

Automatic Detection of Malaria Using Convolutional Neural Network

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Abstract: It's an infectious parasitic disease spread by mosquitoes that's caused by a parasite called (Plasmodium). This parasite infects and destroys red blood cells in the human body. Fever, anemia, spleen enlargement, headache, chills, and exhaustion are among the most common symptoms. Blood smears from individuals are still planned under the microscope by competent professionals and technicians for red blood cells contaminated with parasites, which is the classic approach for diagnosing malaria. The wait for a diagnosis is long. Deep learning algorithms have been used to infected human blood signals as a result, however the practical performance has so far been insufficient. The architecture of the convolutional neural network (CNN) was used in this paper to diagnose malaria illness using deep learning. , which includes 43,828 pictures of infected and uninfected blood cells were used to train on various weighted layers to assess the presence or absence of malaria in red blood cells from microscopic images collected of blood smears of infected human persons. A total accuracy of 96.87 was attained when the CNN model was tested. The proposed deep learning approach for detecting malaria was found to be effective based on the experimental results on the clinical data set.

Keywords: malaria ,deep learning , machine learning ,CNN

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1. Introduction :

Malaria is a contagious disease, and the World Health Organization has announced an outbreak of the disease in many countries. The cause of this disease is a virus known as Plasmodium, which lives in the mother mosquito's body and is injected into the blood of a healthy person as soon as a mosquito bites a person, and humans are often susceptible to infection with this disease according to World Health Organization reports, the number of Another interesting fact from these studies is that 272,000 people died of malaria[1][2], with children under the age of five being the most affected.. As a result, it is one of the leading causes of death for children around the world, particularly in African countries. Despite the fact that there are therapeutic medications to treat malaria, there is currently no viable vaccine, making it a very effective sector. [3] Malaria is a disease that can be passed from one person to another; therefore, it is vital to protect yourself. It is still a method to diagnose it early and hence prevent its spread. The importance of a correct and successful malaria diagnosis cannot be overstated. Only a skilled microscope can accomplish this. Hundreds of billions of microscopic images are examined by the microscope. Furthermore, the number of microscopes available is insufficient, resulting in a serious shortage in the fight against malaria, particularly in developing nations[3]When opposed

to qualitative microscopic diagnosis, deep learning detection of malaria is more objective and quantitative. As a result, there is a growing demand for computer-aided diagnosis, which is quick and accurate in diagnosing and performing health-related tasks. Large quantitative virus insights, annotated data sets for deep training, and learning models for disease identification have all advanced as a result of increasing uses of these automated algorithms in electronic records and imaging-based diagnosis[3][4].The second half of this study will discuss past research, the third will discuss the processes that occur prior to training, and the fourth will discuss the construction of the convolutional neural network.

2.Methods

This part will assist you in comprehending all of the pre-processing procedures employed in the research to record the model's various performances. This research makes use of a data set from the Kaggle website. It contains 43,827 photos of malaria parasite-infected and healthy cells. It was separated into 21900 infected cell images and 21900 infected cell images a picture of cells that haven't been infected. Figure (1) shows the structure of the suggested algorithm in a sequential manner

