

Automatic Red Blood Cell Counting using Watershed Segmentation

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Abstract— The major issue in clinical laboratory is to produce a precise result for every test especially in the area of Red Blood Cell (RBC) count. The number of red blood cell is very important to detect as well as to follow the treatment of many diseases like anaemia, leukaemia etc. Red blood cell count gives the vital information that help diagnosis many of the patient's sickness. The old conventional method of RBC counting under microscope gives an unreliable and inaccurate result depends on clinical laboratory technician skill. This method puts a lot of strain on the technician. Another method for RBC counting uses the automatic hematology analyzer, this machine is very costlier. So it is not possible all the hospital's clinical laboratory implement such an expensive machine to count the blood cell in their laboratory. This paper introduces an efficient and cost effective computer vision system for automatic red blood cell counting using image based analysis.

Keywords— Health care applications, Red Blood Cell count, MATLAB, Morphological operations, Hough Transform.

I. INTRODUCTION

Content-based image indexing and retrieval has been an important research area in computer science for the last few decades. Many digital images are being captured and stored such as medical images, architectural, advertising, design and fashion images, etc. As a result large image databases are being created and being used in many applications. In this work, the focus of our study is on medical images. A large number of medical images in digital format are generated by hospitals and medical institutions every day. Consequently, how to make use of this huge amount of images effectively becomes a challenging problem [1]. In the field of biomedicine, because of cell's complex nature, it still remains a challenging task to segment cells from its background and count them automatically [2-5]. Among all of the body's tissues, blood is unique due to its existence as the only fluid tissue. A blood cell can be any type of cell normally found in blood which falls into four categories which are red blood cell (RBC), white blood cell (WBC), platelet and plasma [6]. The differences between these groups lie on the texture, color, size and morphology of nucleus and cytoplasm. In blood smear, number of red cells is many more than white blood cells. For example an image may contain up to 100 red cells and only 1 to 3 white cells. Platelets are small particles and are not clinically important [7]. Blood cells form in the bone marrow, the soft material

in the center of most bones. Leukocytes or WBC are cells involved in defending the body against infective organisms and foreign substances. Leukocytes cells containing granules are called granulocytes (composed by neutrophil, basophil, eosinophil). Cells without granules are called agranulocytes (lymphocyte and monocyte) [6]. These cells provide major defense against infections in organisms and their specific concentrations can help specialists to discriminate the presence or the absence of very important families of pathologies [8]. When infection occurs, the production of WBCs increases [6]. Abnormal high or low counts may indicate the presence of many forms of disease, since blood counts are amongst the most commonly performed blood tests in medicine.

Current research is doing on blood counting application in the image segmentation. It is an implementation of automated counting for blood cell which manually done by hemacytometer by using counting chamber. Blood counting is synonym with the complete blood count or CBC which refers to compilation test of red blood cell (RBC), white blood cell (WBC), platelet, hemoglobin and hematocrit. Each of them has their role in the body system and the counting result is important to determine the capability or deficiency of the body system. In short, any abnormal reading of CBC can give a sign of infection or disease. For example, the presence of bacterial infection is diagnosed from increasing WBC count. Plus, specific low vitamin may come from a decreased RBC and thrombocytopenia is referring to low platelet count. The result can influence physician to make the best response and monitor the drug effectiveness from the blood count [14].

CBC consists of several counting of the main component in the blood cell. Each of them has a standard quantity range as a reference for a healthy woman and man. Any counting value out of the range is considered abnormal and physician will interpret the result for further action. In addition, differential count also includes in the measurement of CBC as a division of WBC count for five different types of WBC. They are neutrophils, lymphocytes, monocytes, eosinophils and basophils. The standard count for them is 60%, 30%, 5%, 4% and below 1% respectively from the total WBC counts. Table 1 shows the standard CBC for the healthy person divided by gender.

