

A Novel DIP Method based Detection of Malaria Parasites

¹Vidya V, ²Priyanka R, ³Shaik Mohammed Anjum Naaz, ⁴Shubh Laxmi, ⁵Bhgya Shree R Hawde

^{1,2}Asst. Professor and ^{3,4,5}Students

Department of Computer Science Engineering, Nagarjuna College Of Engineering & Technology,
Devanhalli, Bengaluru, Karnataka, India-562164.

Abstract— One of the most profoundly powerful and an astoundingly transmissible infection achieved by a blood parasite which has a spot with kind of Plasmodium is Malaria. Considering the customary and the general microscopy technique, which is "The Gold Standard" for intestinal sickness identifying has been found contrasting, in light of the fact that the method requires significant proportion of time and the outcomes are difficult to repeat as and when required. intestinal sickness is causing a veritable overall clinical issue, and the evaluated methodology is of high significance. The proposed work here presents a model that uses Image processing techniques and counts that are, definite, quick and cost-effective detecting of intestinal infection (malaria) by means of getting ready and testing over the gathered recolored blood smear images. Datasets including pictures of affected and non-affected erythrocytes are accumulated, preprocessed, immovably related features are isolated from the received images and finally, it is confirmed whether the test picture is infected or not taking into account the features that are gathered from them. A great deal of qualities depending upon the features are proposed, and the strategy proposed from the created database of the attributes on the erythrocytes tests is classified using a SVM and MSVM classifier.

IndexTerms - Intestinal sickness, Malaria, Parasite, Digital Image Processing, erythrocyte, grayscale image.

I. INTRODUCTION

Malaria is a disease caused due to the protozoan parasite called plasmodium and mainly responsible of transmission and infection of the earlier mentioned disease. Anopheles mosquitos have been found in around 400 different species, and in this 30 are malarial vectors. They bite during nightfall and first light. The scope of infusion relies upon the encompassing and nature where the individual lives. The encompassing must be very much kept up with appropriate inclusion of sewage section and neatness. Few the spices to name a few are, Plasmodium Falciparum, Plasmodium Malariae, Plasmodium vivax, and Plasmodium Ovale. The plasmodium parasite is carried to the human body using anopheles mosquitoes. RBC are affected by these parasites and they complete their life cycle using the hosting in RBC and finally rupture the RBC [11]. Once the RBC is ruptured, it increases the severity of infection of intestinal sickness which is also known as malaria and is an intense irresistible ailment brought about by a fringe blood parasite of the class plasmodium. As per the World Health Organization insights, in 2000, it was assessed that there were 262 million instances of malaria universally, prompting 839.000 people to die. But by 2015, it was evaluated that the quantity of intestinal sickness cases had diminished to 214 million, and the total number of deaths diminished to 438.000. Most of these deaths included youngsters from Sub-Saharan Africa. This is because of the way that, the natural conditions are appropriate for mosquitoes, not with-standing the poor financial conditions which makes access to human services and sickness, disease prevention troublesome [1].

To analyse malaria there are various methods out of which manual microscopy is viewed as "the gold standard". But due to the quantity of procedures required in manual evaluation, this demonstrative technique is tedious (prompting late determination) and inclined to human blunder (prompting mistaken finding), even in experienced hands like technicians and pathologists. As mentioned earlier, the manual methodology of determination is tedious and may prompt irregularity. Thus the proposed method once digitalized should be able to bring down the amount of time taken for screening the infection. This will improve the consistency in analysis [2]. Hence the proposed work investigates the utilization and use of image processing for distinguishing intestinal sickness parasites utilizing microscopic color images. A productive strategy is proposed for parasite recognition dependent on intensity and color features. Parasite location and detection is the main aim of this semi-automatic and diagnostic analysis. The proposed work is to develop a fully functional and user interactive application which can detect the presence of malaria in blood cells using microscopic images which does not require a skilled pathologist to test the disease, instead a person with average knowledge of computer can easily detect the infection degree. The proposed system is faster than the normal conventional methods of detecting the presence of malaria in blood.

The rest of this paper is organized as follows. Section II presents a discussion of related studies, Section III describes the scope of work and system requirement, Section IV system design and testing, Section V experimental results, and finally Section V concludes the paper.

