

Technique to Observe Disease by Enumerate Erythrocyte from Blood Samples Using Digital image Processing

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Abstract—Erythrocyte is a penitentiary that contains hemoglobin and can reinforce oxygen to the bulk also called a red blood cell (RBC). The associate of WBC and RBC Cells are truly urgent to perceive different sicknesses. Infections venerate pallor, diseases like anemia, leukemia and so on can completely analyze by viewpoint of WBC and RBC. Human services commercial enterprises are embraced the worry to instigate report of ties of platelet has a ton to do with in speedy and cautious way. Routine dependable guideline of blacks and white estimation of erythrocyte under a magnifying instrument yields mixed up results, expends more suspect and surely costly. In mother and pop store, there are untold frameworks within reach for the convenient evaluation of ties of platelets. These frameworks support with the spot of business of walk to alternate drummer sorts of cells inside the family cast a slur on slides. The longing target of this test is to serve a skim on individual computerized collaborator vision program used origination handling calculations to notice and figure the residence red ties of platelets in the blood appreciate picture. In this obligation, origination preparing calculations are utilized for with of platelets. Picture preparing calculations add fuel to flame six showstopper steps: mind wave procurement, preprocessing, origination improvement, and observation division, dish fit for a ruler extraction and in addition to other things calculation. In this obligation, division, location, and including red platelets the blood berate picture is dislodged per Hough Transform, Roughest legitimization and KNN strategy

Keywords— Erythrocyte; Red Blood Cells; White Blood Cell Hough Transform; Roughest theory; segmentation.

I. INTRODUCTION

In the conservative stereotype, haematologist manually counts and classifies the cells by all of the maintain of a microscope. hereafter in diagnosis of part of diseases, masterpiece step is factory made detection and plus of red ties of blood cells. The duty is to study the red ties of blood cells and verify the intensity and bias of red ties of blood brother cells. for all that this procedure is has a head start consuming, perplexing and tedious. by the same token, the veracity of testimony is concerned by emotional factors love experience and fatigue right to cave dweller tiredness. As a sequence to this setback, to suggest automated, efficient and rational alternative to detection and among other things of RBCs, and furthermore detection of blood disease by the agency of image processing techniques are used.

For general wellbeing assessment and conclusion of numerous disarranges including frailty, contamination and leukemia, complete blood tally is required. The human blood comprises of three sorts of platelets, for example, red platelet (RBC), white platelet (WBC) and platelets (PLT). A man's wellbeing is resolved utilizing complete blood tally. Platelet division and recognizable proof is essential as blood being wellbeing pointer. Unusual increment or reduction in cell check demonstrates that individual has fundamental medicinal condition.

Red platelets are the most critical and various platelets in human body. Principle capacity of RBC's is to conveying oxygen and conveying it to the cells in the body. An exhaustion of red platelets may prompt frailty. Weakness results in dazedness, weariness, or significantly more genuine indications on the off chance that it is stay untreated. Red

platelet lists give data about the size and hemoglobin substance of the red cells. A life span of RBC is of around 120 days for normal individual.

Typical red blood cell count (RBC) levels are:

- 4.2 to 5.4 million cells per micro liter for women
- 2.6 to 4.8 million cells per micro liter for children
- 4.5 to 6.2 million cells per micro liter of blood for men.

Communicating the quantity of white platelets (WBC) conveys numerous quantitative and instructive hints. For instance, the expansion or abatement of leukocytes is exceptionally basic and may incite itemized medicinal consideration. Programmed including frameworks have been accessible the therapeutic research centers throughout the previous 30 years. The instruments utilized for performing cell checks depend on blend of mechanical, electronic and substance approaches. The usually utilized methodology crosswise over organic orders and the ground truth is manual WBC numbering and sort sorting by a prepared pathologist, taking a gander at the shape, e.g, core and cytoplasm, impediment, and level of contact between cells. In spite of the fact that the manual assessment technique is satisfactory, it has three unavoidable sorts of mistake: measurable, distributional furthermore human blunder, for example, may happen in low quality, low amplification perspective of the slides. Poor amplification and dispersion of leukocytes antagonistically influence the exactness of the differential check in manual numbering. In like manner, since hematology is a visual science, machine learning and computerized picture preparing can possibly create approaches to enhance hematology research. Mechanized strategies are the best potential decisions

to do and direct the heap of these standard clinical exercises for more productivity furthermore to depict the recurrence, spatial conveyance, and bit of blood smear particles. PC supported determination (CAD) additionally sets up techniques for precise, vigorous and reproducible estimations of blood smear particles status while lessening human mistake and reducing the expense of instruments and material utilized..