Table 1: Normal Blood Count Differentiate by Gender

| Blood Cell Types | Gender | |
|------------------|---------------------------------|---------------------------------|
| | Men | Women |
| RBX | 5.4-6.0 Million/micro liter | 4.0-5.0 Million/micro liter |
| WBC | 4.5-11 Thousand/micro liter | 4.5-11 Thousand/micro liter |
| Platelet | 150-450 Thousand/micro liter | 15.0-45 Thousand/micro liter |
| Hematocrit | 42%-50% | 36%-45% |
| Hemoglobin | 14-17 Grams/100 milli liters | 12-15 Grams/100 milli liters |

II. IMAGE SEGMENTATION

In the segmentation process, morphological technique is major used because the mathematical morphology offers a powerful tool for segmenting images and useful to describe the region shape, such as boundaries, skeletons and texture. The first method in this process divides saturation, S image into two images output by applying the thresholding process. Thresholding is one of the methods to extract and segment the object from the background by selecting any point, T [15].

Any point for which is called an object point, otherwise the point is called background point. Thresholding normally results in binary image and the mathematically; the operation can be expressed as;

$$g(x, y) = \begin{cases} 1 & \text{if } f(x, y) \geq T \\ 0 & \text{if } f(x, y) < T \end{cases}$$

where the pixels labeled 1 is corresponded to object whereas the pixels labeled 0 are corresponding to the background.

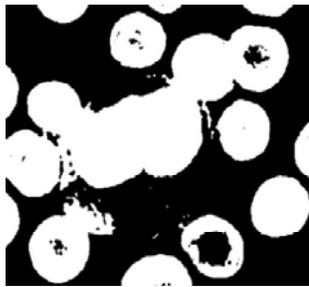


Figure 2(a)



Figure 2(b)

- (a) Morphological area closing on lower pixel value image.
- (b) Morphological dilation and area closing on higher pixel value image.

Jianhua et al. [21] stated that in the case of cell segmentation for blood, edge detection performs poorly on cell images because not all boundaries are sharp and it is difficult to get all edge information and locate cells accurately. They developed an iterative Otsu's approach based on circular histogram for the leukocyte segmentation. Otsu's approach is generalized on the base of least square method. R. Suresh Kumar et al. [22] discussed about an approach for color image segmentation using higher order entropy as a textural feature for determination of thresholds over a two dimensional image histogram. Two basic models for color images are the RGB (Red, Green, Blue) color model and the HIS (Hue, intensity, saturation) color model. Two methods of color image segmentation used RGB space as the standard processing space. These techniques might be used in blood cell image segmentation. Color images are very rich source of information, because they provide a better description of a scene as compared to grayscale images. Hence, color segmentation becomes a very important issue [22]. Khoo Boon et al. [23] performed comparisons between nine image segmentation methods which are gray level thresholding, pattern matching, morphological operators, filtering operators, gradient-in method, edge detection operators, RGB color thresholding, color matching and HSL (hue, saturation, lightness) and color thresholding techniques on RBC and concluded that there is no single method can be considered good for RBC segmentation [23].

In image enhancement process, there are two common image processing techniques used in order to reduce the noise and at the same time to enhance the image. Figure 3 shows the flow process in enhancement processing which are analysed in hue-saturation value color space (HSV) and the green component image. For HSV, we proceed with analyses in saturation component, S, because this S image shows clearly the bright objects such as white blood cell and parasites, therefore, it's easy to distinguish the red blood cells with another cell.