II. REVIEW OF PREVIOUS WORK

It is always necessary to review the work of the researchers done previously to deduce out the problem statement and solution for it. This leads to the development of device which overcomes the problems posed by the earlier ones.

Ashwini Awchite et al., it presents a computerized system using image analyzation, hence eliminating the need for manual approach for locating, identifying and testing of the parasite species on thick blood [3]. Nicholas E. Ross et al., here a systematic method is proposed for the errors to be identified which are caused by an individual during the identification of the malaria parasite which are taken through the blood sample carried out using automated image processing thus supervising and checking the errors [4].

S. S. Savkare Moze et al., the system is developed using an algorithm for helping in finding the point by allocating near to a two side by side (parallel) planes and Otsu's threshold thus helping in getting the binary images of the cells, separation of the cells which are overlapping and finally it helps in the discovery of the infected and non-infected images with the study of SVM networks [5]. Hanung Adi Nugroho et al., here different methods like classification, ANN algorithm, and diafeature extraction methods are used. Segmentation and enhancement of image was developed earlier. The main aim here is to improve the accuracy of the result obtained, by developing a computerized system [6]. Deepali A. Ghate et al., here a programmed Detection of malarial parasites from blood pictures is presented. It checks the human blunder while distinguishing the nearness of intestinal sickness parasites in the blood test by utilizing Image acquired and diminishing procedures to identify malaria samples [7]. Pranati Rakshit et al., a novel strategy is built up that identifies all the existence phases of *P. Vivax* also, leukocytes from tiny microscopic of the thick blood smear. The parasite number tally by the proposed calculation additionally the proposed technique, parasite thickness can be assessed utilizing all the existence phases of malaria parasite, in contrast to condition of workmanship calculations, that can assess tally just from ring life stages [8] Ankit Sanadhya et al., areas of intrigue are classified in five classes: ring structure, early trophozoite (ET) structure,

late trophozoite (LT) structure, leukocyte and non-parasite. By decreasing various highlights utilizing head part examination (PCA), the exactness in the parasite tally diminishes. The calculation performs at its best when each of the 123 highlights are chosen for preparing, precision might be improved by utilizing more pictures for preparing,

by improving slide quality and improving picture obtaining technique. This work can be stretched out for identifying different Plasmodium species and their different life stages [9].

Najeed Ahmed Khan et al., Location of jungle fever parasite species and its life cycle stages requires an exceptionally prepared pathologist, as there exists countless variable, highlights making it difficult to recognize these parasites. Efficiency and exactness of proposed algorithm is improved as just the highlights of the contaminated cells are thought about [10]. J.Somasekar et al., The essential idea here is, at first transformation of the info shading blood picture to grayscale, and afterward ascertaining the scope of yth request of a grayscale image. Here right now Effective calculation, with the gamma equalization (GE) technique is executed whose principle rationale is to ensure the major structures of the obtained blood pictures which is contaminated by jungle fever [11].

Snehal Suryawanshi et al., This paper presents upgraded system for Malaria Parasite Detection, where cell division process consists of different features, for example, image banalization utilizing Poisson's dissemination based Minimum Error threshold, trailed by Morphological Opening with the output goal of refinement [12].

III. SCOPE OF PROPOSED WORK AND SYSTEM REQUIREMENT

The main aim of the proposed work is to finish detection of Malarial Parasites in Blood Using DIP are as per the following: Identifying the Challenges and openings in Malarial Parasites in Blood are detected Using Image Processing. Image pre-handling on the information Identification of Malarial Parasites in Blood Using Image Processing. Division and Analysis of Detection of Malarial Parasites in Blood Using Image Processing. Classification of the platelets. Identification of malarial parasites. This product recognizes the nearness of intestinal sickness in minuscule blood pictures. The software:

1. Is able carry out segmentation of the given blood image.
2. Segmented image is applied with image enhancement operations.
3. Reducing the presence of noise by converting the enhanced image to binary image.
4. Reducing the amount of data to be handled for analysis by extracting the features of the malarial parasites.
5. Negative and positive cases of malaria Classification is performed.