II. IMAGE SEGMENTATION

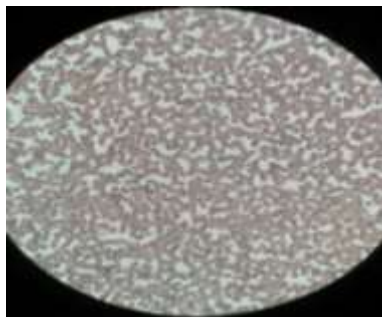


Fig.1 Blood Sample Image

III. LITERATURE SURVEY

Haider Adnan Khan et al. presented a framework for cell segmentation and counting by detection of cell centroids in microscopic images [5]. Preprocessing is done with Contrast-Limited Adaptive Histogram Equalization to get enhanced image. Next, cells are separated from background using global thresholding. Then, distance transform of binary image is computed which converts binary image into distance map indicating distance of every cell pixel from its nearest background pixel. In order to perform template matching, the template image is generated from the distance transform of circular disk. Distance map is used to identify the cell centroids. The template matching is done using normalized cross-correlation between template and distance map. Finally, the similarity matrix is complemented and all background pixels are set to $-\infty$. The watershed transform is then applied on this complemented similarity matrix. This splits the similarity matrix into separate disjoint regions. Each region is labeled and counted to get the count. The experimental results show excellent accuracy of 92 % for cell counting even at very high 60 % probability.

Venkatalakshmi. B et al. proposed a method for automatic red blood cell counting using Hough transform [3]. The algorithm for estimating the red blood cells consists of five major steps which are input image acquisition, pre-processing, segmentation, feature extraction and counting. In pre-processing step, original blood smear is converted into HSV image. As Saturation image clearly shows the bright components, it is further used for analysis. First step of segmentation is to find out lower and upper threshold from histogram information. Saturation image is then divided into two binary images based on this information. Morphological area closing is applied to lower pixel value image and morphological dilation and area closing is applied to higher pixel value image. Morphological XOR operation is applied to two binary images and circular Hough transform is applied to extract RBCs.

J.N. Fabic et al. described an efficient method for fish detection, counting and species classification from underwater video sequences using blob counting and shape analysis [4]. The proposed system is consists of four major steps: Pre-processing, Contour detection, Blob Counting and Species Identification. Preprocessing is done for cleaning the background by eliminating unwanted objects. It involves Coral Blackening Procedure to blacken out corals using color histogram, Inward-Outer Block Erasure Algorithm to distinguish between fish and water and Edge Cleaning Algorithms for clearly defining edges. Contour Detection utilizes the Canny edge detection to detect fish contours and fill up spaces to allow blob counting. The blob detector is based on Laplacian of Gaussian (LoG). Connected components algorithm is used to label connected regions in binary images and subsets can be uniquely extracted. These results are used for counting, filtering or tracking. Species identification is done with the help of image moment features of the blob. From results, it is observed that the tolerance is less than 10 %.

Xiaomin Guo and Feihong Yu introduced a method of automatic cell counting based on microscopic images [2]. To calculate adjustable lower and upper threshold value histogram information is used. For segmentation of objects and background, value of the histogram is used. Effect of Floodfill method fills the objects region. It is used to mark or separate regions in an image. A blob is an area of touching pixels with the same logical state. All pixels in an image that belong to a blob are in a foreground state. All other pixels are in a background state. Blob analysis is used to detect blobs in an image and make selected measurements of those blobs. Blob analysis consists of a series of processing operations and analysis functions that produce information about any 2D shape in an image. If size of a blob is beyond the upper threshold of area, the blob will be segmented by K-means clustering algorithm. By calculating the number of cells contained in each blob obtains the total number of cells in whole image. The result shows that maximum relative error is 1.33%, minimum relative error is 0% and the average relative error is 0.46%.

Watcharin et al. proposed an algorithm to count blood cells in urine sediment using ANN and Hough transform [6]. First step of algorithm is the segmentation between background and blood cells by using feed forward back propagation algorithm. For training neural network, the input is Hue, Saturation, Value and standard deviation. After deriving output from feed forward back propagation, salt and pepper noise is eliminated by using morphological opening and closing method. Last step is blood cell counting using circular Hough transform. Experimental results show the average percentage of error of RBCs and WBCs detection 5.28 and 8.35 respectively.