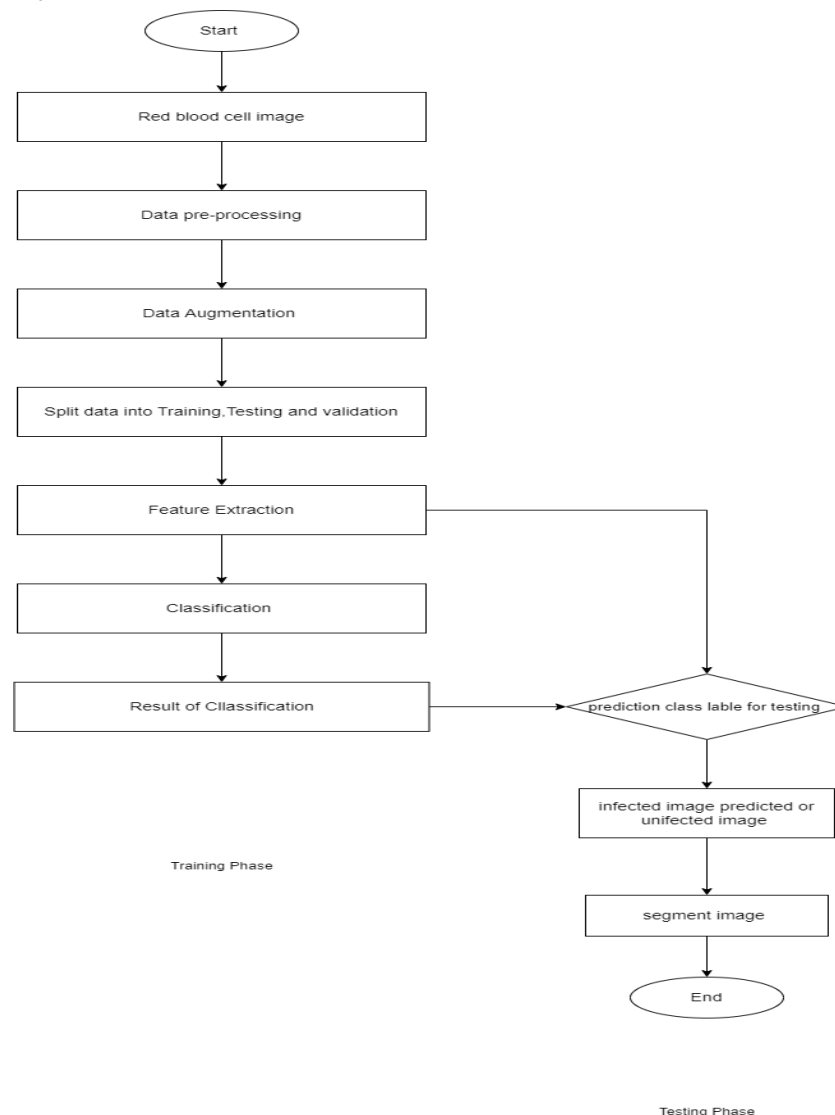


Figure .1: Block Diagram for Malaria Detection

2.1.Re-scaling

Although image scaling improves performance, it takes a long time to compute and consumes a large amount of memory. It's ideal to keep this trade-off between accuracy and computation. All of the photographs in this study were resized to [145 145]. As a result, the classification score increased and the processing time decreased. Because the neural network gets inputs of the same size, all photographs must be scaled to a specified size before being sent to Cnn. Large images, on the other hand, require more memory space as well as a larger neural network, which takes up more space and time. Today, it's clear that choosing a specific image size involves a trade-off between computational efficiency and accuracy figure (2) shows the image of blood cell and one healthy

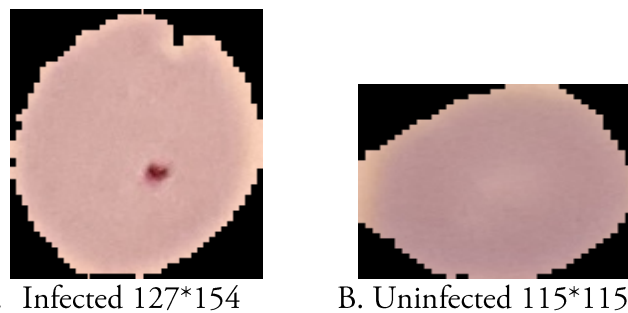


Figure.2.A. the a image of an infected red blood cell and uninfected one when the image size are different.