Programming Requirement Specification absolutely defines how the anticipated programming acts without unfurling how the product will perform it. The rudimentary goal of the prerequisite stage is to yield the product necessity specification that assigns the fringe execution of the anticipated programming. Programming necessity can be well-defined as a state of a capacity required by a client to take care of an issue or achieve a goal.

User Characteristic

Here the characteristics of the user is covered in three senses: machine translation system is handled by the end user; translation of the final product process also which may include for example post editing is also taken care by end user, the deployment of machine translation system by the organization.

Lab Technician

The blood images of the patients are uploaded to the software for detection of infection to be positive or negative is done by lab Technicians.

Interfaces

A periphery where two independent systems actually meet and an interaction or communication with each other. The various types of interfaces are:

1. User Interface – which consist of the keyboard, mouse, menus of a computer system. Communication with the operating system is done using the user interface. Also there is GUI (Graphical user interface)
2. Software Interface – This consist of the codes and programs in different languages which help the application to communicate with each other and other hardware devices.
3. Hardware Interface – It consists of devices like plugs, socket and wires which these hardware devices use to communicate with each other.

Hardware Interfaces

1. Processor: Any processor above 500 MHz
2. RAM: 4GB
3. Hard Disk: 5GB
4. Input Device: Standard keyboard and Mouse
5. Output Device: Monitor

Software Interfaces

1. Database: Local Database
2. Programming Language: MATLAB
3. Application Software: MATLAB R2016b

Functional Requirements

1. Input: The system is fed with train and test dataset which consist of 90 images for training the system. The blood samples in microscopic image form is the input to the software.
2. Processing: Software part of the system performs the following:
 - a. Image Processing
 - b. Removal of Image Noise
 - c. Image Segmentation
 - d. Image enhancement
 - e. Feature extraction
 - f. Classification
3. Output: it predicts if the output which is malaria is infected or not. It uses SVM or MSVM classifier to identify.

Non- Functional Requirments

1. Security
2. Availability
3. Portability
4. Scalability
5. Reliability

IV. SYSTEM DESIGN AND TESTING

The process of defining the architecture modules, interfaces, and data for system which is known as system design and used for specified requirement. It can be used for the integration of systematic theory for application of product development. There by working together with system analysis, system engineering and system architecture.

Architecture Design

Architecture structure and the dataset is gathered which is isolated into preparing and testing datasets. The prepared datasets incorporate the tiny pictures of platelets of different patients. The Microscopic picture is first pre-handled by utilizing different commotion filtering strategies. The Microscopic pictures of patient's platelets are given as a contribution to the framework. Figure 1 shows the architecture design.

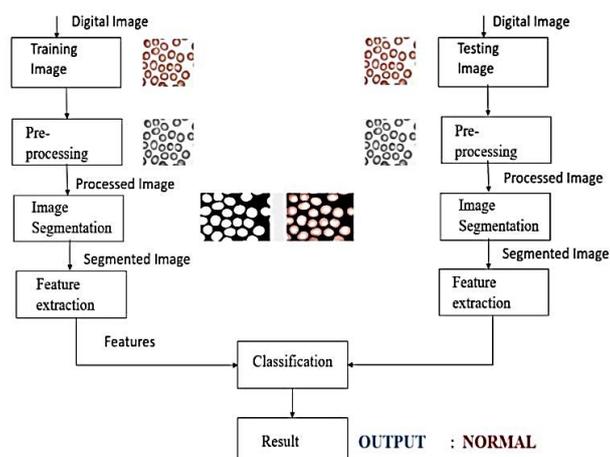


Fig. 1 Architecture Design

Data flow Design

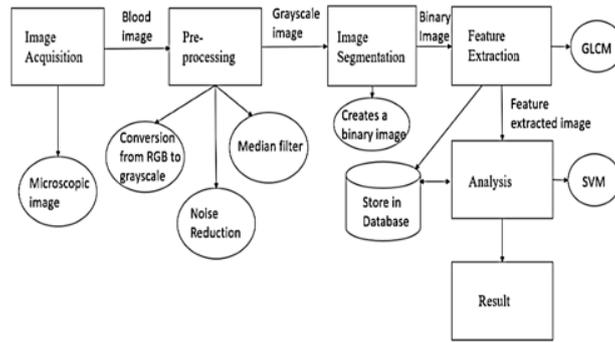


Fig. 2 Data Flow Diagram

Flow of information through a data framework is done using DFD is nothing but a graphical portrayal of information flow. DFD's can be used for representing the information preparing also. Information things on a DFD flow from an exterior information source or an internal information store to an inner information store or an outside information sink, by means of an interior procedure. Details on planning, or requesting of procedures or whether a procedure will work or not , these data are given from DFD.