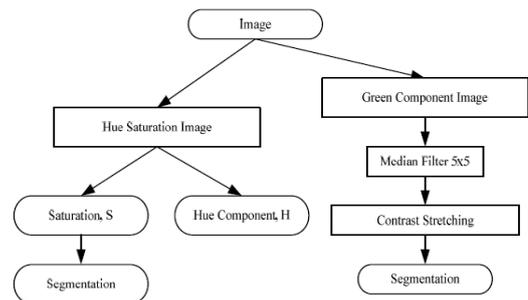


Figure 3: Image Enhancement

III. COUNTING METHODS

Roy A. Dimayuga et al [13] used the histogram thresholding to distinguish the nucleus of the leukocyte or white blood cells from the rest of the cells in the image. Ramin Soltanzadeh [9] has proposed feature extraction technique based on morphology in his three blood cell's experiments. Based on morphology of the cells, the mass center of each cell in the images and then find the distance

of each pixel on an edge from the center. Heidi Berge [10] has purposed thesegmentation red blood cells in a thin blood smear image which is based on the Zack's Method[11]. This method is one of the approached to determine the red blood cells tresholding where aline is drawn between the two peaks and between these two peaks, they used the point which isfurthest from the drawn line as a threshold for red blood cell. In the conclusion for this technique,the segmentation result is better to the blood smear which in case red blood cells is sparse and inthe image. However, in images with high Red blood cell concentrations, large clumps may resultand this method is less accurate. Guitao et. al [12] purposed the Hough transform in detecting andextracting the red blood cells in the urine micrograph. Based on Hough transform, Guitao hasused the geometrical feature to detect the circle center in the image.

Haider Adnan Khan et al. presented a framework for cell segmentation and counting by detection of cell centroidsin microscopic images [16]. Preprocessing is done with Contrast-Limited Adaptive Histogram Equalization to getenhanced image. Next, cells are separated from background using global thresholding. Then, distance transform ofbinary image is computed which converts binary image into distance map indicating distance of every cell pixelfrom its nearest background pixel. In order to perform template matching, the template image is generated from thedistance transform of circular disk. Distance map is used to identify the cell centroids. The template matching isdone using normalized cross-correlation between template and distance map. Finally, the similarity matrix iscomplemented and all background pixels are set to $-\infty$. The watershed transform is then applied on thiscomplemented similarity matrix. This splits the similarity matrix into separate disjoint regions. Each region islabeled and counted to get the count. The experimental results show excellent accuracy of 92 % for cell countingeven at very high 60 % probability.

Watcharin et al. proposed an algorithm to count blood cells in urine sediment using ANN and hough transform [17].First step of algorithm is the segmentation between background and blood cells by using feedforwardbackpropagation algorithm. For training neural network, the input is Hue, Saturation, Value and standarddeviation. After deriving output from feedforward backpropagation, salt and pepper noise is eliminated by usingmorphological 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 8.35respectively.

6. J. G. A. Barbedo presented a method for counting of microorganisms that use a series of morphological operationsto create a representation in which objects of interest are easily isolated and counted [18]. First step of this method isRGB to gray conversion. After that, two-dimensional median filter is applied, in order to eliminate noise and otherartifacts. Ideal size of the neighborhood over which filter should be applied depends on three main

factors: size ofobjects of interest, size of spurious artifacts and resolution of the image. The program has two approaches fordeciding neighborhood. In the first approach, user enters estimate of diameter of objects and artifacts. In the secondapproach, estimation using multiple counts is done. Then, contrast is adjusted in such a way the brightest pixelassumes the full-scale value 255 and darkest pixel equal to zero. In following, the algorithm verifies if thebackground is brighter or darker than the objects. If the background is brighter, a complement operation isperformed. The image is then submitted to top-hat morphological filtering. Image is binarized with threshold in128. After that object counting becomes trivial. By observing results, it can be seen that, except for the case ofmerged objects, the method identifies the objects correctly in more than 90 % of the cases, and the number of falsepositives is always low. The overall deviation was 8 %; such a number falls to 2.5 % if the images with mergedobjects are not taken into account.