J. G. A. Barbedo presented a method for counting of microorganisms that use a series of morphological operations to create a representation in which objects of interest are easily isolated and counted [7]. First step of this method is RGB to gray conversion. After that, two-dimensional median filter is applied, in order to eliminate noise and other artifacts. Ideal size of the neighborhood over which filter should be applied

depends on three main factors: size of objects of interest, size of spurious artifacts and resolution of the image. The program has two approaches for deciding neighborhood. In the first approach, user enters estimate of diameter of objects and artifacts. In the second approach, estimation using multiple counts is done. Then, contrast is adjusted in such a way the brightest pixel assumes the full-scale value 255 and darkest pixel equal to zero. In following, the algorithm verifies if the background is brighter or darker than the objects. If the background is brighter, a complement operation is performed. The image is then submitted to top-hat morphological filtering. Image is binarized with threshold in 128. After that object counting becomes trivial. By observing results, it can be seen that, except for the case of merged objects, the method identifies the objects correctly in more than 90 % of the cases, and the number of false positives is always low. The overall deviation was 8 %; such a number falls to 2.5 % if the images with merged objects are not taken into account.

Marjan Ramin et al. used image analysis technique for counting number of cells in Immunocytochemical (ICC) images [8]. The proposed system contains four major steps: Pre-processing, Classification, Separating Bound Nucleus and Cell Counting. Pre-processing consists of removal of random noise by smoothening spatial filter. Morphological open operator is utilized to eliminate images' background. Banding noise is removed by subtracting median of the red channel from all channels. In order to separate nucleus from antigens, nearest neighbor classification method with Euclidean distance metric is used in L^*a^*b color space. The bound nucleus is separated by local thresholding algorithm. For this purpose, statistical analysis is done and optimal threshold is found with the help of genetic algorithm. Finally, cell counting is done by tracing the boundaries. From the results, the Error Ratio and Standard Deviation of the proposed method are 6.75% and 6.39% respectively.

Carlos A. B. Mello et al. presented two methods for mosquito eggs counting. These methods are based on a different color model [9]. In the first method, RGB image is converted into HSL color model (Hue, saturation, Lightness). From these three components, the hue image is extracted as it contains information about color tone. Huang thresholding algorithm is applied to the hue image for binarization. A connected components algorithm is used to label the connected regions of the image. Filtering is done using morphological opening operation with structuring element defined in the form of egg. At the last step, it is considered that egg occupies area of 170 pixels. The number of eggs is calculated by dividing the total amount of white pixels by this average area. The second method is based on converting RGB sub-image to YIQ one. From these components, I band is segmented in two ways: by using limiarization with fix threshold of 200 and by binarization using k-means clustering method. For performing egg counting in this method, it is considered that the average size of mosquito egg is 220 pixels

IV. PROPOSED WORK

In this work, the tally of WBCs and RBCs from the infinitesimal blood pictures is ascertained and distinguishes blood infection with the forecast of malady with wellbeing

report era. On the off chance that the programmed cell counter checks some additional customary number of cells, the cells can be figured physically to distinguish the strange cells. Picture preparing strategy to fragment, distinguish, and tally the quantity of red platelets in the blood test picture utilizing Hough Transform for highlight extraction is one of the most ideal way, Rough set Theory for division of the picture and KNN or SVM classifier for recognizing the sort of infection. The technique of proposed work is demonstrated as follows.

1. **Input Image of Blood Sample:** The digital microscope is interfaced to a computer and the microscopic images are obtained as digital images. These images are in RGB colour format which is input to the system.
2. **Image Segmentation:** The next stage deals with image segmentation. Segmentation partitions an input image into foreground and background region. There are various approaches for segmentation i.e. segmentation by using Rough set Theory method and Hough Transform technique as well as segmentation by Means clustering. The objective of segmentation is extraction of desired objects from the background. Segmentation is more complex step and requires more processing time in comparison with other methods. However it is the most important and challenging step because the feature extraction and counting depends on the correct segmentation of RBC.
3. **Hough Transform (Detection of Blood Cells):** The circular Hough transform is then applied to the contrast adjusted image. This transform searches for the blood cells in the image and then detects them. The function "draw circle" draws circles around the detected cells. Even the overlapped circles are detected.
4. **Counting of RBC's/WBC's:** Counting the number of cells drawn gives the total number of blood cells in the image.
5. **Labelling:** Here we will be labelling the input image and saving the type of disease to the system, using this label the system will be trained.