Sequence Diagram

A sequence diagram in Unified Modeling Language(UML) is a type of collaboration outline that shows how procedures work with each other and in what request. It is a developed message Sequence Chart. A grouping chart appears, as equal vertical lines ("lifelines"), different procedures or items that live at the same time, and, as also horizontal arrows. The messages traded between them, in the request where they happen. Figure 3 shows the arrangement of steps for the proposed sequence diagram.

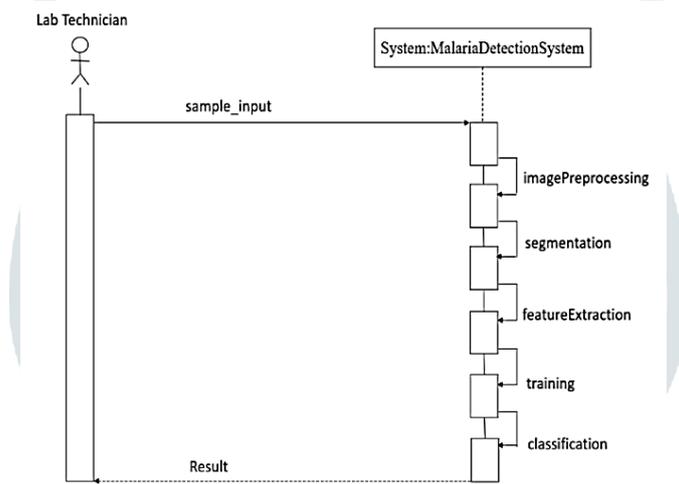


Fig. 3 Sequence Diagram

Decomposition Description

Graphic representation of logic sequences, work or manufacturing process, organization chart, or similar formalized structure is known as a flowchart which is formalized based on requirement. Figure 4 represents the flow chart of the proposed system.

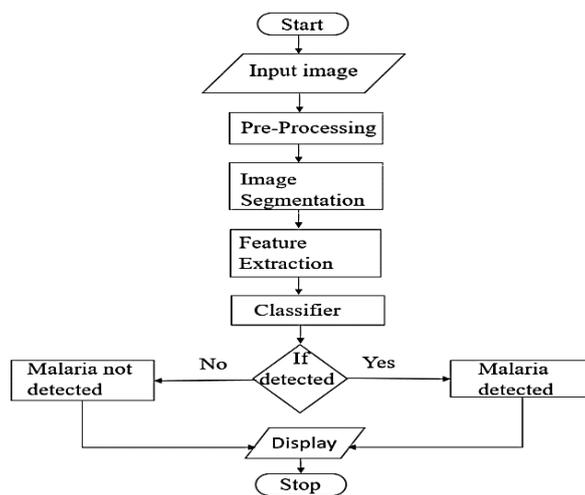


Fig. 4 Flow Chart

Types of testing

The main reason for testing is to show that the developed product meets the expected conditions and requirements under the set environmental conditions.

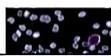
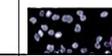
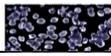
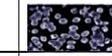
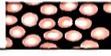
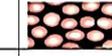
F. Types of testing

1. Unit testing
2. Integration Testing
3. System Testing

Test Cases

For a particular output and ideally appropriate output, number of test cases have been tested and presented in below Table 1.

TABLE I – TEST RESULTS

TC#	Description	Expected Result	Actual Result	Status
TC-1				Pass
TC-2				Pass
TC-3				Pass
TC-4				Pass
TC-5				Fail
TC-6				Fail

EXPERIMENTAL RESULTS

Experimental results are validated using successful detection of the malaria parasites in blood. The progress of the proposed system is monitored using precision and recall graph. The user interface is as shown in the Figure 5 above, which is mainly used for displaying graphical data in one or more window having control, called components thereby helping the user to perform interactive tasks.