IV. RESULTLS AND DISCUSSION

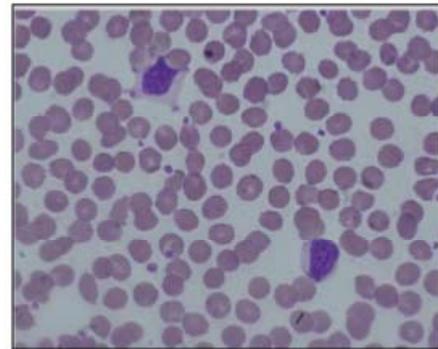


Figure 4: Original Image of Blood for 40X object

When the operation of masking is applied, the maskedimage has diminished the WBC nucleus morphological view.After morphological operation involving binary erosion andfilling holes, the RBC can be viewed accordingly. In thisstudy, masking has been used to remove WBC and platelet issubtracted by morphological operators. The left one will beRBC which represent the RBC segmentation. Figure 5 showsthe result of the RBC segmentation from the elimination ofWBC nucleus and small particles including platelets.

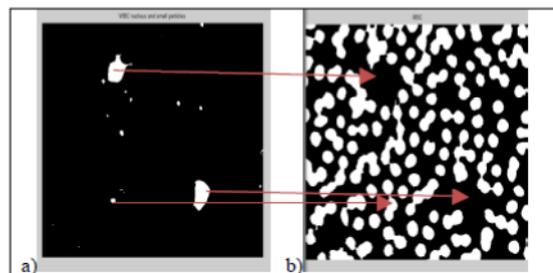


Figure 5: a) Segmented WBC nucleus b) RBC segmentation from theelimination of WBC nucleus and small particles.

To handle overlapping RBC, it involves Laplace ofGaussian (LoG) edge detection, morphological

operation, gradient magnitude and marker controlled watershed algorithm. The result from erosion on RBC segmentation result and gradient magnitude has being used together as mask with watershed algorithm to form marker controlled watershed algorithm. This could avoid oversegmentation which often occurs for watershed algorithm. After dilation, it being used together with LoG edge detection on the Ycbr second component of the image as mask and segmented RBC as marker. Lastly, it being superimpose to the original image. The result of [19] is given in figure 6.

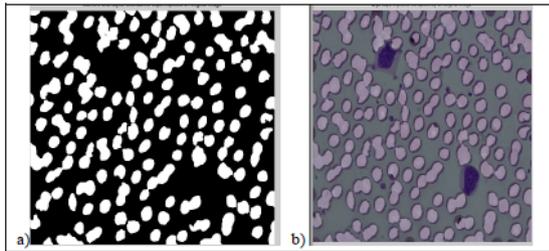


Figure 6: a) Separation of overlapping cell b) Superimpose on original image.

The circular hough transform is applied to the contrast adjusted image by some of researchers.