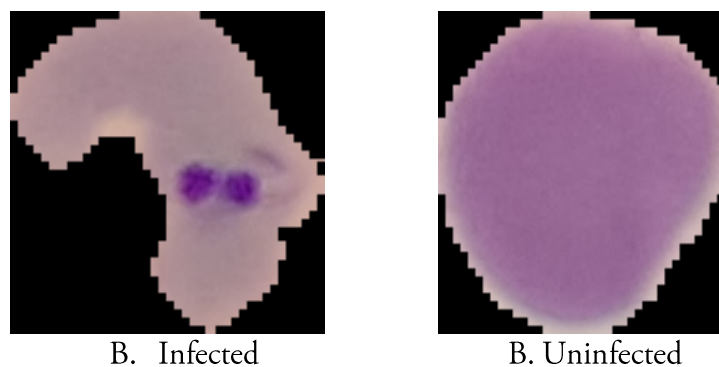


Figure.2.B. the a image of an infected red blood cell and uninfected one when the image size is standardized to 145*145

2.2.Data splitting

After the data is collected, it is separated into three groups: training, validation, and testing. Only 70% of the data set is being used for training. Validation had a percentage of 20% and a test of 10%.

3. CNN Architecture

CNN is a deep learning technique for classifying images that takes the image as input, assigns learnable weights, and outputs features or objects in the image that may be used to differentiate one from the other. The purpose of this ConvNet layer is to minimize the size of the photos to a manageable amount Build up a lot of hidden layers, but only one input and output layer. Multiple convolution layers and max pooling layers, two dense layers, one flattening layer, and the fully linked layer make up the CNN. This model is composed of four convolutional

layers, each of which contains the following filters: kernel size = (3,3), padding = same, stride= (2,2), activation ("relu"), and dropout layer = (0.3). At the input layer, images were transformed to $145 \times 145 \times 3$, with 3 denoting the channel number. The tensor was sequentially passed through (32,64,128,256) filters in each layer, starting with 32 in the first layer, and the results were satisfactory. Finally, the output neurons of the flattening layer were fed into the max pooling layer. We used a batch size of 32 and received a moderate training result after 180 epochs and 40 per epochs using the output from the dense layer as input to the activation function, which was a sigmoid activation function, in order to get the output probability of the classification task. figure 3 show Details of Layers in Deep CNN system design