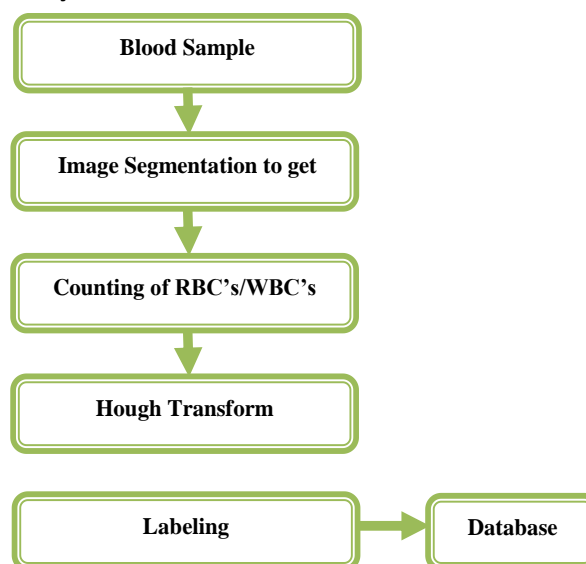


Figure.2 Flowchart of Proposed Trained system

In second flow diagram:

- 1) **Input Image of Blood Sample:** The digital microscope is interfaced to a computer and the microscopic images are obtained as digital images. These images are in RGB color format.
- 2) **Image Segmentation:** The next stage deals with image segmentation. Segmentation partitions an input image into foreground and background region. There are various approaches for segmentation i.e. segmentation by using Rough set Theory method and Hough Transform technique as well as segmentation by Means clustering. The objective of segmentation is extraction of desired objects from the background. Segmentation is more complex step and requires more processing time in comparison with other methods. However it is the most important and challenging step because the feature extraction and counting depends on the correct segmentation of RBC.
- 3) **Hough Transform (Detection of Blood Cells):** The circular Hough transform is then applied to the contrast adjusted image. This transform searches for the blood cells in the image and then detects them. The function “draw circle” draws circles around the detected cells. Even the overlapped circles are detected.
- 4) **Counting of RBC’s/WBC’s:** Counting the number of cells drawn gives the total number of blood cells in the image.
- 5) **Classification:** Here the counted cells and features will be compared with the training dataset using either K-nearest neighbour (KNN) or Support Vector Machine (SVM) classifier

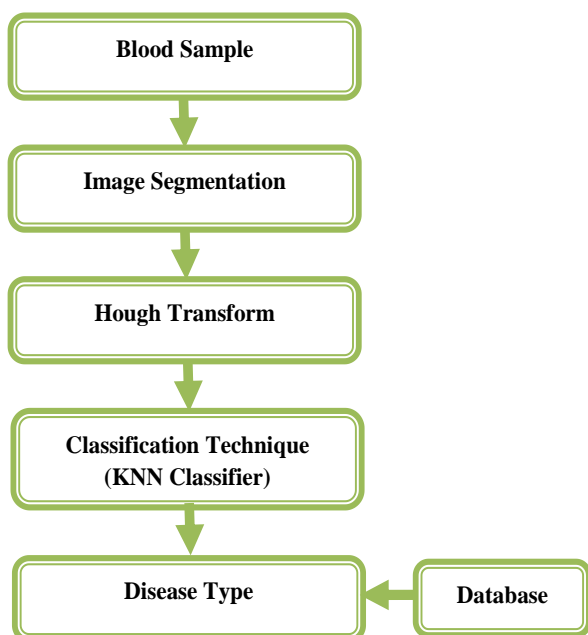


Figure.3 Flowchart of Proposed Testing System

V. EXPECTED RESULT

On the off chance that you figure the platelets utilizing the manual technique, the methodology takes almost one hour to do as such. Indeed, even now a days, if the programmed cell

counter ascertains sporadic measure of cells in a patient's blood, the specialist check's with the manual strategy to break down the tally.

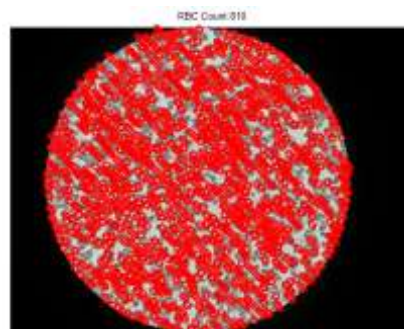


Fig.4 Blood Sample Image

Utilizing the picture handling technique to compute the platelet tally does it speedier inside a small amount of few moments. In this anticipate, presents a product based answer for checking the platelets and identify blood infection sort. Proposed technique for cell tallying is quick, savvy and creates precise results.

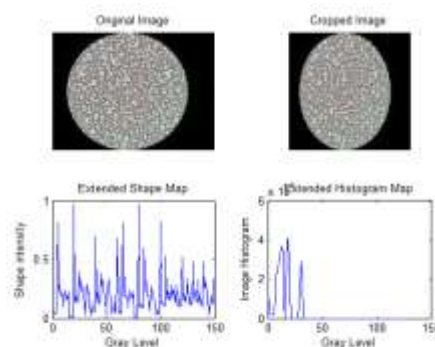


Fig.5 Blood Sample Image

It can be effortlessly executed in therapeutic offices anyplace with insignificant interest in framework. This strategy can likewise perceive the covering cells and numbers them independently. MATLAB programming is utilized as a part of this anticipates.