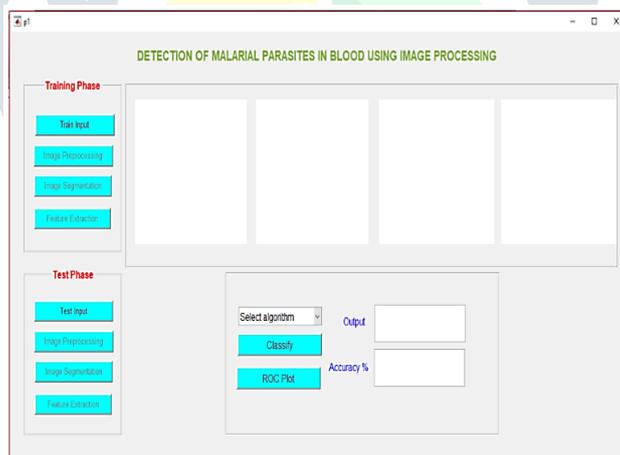


Fig. 5 Snap shot of GUI

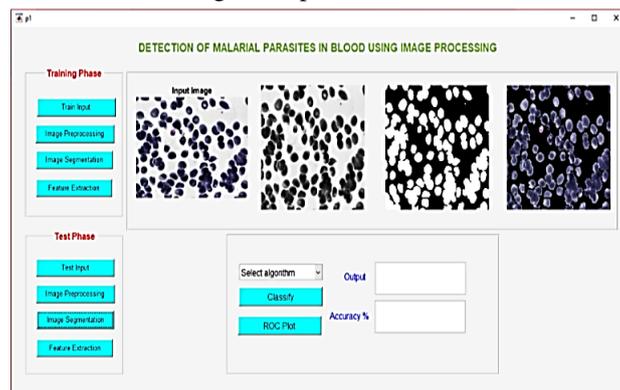


Fig. 6 Preprocessing and segmentation process

The above Figure 6 shows the software setup of training phase of the proposed system, here multiple images which are taken from the database are trained as per the procedures. As can be seen from the figure above, the first image is taken, that is the original image is taken for the input, then the second block is also preprocessed after the noise reduction and then the third block is the image which is obtained after segmentation technique.

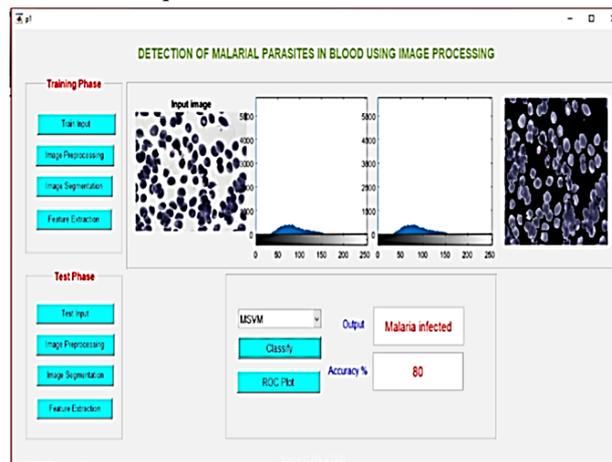


Fig. 7 Result obtained using MSVM algorithm

Here a random input type image is uploaded to the system and even for the single test input the same procedure is repeated. The algorithm is selected and the classifier will segregate the image and find whether the given image is actually infected with malaria or not. Figure 7 shows the actual results obtained using MVSM classifier. Extraction process usually take the features from the image and then calculates the values and compares it with the stored value which is in the database. Thus it is used to predict the presence of malaria or not. The below Figure 8 shows the feature extraction process that is represented in form of histograms.

