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INTRODUCTION

Analysis of microscopic images of blood smears for parasiteinfected red blood cells can assist with screening and monitoring malarial infection. Millions of blood slides are manually screened for parasites every year, which is a time consuming and subjective process. We have developed a smartphone-based software to perform this task using machine learning and image analysis methods for counting infected red blood cells automatically. The software runs on a standard Android smartphone attached to a microscope by a low-cost adapter. Images of thin blood smear slides are obtained through the eyepiece of the microscope using the smartphone's built-in camera. The method we have implemented first needs to detect and segment red blood cells. However, the presence of white blood cells (WBCs) is adversely affecting the accuracy of red blood cell detection and segmentation since WBCs are often mistaken for red blood cells by current automatic cell detection methods. As a result, a pre-processing step for WBC elimination is necessary. Segmentation of WBCs is a complex process due to the morphological diversity of WBCs and staining differences. We propose a novel method for white blood cell segmentation in microscopic images of blood smears that combines a WBCs detection method based on range filtering with a customized level-set algorithm to estimate the boundary of each WBC in an image.

WHITE BLOOD CELLS

- Also known as "Leukocytes"
- > Nucleated cells in the blood that protect the body against infectious disease
- > Five main types:





Monocytes



Segmentation challenges in blood smear images

Cell staining differences

- Uneven illumination
- Cell texture complexity
- Shape diversities
- Size variation



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Value
96.37
98.37
97.36
-

Measure	Value
Jaccard Index	82.28
Dice Index	78.33



We introduced an algorithm that successfully detects white blood cells in microscopic images of thin blood smears following a range filtering approach. This method is capable of detecting different types of white blood cells with different shades of staining. The proposed algorithm accurately estimates the boundary of each cell utilizing a level-set algorithm. The results demonstrate that our algorithm provides promising segmentation outcomes. In future work, we will add this algorithm to our current smartphone application for malaria cell counting.

hematology/white-blood-cells [2] M. Poostchi, Stefan Jaeger, et.al., "Image Analysis of Blood Slides for Automatic Malaria Diagnosis", NIH-IEEE Strategic Conference on Healthcare Innovations and Point-of-Care Technologies for Precision Medicine (HI-POCT), 2015 [3]Z. Liang, A. Powell, I Ersoy, M. Poostchi, K. Palaniappan, "CNN-Based Image Analysis for Malaria Diagnosis", IEEE International Conference on Bioinformatics and Biomedicine (BIBM), 2016. [4] S. Jaeger, K. Silamut, H. Yu, M. Poostchi, I. Ersoy, A. Powell, Z. Liang, M. Hossain, S. Antani, K. Palaniappan, R. Maude, G. Thoma, "Reducing the Diagnostic Burden of Malaria Using Microscopy

Image Analysis and Machine Learning in the Field". Annual Meeting of the American Society of Tropical Medicine & Hygiene (ASTMH), Atlanta, USA, 2016.





RESULTS (cont.)

> The outcome of the proposed method for sample images from our dataset is demonstrated below.

CONCLUSIONS

REFFERENCES

[1] <u>http://www.uwosh.edu/med_tech/what-is-elementary</u>

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