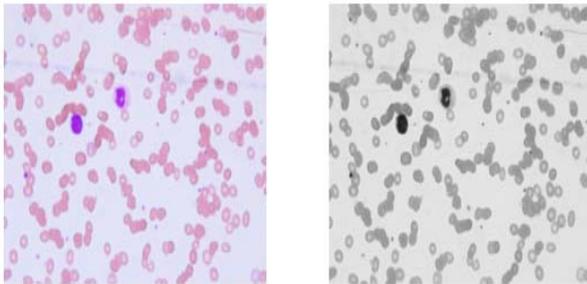


Figure 7(a) Microscopic Image and (b) Green Plane Extraction – I

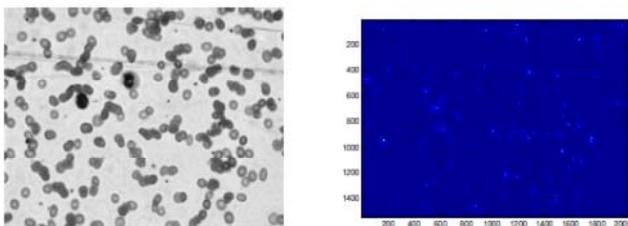


Figure 7(c) Contrast adjustment-1 and (d) Accumulation array-1

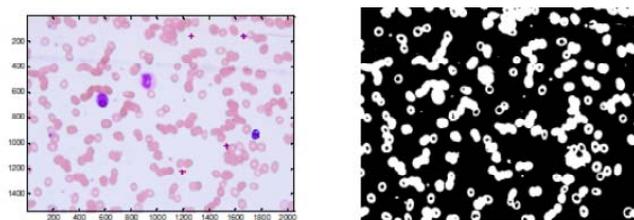


Figure 7(e) Detected blood cells-1 and (f) Binary image-1

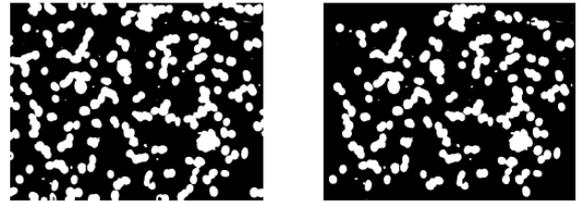


Figure (g) Holes filled-1 and (h) Borders cleared-1

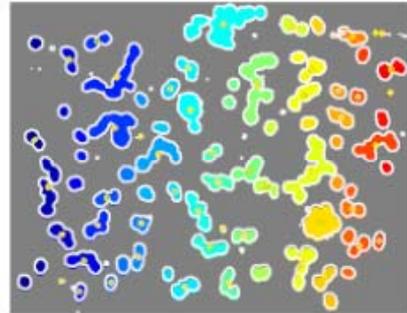


Figure 7(i) Labelled Image

The results of the image can be used as good input in determining the number of red blood cells by using Hough transform technique. By using the MATLAB, all the importance's aspects like correct algorithm and system has been successfully produced. With correct algorithm, the red blood cells can be detected and segmented as well as estimated the number of the red blood cells. Through system created using MATLAB, it also enable the study of the morphological features of the red blood cells image, thus, can determine whether the person is normal or otherwise by referring amount of red blood cells in human blood [15].

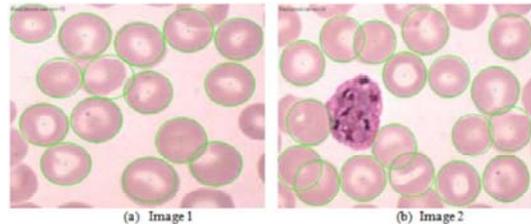


Figure 8 (a) and (b)

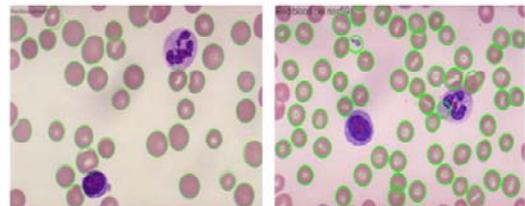


Figure 8(c) and (d)

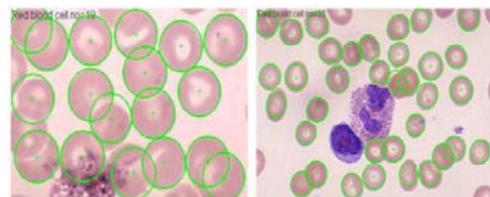


Figure 8(e) and (f)

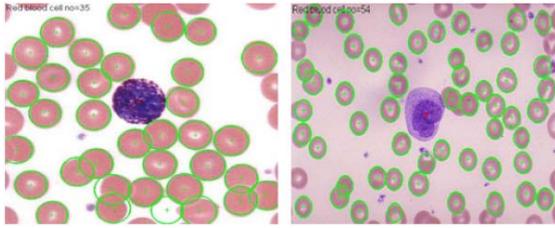


Figure 8 (g) and (h)

Figure 8: results of 8 samples of red blood cells after performing Hough transform technique.

Object counting using image processing has huge applications where automation is to be introduced and time of counting is to be reduced. Some of the main applications of object counting in industrial systems are packaging, quality control, and so on. It is helpful in the research areas where objects are of very small size. Object counting algorithm can be also used to track and identify objects. The present methods can be extended to have counting system based on user selected attributes.