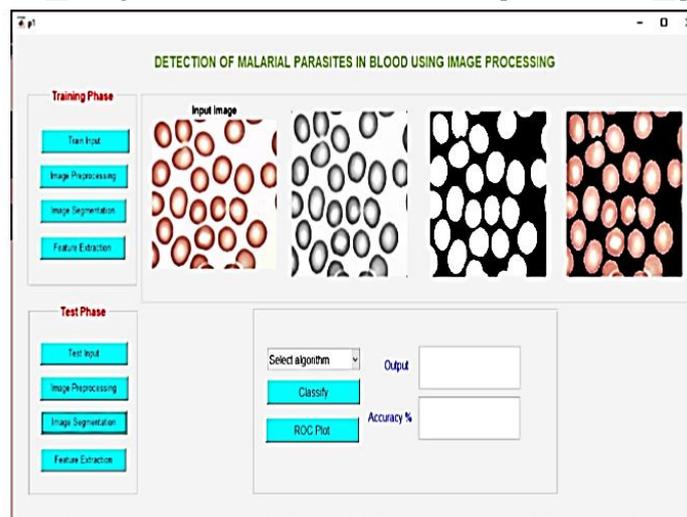


Fig. 8 Snapshot of Test input for an uninfected image

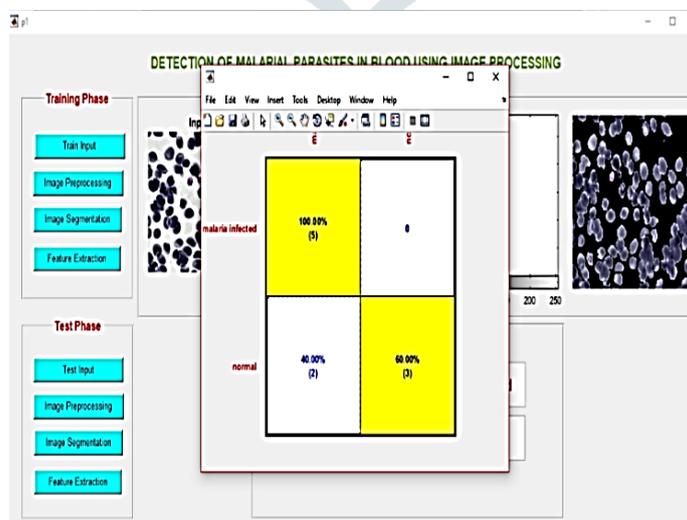


Fig. 9 Confusion matrix for the uninfected image using SVM

Using the method of confusion matrix for the uninfected blood image, which is used to calculate the against the recall and recession. Figure 9 shows the results of the confusion matrix for the uninfected image using SVM. The Figure 10 and Figure 11 shows that once the confusion matrix is done the ROC plot using the 2 classes, then the browsing for input image for testing and pre-processing for the segmentation process in initiated.