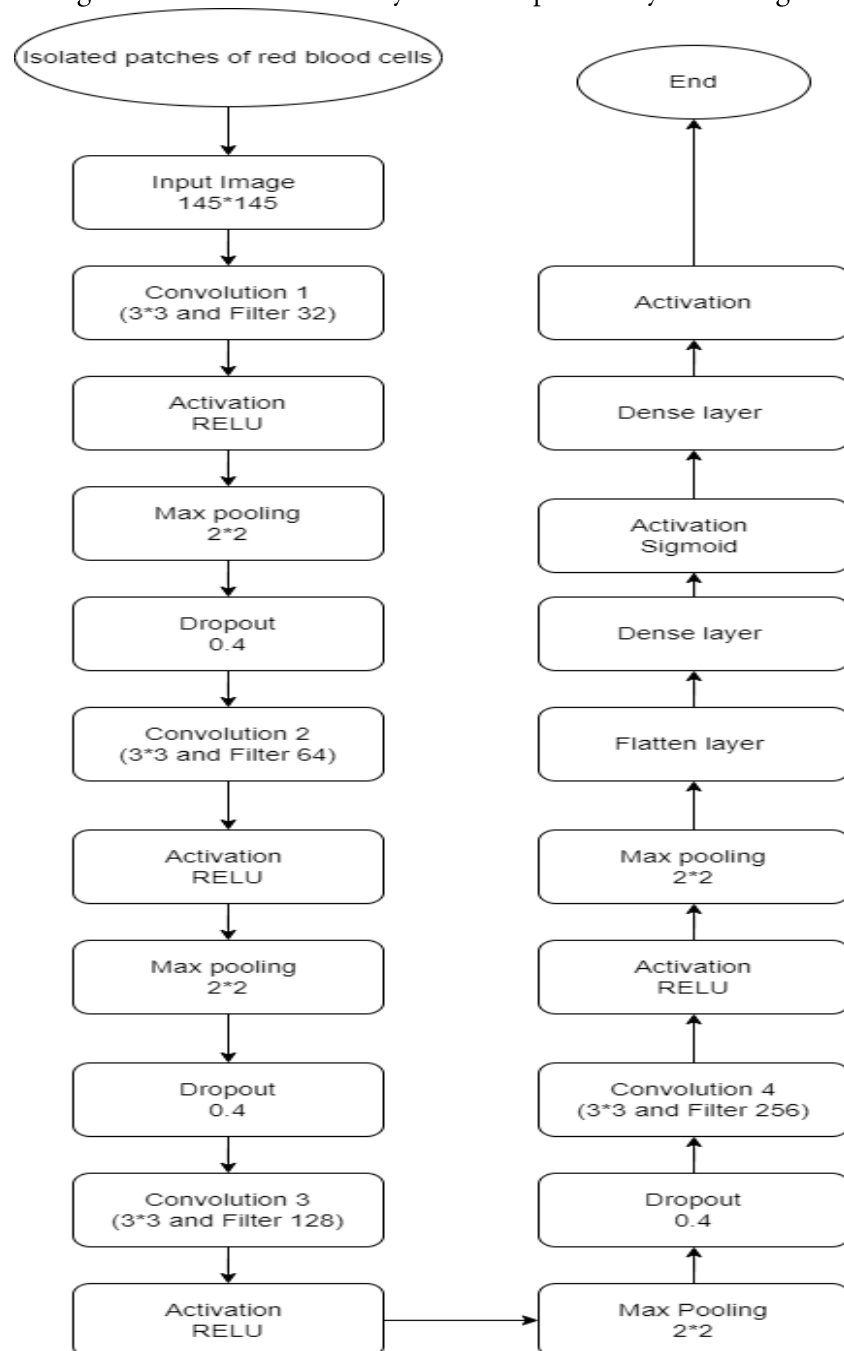


Figure 3. Convolutional Neural Network Architecture

3.1.convolution layer:

This is the first layer of the CNN architecture, which is used to extract picture properties. An picture is convoluted with various filters to perform actions such as sharpening and edge detection.

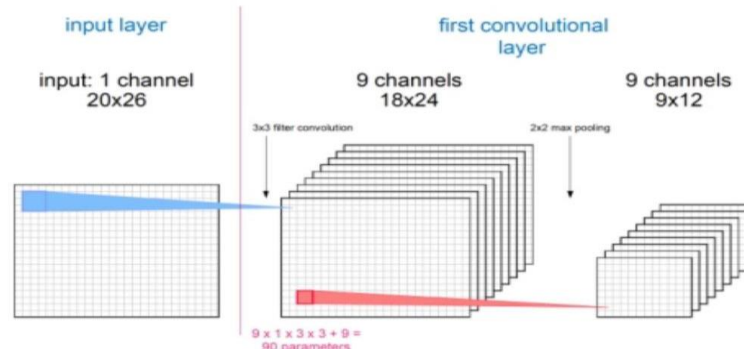


Fig.3.Example of a Convolution layer

3.2.Pooling layer:

When an image is too huge, this layer is used to train the algorithm to take a few numbers of parameters. There are three distinct types. The first is maximum pooling, which takes the largest element from a feature map into account. Another is Average pooling, which takes into account the average of all the components in a feature map, and Total pooling, which takes into account the sum of all the elements in a feature map.

3.3. ReLU Layer:

Rectified Linear Unit (ReLU) is an abbreviation for Rectified Linear Unit. It's an activation function that's applied to CNN neurons' outputs $y = \max(0, x)$ is the mathematical formula for the Rectified Linear Unit (0, x).

3.4 Fully Connected layer:

A fully connected (FC) layer is an input feature vector. The FC layer's function is to smooth high-level properties that can be learned by Convolutional layers and merge features. It sends the resulting flattened output to the output layer.