V. CONCLUSION

Image processing techniques are helpful for object counting and reduce the time of counting effectively. Proper recognition of the object is important for object counting. The accuracy of the algorithm depends on camera used, size of objects, whether or not objects touching and illumination conditions.

REFERENCES

- [1] Lehmann T.M., Wein B., Dahmen J., Bredno J., Vogelsang F. & Kohnen M. : Content based image retrieval in medical applications : a novel multi step approach. International Society for Optical Engineering (SPIE), 3972, pp.312-320.(2000)
- [2] Dwi Anoragaingrum : Cell segmentation with median filter and mathematical morphology operation, proceeding of the IEEE 10th International Conference on Image Analysis and Processing (ICIAP), pp. 1043-1046 (1999).
- [3] Keng Wu et al.: Live cell image segmentation, IEEE Trans on Biomedical Engineering, 42(1), pp.1-12.(1995).
- [4] Mark B. Jeacocke, Brian C. Lovell : A Multi-resolution algorithm for Cytological image segmentation, The second Australian and New Zealand conference on intelligent information systems, 322-326 (1994).
- [5] Choi H, Baraniuk R., Multiscale : Image segmentation using wavelet-domain hidden Markov models, IEEE Transaction on image processing, 10(9), pp.1309-1321 (2001).
- [6] H Elaine N. Marieb (2006): Essentials of Human Anatomy & Physiology, 8th ed. Pearson Benjamin Cummings
- [7] Fatemeh Zamani, Reza Safabakhsh: "An Unsupervised GVF Snake Approach for White Blood Cell Segmentation Based on Nucleus", Signal Processing, The 8th International Conference on Volume 2, 2006.
- [8] Vincenzo Piuri, Fabio Scotti: "Morphology Classification of Blood Leucocytes by Microscope Images", IEEE International Conference on Computational Intelligence for Measurement Systems and Applications, Boston, MA, USA, 14-16 July 2004.
- [9] Ramin Soltanzadeh. "Classification of Three Types of Red Blood Cells in Peripheral Blood Smear Based on Morphology. Proceedings of ICSP, 2010.
- [10] Heidi Berge, Dale Taylor, Sriram Krishnan, and Tania S. Douglas. Improved Red Blood Cell Counting in thin Blood Smears. Proceedings of ISBI, 2011. pp.204-207.
- [11] Zack G.W., Rogers W.E. and Latt S.A. "Automatic-measurement of sister chromatid exchange frequency." Journal of Histochemistry & Cytochemistry 25, 1977, 741-753.
- [12] Guitao Cao, Cai Zhong, Ling Li and Jun Dong. "Detection of Red Blood Cell in Urine Micrograph". The 3rd International Conference on Bioinformatics and Biomedical Engineering (ICBBE). 2009.
- [13] Roy A. Dimayuga, Gerwin T. Ong, Rainier Carlo S. Perez, Gefferson O. Siy, Saman C. Sohrabi Langroudi and Miguel O. Gutierrez. "Leukemia Detection Using Digital Image Processing in Matlab". ECE Student Forum, De La Salle University, Manila. March 26, 2010.
- [14] Miswan, M. F., et al. "An Overview: Segmentation Method for Blood Cell Disorders" 5th Kuala Lumpur International Conference on Biomedical Engineering 2011 Springer Berlin Heidelberg, 2011
- [15] Mahmood, Nasrul Humaimi, and Muhammad Asraf Mansor. "Red blood cells estimation using Hough transform technique." Signal & Image Processing: An International Journal (SIPIJ) 3.2 (2012): 53-64.
- [16] 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.
- [17] 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
- [18] 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.
- [19] Sharif, J. MISWAN, et al. "Red blood cell segmentation using masking and watershed algorithm: A preliminary study." Biomedical Engineering (ICoBE), 2012 International Conference on. IEEE, 2012.