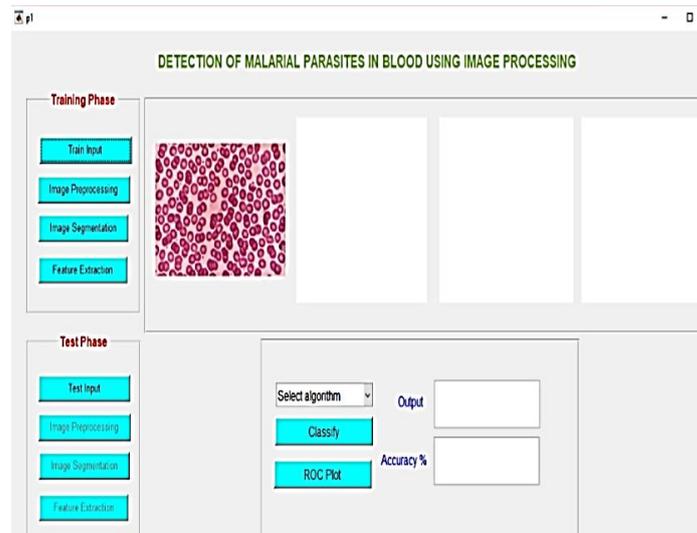


Fig. 10 Browsing the single input image for testing

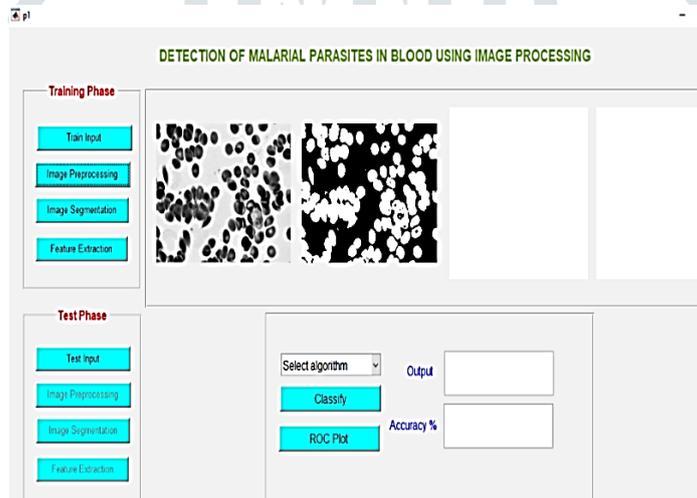


Fig. 11 Pre-processing and the segmentation process

V. CONCLUSION

The proposed work here is to build up an exact, quick and low cost economic device for the finding and location of parasites of intestinal sickness also known as malaria, using thin microscopic images, and utilizing Image Processing. Here, tiny blood Images of ordinary, malaria contaminated are taken, which is precisely pre-processed and, then finally segmented. Main aspects are separated from the segmented image in order to reduce the measure of information to hold on for examination and analysis. From that point SVM and MSVM classifiers are utilized to identify whether the microscopic image of the platelets is intestinal sickness (malaria) tainted or not upon which the final result is concluded.

The future improvements and developments are as mentioned:

1. Validation of info pictures: Ability of the system that it is able to recognize the images that are not blood cells.
2. Ongoing or Real time: It can be utilized in real time by interfacing an outer magnifying instrument legitimately to the PC so that to get a prompt yield without wasting time for advanced images from magnifying instruments.

REFERENCES

- [1] W. World Health Organization, "Who Report 2015," 2015.
- [2] N. R. Shet and N. Sampathila, "An Image Processing Approach for Screening of Malaria," *Canar. Eng. Coll. Mangalore*, pp. 395–399, 2015.
- [3] A. Awchite, K. Pasalkar, M. Killedar, S. Dongre, and S. Godse, "Detection of malaria and anemial parasites in blood using image processing,"
- [4] N. E. Ross, C. J. Pritchard, D. M. Rubin, and A. G. Duse, "Automated image processing method for the diagnosis and classification of malaria on thin blood smears," *Medical and Biological Engineering and Computing*, vol. 44, no. 5, pp. 427–436, 2006.

- [5] S. Savkare, S. Narote, et al., "Automatic detection of malaria parasites for estimating parasitemia," *International Journal of Computer Science and Security (IJCSS)*, vol. 5, no. 3, p. 310, 2011.
- [6] H. A. Nugroho, S. A. Akbar, and E. E. H. Murhandarwati, "Feature extraction and classification for detection malaria parasites in thin blood smear," in *2015 2nd International Conference on Information Technology, Computer, and Electrical Engineering (ICITACEE)*, pp. 197–201, IEEE, 2015.
- [7] D. A. Ghate, C. Jadhav, and N. Rani, "Automatic detection of malaria parasite from blood images," *Int J Comput Sci Appl*, vol. 1, 2012.
- [8] P. Rakshit and K. Bhowmik, "Detection of presence of parasites in human rbc in case of diagnosing malaria using image processing," in *2013 IEEE Second International Conference on Image Information Processing (ICIIP-2013)*, pp. 329–334, IEEE, 2013.
- [9] A. Sanadhya, A. Mathur, M. S. Chouhan, and I. Suwalka, "Detection of malaria parasite using digital image processing,"
- [10] N. A. Khan, H. Pervaz, A. K. Latif, A. Musharraf, et al., "Unsupervised identification of malaria parasites using computer vision," in *2014 11th International Joint Conference on Computer Science and Software Engineering (JCSSE)*, pp. 263–267, IEEE, 2014.
- [11] J. Somasekar, B. E. Reddy, E. K. Reddy, and C.-H. Lai, "An image processing approach for accurate determination of parasitemia in peripheral blood smear images," *International Journal of Computer Applications*, pp. 23–28, 2011.
- [12] S. Suryawanshi and V. Dixit, "Comparative study of malaria parasite detection using Euclidean distance classifier & svm," *International Journal of Advanced Research in Computer Engineering & Technology (IJARCET)*, vol. 2, no. 11, pp. 2994–2997, 2013.