4.Evaluation Metrics

The following standard metrics are used to evaluate the proposed model: accuracy, sensitivity, specificity, precision, and classification error rate. The accuracy and sensitivity parameters are utilized as metrics in the experimental setting, and the following metrics are employed as satisficing:

4.1. **Accuracy** : calculates accuracy by adding true positives and true negatives to the total number of datasets Eq. 1

$$Accuracy = \frac{True\ positive(TP) + True\ Negative(TN)}{TP + TN + FP + FN}$$

4.2. **Recall** : This is the ratio of real positive examples divided by the total number of positive cases, as calculated using the equation in Eq.2

$$Sensitivity = \frac{True\ positive(TP)}{Actual\ Positive(TP + FN)}$$

4.3. **Specificity** : The ratio of real negative to actual negative, as given by equation in Eq.3 is used to calculate specificity.

$$Specificity = \frac{True\ Negative(TN)}{Actual\ Negative(FP + TN)}$$

4.4. **Precision** : Divide the genuine positive by the projected positive using the equation in Eq.4 to find precision.

$$Precision = \frac{True\ positive(TP)}{Predicted\ Positive(TP + FP)}$$

4.5. **F1- Measure** : It's the weighted average of accuracy and recall that's been resolved. It is used to statistically assess an individual's performance. The F1 score, as previously stated, is an evaluation metric that is used to express the performance of a machine learning model by providing aggregate information on the model's accuracy and recall. Eq.5 find f1

$$F1 = 2 \cdot \frac{Precision \cdot recall}{Precision + recall}$$

Table 1. Analysis of parameters for malaria detection

Parameter	Accuracy	Recall	Specificity	Precision	F1
Values	96.89	97.77	96	95.94	96.85

5. Result:

The main goal of this project is to create a useful deep learning model for predicting malaria illness. The data set used in Figure 1 in our proposed work is made up of 43,827 pictures of infected and uninfected blood cells. The data was imported from the website kaggle.com, as previously stated. The cnn model appears in the drawing where the images are classified as healthy or unhealthy in the following graph figer 3 , the validation line and the training line meet, and the model produces accurate and high results

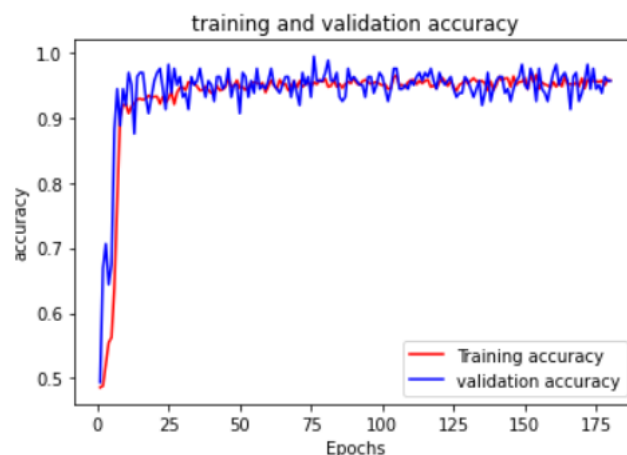


Fig.4. the accuracy graph

after which a confusion matrix is formed for each convolutional neural network for a better understanding. A two-dimensional matrix containing four attributes is used to measure machine learning classification. The confusion matrix is used to generate the precision values for true positive (TP), false positive (FP), false negative, true negative (FN), and true negative (TN). Accuracy, recall, and the F-measure are all important factors to consider.

		Actual	
		(Positive)	(Negative)
Predicted	(Positive)	True Positive (TP)	False Positive (FP)
	(Negative)	False Negative (FN)	True Negative (TN)

Fig 4. The confusion matrix

6. conclusion

Traditional methods for detecting malaria entail collecting samples and analyzing cell proliferation, which takes time and effort. In the proposed work, a deep learning model was developed to predict malaria with high accuracy and in a short amount of time. A data collection from the Kaggle.com website was used to develop and train the model by utilizing the CNN convolutional network's characteristics, and an accuracy of 96.87 was attained, demonstrating that this model is superior to other models in the future. CNN can be used to detect various disorders such as pneumonia and breast cancer.

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