product plans. Caffe fits industry and web scale media needs by CUDA GPU computation, planning in excess of 40 million pictures each day on a single K40 or Titan GPU (= 2.5 ms per picture). By disengaging model depiction from authentic execution, Caffe licenses experimentation and steady trading among stages for effortlessness of progress and sending from prototyping machines to cloud conditions. Caffe is stayed aware of and made by the Berkeley Vision and Learning Center (BVLC) to help working neighborhood givers on GitHub. It powers advancing examination projects, gigantic degree mechanical applications, and startup models in vision, talk, and media proposed by S., Long, et al., [7]. A CNN with nearby differentiation standardization layer is intended for parallel order with ReLu as initiation work Yusuke Kawasaki et al., [8]. The creator presents a novel plant sickness discovery framework dependent on convolutional neural organizations (CNN). Utilizing just preparing pictures, CNN can naturally obtain the essential highlights for order and accomplish high arrangement performance. The creator utilized an aggregate of 800 cucumber leaf pictures to prepare CNN utilizing our inventive strategies. Under the 4-overlay cross-approval procedure, the proposed CNN-based system (which likewise broadens the preparation dataset by creating extra pictures) accomplishes a normal precision of 94.9 % in grouping cucumbers into two ordinary sickness classes non-unhealthy class. AlexNet and GoogleNet are ready and adapted to the portrayal of sickness regions and incidental effects [2]. On the greater part of them, DenseNets achieve impressive updates over the bleeding edge while requiring less computation to achieve high capability. Caffe outfits media specialists and specialists with a perfect and adaptable framework for state of the art significant learning estimations and a grouping of reference models. The construction is a BSD-approved C++ library with Python and MATLAB ties for getting ready and passing on comprehensively valuable convolutional neural associations and other significant models gainfully on product plans. 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Execution of various CNN for plant disease conspicuous verification depends upon various segments: availability of a foreordained number of remarked on; powerless depiction of disorder incidental effects, picture establishment and getting conditions; confined assortments in sickness signs [21].
III. PROPOSED METHODOLOGY

A. Dataset Collection

Pictures of Tomato sickness have been taken from the Plant Village dataset. The dataset incorporates more than 50,000 pictures of 14 yields, like tomatoes, potatoes, grapes, apples, corn, blueberry, raspberry, soybeans, squash, and strawberry. We chose tomato as our objective harvest. The tomato leaf infection pictures have been taken from the Plant Village storehouse.

The gained dataset comprises of around 10000 pictures having a place with 10 distinct classes. The dataset incorporates pictures of all significant sorts of leaf sicknesses that could influence the tomato crop. Every one of the downloaded pictures has a place with the RGB shading space naturally. The size of the relative multitude of pictures is 256×256, and the configuration is jpeg.

B. Data Pre-Processing

Pre-processing is an initial step in image processing. Resizing the original image and then those image edges are detected by using the canny edge detection method.

1. Resizing

The pictures in the dataset were resized to 60×60 goal to accelerate the preparation cycle and make the model preparing computationally practical.

2. Edge Detection

The Canny edge recognition technique is utilized for edge identification. The Canny edge indicator is an edge recognition administrator that utilizes a multi-stage calculation to recognize a wide scope of edges in pictures. At first, perusing the first picture followed by changing it over to a grayscale picture at last edges are distinguished utilizing the watchful edge discovery technique.

C. Convolutional Neural Networks

Convolutional neural organizations (CNNs) are a type of profound neural organization (DNN) motivated by the human visual framework and intended to get pictures. A CNN is comprised of an assortment of convolutional and spatial pooling layers that are on the other hand stacked. The convolutional layer utilizes direct convolutional channels and nonlinear enactment capacities to remove highlight maps (e.g., rectifier, sigmoid, tanh, and so forth) Neighborhood highlights got from spatially nearby pixels are amassed utilizing the spatial pooling layer. This layer is regularly used to expand the target's protection from minor disfigurements. Convolution, ReLU Layer, Pooling, Completely Connected, Flatten, and Normalization are a portion of the layers in the CNN model. Every part is
alluded to as a capacity or channel. CNN utilizes the weight lattice to get novel provisions from the info picture without losing data about its spatial course of action.

1. Convolution Layer

The essential and first layer of the CNN design is the convolution layer. It is utilized to separate characters from a picture input. By taking in picture highlights from input information, the convolutional layer protects the connection between pixels. Two information picture lattices and a 3×3 part channel are utilized in the numerical interaction. The image pixel upsides of 5×5 and the 3×3 convolution channel. In the wake of completing the duplication of the relating framework, the provisions. The CNN increases the outcome by the all out number of pixels, then, at that point makes a guide and places the channel's worth. The element is then moved to each and every area in the picture, the lattice yield is acquired, and the cycle is rehearsed for different channels. This layer smoothes the channel on the image in each conceivable spot.