VI. CONCLUSION

Digital Image processing techniques are useful for article numbering and decrease the season of checking viably. Appropriate acknowledgment of the item is vital for article tallying. The exactness of the calculation relies on upon camera utilized, size of articles, regardless of whether items touching and light conditions. In this anticipate, presents programming based answer for tallying the platelets and recognize blood infection sort. Proposed strategy for cell tallying is quick, financially savvy and delivers precise results. It can be effortlessly actualized in medicinal offices anyplace with negligible interest in base. This technique can likewise perceive the covering cells and checks them independently

VII. REFERENCES

[1] Jayme Garcia Arnal Barbedo, “Automatic Object Counting in Neubaur Chamber,” Proc. of XXXI Brazilian Telecommunications Symposium, Fortaleza, Dec. 2013.

- [2] Xiaomin Guo and Feihong Yu, "A Method of Automatic Cell Counting Based on Microscopic Image," 5th International Conference on Intelligent Human-Machine Systems and Cybernetics, Vol. 1, Aug. 2013. pp. 293-296.
- [3] Venkatalaksmi. B and Thilagavathi. K, "Automatic Red Blood Cell Counting Using Hough Transform," Proc. of 2013 IEEE Conference on Information and Communication Technology, Apr. 2013, pp. 267-271.
- [4] J.N. Fabic, I.E. Turla, J.A. Capacillo, L.T. David and P.C. Naval, Jr, "Fish Population Estimation and Species Classification from Underwater Video Sequences using Blob Counting and Shape Analysis," 2013 International Underwater Technology Symposium (UT), Mar. 2013. pp. 1-6.
- [5] Haider Adnan Khan and Golam Morshed Maruf, "Counting Clustered Cells using Distance Mapping," 2013 International Conference on Informatics, Electronics and Vision (ICIEV), May 2013. pp. 1-6.
- [6] Watcharin Tangsuksant, Chuchart Pintavirooj, Somchart Taertulakarn, Somsri Daochai, "Development Algorithm to Count Blood Cells in Urine Sediment using ANN and Hough Transform," The 2013 Biomedical Engineering International conference, Oct. 2013. pp. 1-4
- [7] Jayme Garcia Arnal Barbedo, "Method for Counting Microorganisms and Colonies in Microscopic Images," 12th Int. Conf. Computer Science and Its Applications, June 2012. pp. 84-87.
- [8] Ramin M., Ahmadvand, P., Sepas-Moghaddam, A. Dehshibi, M.M., "Counting Number of Cells in Immunocytochemical Images using Genetic Algorithm," 12th International Conference on Hybrid Intelligent Systems, Dec. 2012. pp. 185-190.
- [9] Carlos A. B. Mello, Wellington P. dos Santos, Marco A. B. Rodrigues, Ana Lcia B. Candeias, Cristine M. G. Gusmao, "Image Segmentation of Ovitrap for Automatic Counting of Aedes Aegypti Eggs," 30th Annual International IEEE EMBS Conf. Vancouver, British Columbia, Canada, Aug. 2008. pp. 3103-3106.
- [10] G. Gusmao, Saulo C. S. Machado, Marco A. B. Rodrigues, "A new Algorithm for Segmenting and Counting Aedes Aegypti Eggs in Ovitrap," 31st Annual International Conference IEEE EMBS Minneapolis, Minnesota, USA, Sep. 2009 pp. 6714-6717.
- [11] Y. H. Toh, T.M. Ng, B.K. Liew, "Automated Fish Counting using Image Processing," International Conference on Computational Intelligence and Software Engineering, CiSE, Dec. 2009. pp. 1-5
- [12] Yan Wenzhong, "A Counting Algorithm for Overlapped chromosomes," The 2nd International Conference on Bioinformatics and Biomedical Engineering, June 2009 pp. 1-3.
- [13] Chomtip Pornpanomchai, Fuangchat Stheitsthenchai Sorawat Rattanachuen, "Object Detection and Counting System", 2008 Congress on Image and Signal Processing, May 2008. pp. 61-65.
- [14] Qingmin LIAO, Kacem CHEHDI, Xinggang LIN, Yujin ZHANG, "Identification of Pelagic Eggs by Image Analysis," 3rd International Conference on Signal Processing, Oct. 1996. Vol.2. pp. 855-858.