2. ReLu Layer

Rectified Linear Unit (ReLu) is an abbreviation for Rectified Linear Unit. It is the most regularly utilized initiation include in neural organization covered up layers. The separated picture's negative worth will be supplanted with zeros in this layer is given in condition (1).

\[ A(x) = \max(0, x) \]  

3. Pooling Layer

This layer reduces the image's size by extracting the full value from the filtered image and converting it to a matrix. It also holds overfitting at control.

4. Fully Connected Layer

A bunch of totally associated layers make up a completely associated neural organization. In completely associated networks, hubs are regularly alluded to as “neurons” subsequently, totally associated organizations would be alluded to as “neural organizations”. The information measurement decides the capacity of the totally associated layer from \( R^m \) to \( R^n \). This component vector is utilized for characterization, relapse, and other yield structures after the organizations have been prepared. The FC layer stores composite and accumulated information from all pertinent Conv layers. We should take a gander at the numerical type of a totally associated network in more detail.

\[ y_i = (w_1 \times 1 + \cdots + w_m \times m) \]  

Let \( x \in R^m \) reflect the input to a fully attached layer. Let \( y \) \( i \in R \) be the i-th output from the completely attached layer.
5. VGG NET 19

A 224*224 RGB picture is utilized as the contribution to the VGG-based ConvNet. The pre-handling layer takes away the mean picture estimates estimated over the whole Tomato leaves preparing set from the RGB picture with pixel estimates in the scope of 0–255.

After pre-handling, the info pictures are gone through these weight layers. A pile of convolution layers is utilized to move the preparation pictures around. In the VGG16 design, there are an aggregate of 13 convolutional layers and three totally associated layers. Rather than having enormous channels, VGG has more modest channels (3*3) with more profundity. It presently has a similar amazing responsive field as though only one 7 x 7 convolutional layer were utilized. Another VGGNet variation has 19 weight layers, with 16 convolutional layers and three totally connected layers, and five pooling layers. VGGNet has two completely associated layers with 4096 channels each, trailed by another completely associated layer with 1000 channels to anticipate 1000 marks in the two varieties. Soft max layer is utilized for characterization purposes in the last completely associated layer.

**Walkthrough of the Architecture**

The first two layers are convolutional layers with 3*3 filters, and the first two layers use 64 filters, results in a volume of 224*224*64. 3*3 filters with a stride of 1 are used.

Following that, a pooling layer with a max-pool of 2*2 size and stride 2 was used to reduce the volume's height and width from 224*224*64 to 112*112*64.

After that, there are two more convolution layers with 128 filters. The new dimension is 112*112*128 as a result of this.

The volume is reduced to 56*56*128 after using the pooling layer. Two more 256-filter convolution layers are inserted, followed by a down sampling layer that shrinks the size to 28*28*256.

A max-pool layer divides two more stacks, each with three convolution layers.

The 7*7*512 volume is flattened into a Fully Connected (FC) layer with 4096 channels and a soft max output of 1000 classes after the final pooling layer.

**IV. EXPERIMENTAL RESULTS**

The proposed method has been tried on the Plant Village dataset. It contains roughly 18160 photographs of tomato leaf illnesses from ten distinct gatherings. 220 pictures were saved for research and 880 pictures were utilized for preparing out of the 1100 aggregate. Programmed information expansion methods were utilized to expand the dataset by self-assertively pivoting the pictures just barely of 20 degrees, level flipping, and vertical and flat moving of pictures. The improvement was finished with SGD and a misfortune capacity of unmitigated cross-entropy. The model was prepared for 30 ages and utilized a bunch size of 20. The underlying learning rate is set to 0.01 and afterward diminished by a factor of 0.3 when the disappointment arrives at a level. Early halting has additionally been utilized to screen legitimacy disappointment and stop the preparation stage as it rises.

More than 30 ages of preparing, the most elevated approval exactness of 94.8 percent was accomplished, with a high preparing precision of 99.3 percent. The approval precision was observed to be 94% by and large. This is a decent marker of how well the profound learning model characterized the information. The plots of train and test
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precision and misfortune against the ages give a visual portrayal of model combination and a sign of its speed. The model appears to have settled around 20 ages, and the measurements have not worked on fundamentally in the last 10 ages.

V. CONCLUSION

The agrarian area keeps on being one of India’s most significant areas, with most of the populace depending on it. The discovery of illnesses in these harvests is consequently basic to the economy’s turn of events. Tomatoes are a staple yield that is filled in enormous amounts. Thus, this paper expects to distinguish and recognize 10 unique illnesses in the tomato crop. To recognize tomato leaf sicknesses from the Plant Village dataset, the proposed approach utilizes a convolutional neural organization model. To order tomato leaf sicknesses into 10 unique gatherings, a basic convolutional neural organization with various layers was utilized.

REFERENCES