A dataset for automatic contrast enhancement of microscopic malaria infected blood RGB images

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A B S T R A C T

In this article we introduce a malaria infected microscopic images dataset for contrast enhancement which assist for malaria diagnosis more accurately. The dataset contains around two hundred malaria infected, normal, species and various stages of microscopic blood images. We propose and experimentally demonstrate a contrast enhancement technique for this dataset. This simple technique increases the contrast of an image and hence, reveals significant information about malaria infected cells. Experiments on the dataset show the superior performance of our proposed method for contrast enhancement of malaria microscopic imaging.

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1. Data

The dataset in this article describes the malaria infected cells microscopic imaging with various contrast environments. It is difficult to identify the infected cells through automation and manually when the image having low contrast which leads to false diagnosis [2,4]. Fig. 1 describes the low contrast malaria infected microscopic images. Fig. 2 describes the low contrast malaria infected microscopic image and enhanced image by proposed method. Fig. 3 describes the histograms of original and enhanced RGB images. Fig. 4 describes contrast enhancement of low contrast malaria infected images at various stages. Fig. 5 displays comparison of proposed method with other existing methods on contrast enhancement of malaria microscopic imaging.

2. Experimental design, materials and methods

This section brings a brief explanation about the data set used, experiment processing and methodology of the proposed method for contrast enhancement of microscopic blood imaging.

2.1. Data set and experiment processing

The microscopic malaria images dataset for automatic contrast enhancement was performed using MATLAB 7.10.0 (R2010a) Programming software on a personal computer with an AMD Phenom II N830 triple-core processor 2.10 GHz, 3 GB system memory and 64-bit windows-7 operating system. The dataset from the Centers for Disease Control and Prevention (CDC) [https://www.cdc.gov/dpdx/malaria/index.html#tabs-1-2] have been used for the experiments [1]. Few more malaria images in dataset were supplied by Dr. Ashok K. Maiti, Department of Pathology, Midnapur Medical College & Hospital, Midnapur, West Bengal, India and few tested images from research articles [2,8,11].
2.2. Methods

Microscopic blood images are commonly acquired using a digital camera with a blood smear attachment. The input blood color image $f(x, y)$ of size $m \times n$ has three channels Red, Green and Blue, is denoted by the following mathematical expression.

$$f(x, y) = \begin{pmatrix} f_R(x, y) \\ f_G(x, y) \\ f_B(x, y) \end{pmatrix}$$

where, $(x, y) \in \{0, 1, 2, \ldots, m - 1\} \times \{0, 1, 2, \ldots, n - 1\}$
To transfer the image \( f(x, y) \) into an image \( g(x, y) \) so that it retains all the relevant information of the original image to improve the contrast of the image. Therefore, the image \( g(x, y) \) is contrast enhanced image which is considered as the standard version of the original image for further processing which assisting to improve the performance of the diagnosis. The three channels in the contrast enhanced

![Fig. 4. First row indicates low contrast malaria microscopic images at various stages (Ring, Gametocyte, Trophozoite) from dataset and second row indicates the contrast enhancement of the first row images by proposed method.](image)

![Fig. 5. Enhanced results of a low contrast color microscopic malaria image using different methods. (a) Original image (b) HE (c) CLAHE and (d) proposed.](image)
image $g(x, y)$ of an image $f(x, y)$, are constructed by using the following proposed mathematical equation [2,3].

\[
g(x, y) = \begin{pmatrix} g_R(x, y) \\ g_G(x, y) \\ g_B(x, y) \end{pmatrix} = D \begin{pmatrix} \frac{f_R^{\text{max}}(x, y) - f_R^{\text{min}}(x, y)}{f_R^{\text{max}}(x, y) + f_R^{\text{min}}(x, y)} \\ \frac{f_G^{\text{max}}(x, y) - f_G^{\text{min}}(x, y)}{f_G^{\text{max}}(x, y) + f_G^{\text{min}}(x, y)} \\ \frac{f_B^{\text{max}}(x, y) - f_B^{\text{min}}(x, y)}{f_B^{\text{max}}(x, y) + f_B^{\text{min}}(x, y)} \end{pmatrix}^T + \begin{bmatrix} \delta_R \\ \delta_G \\ \delta_B \end{bmatrix}
\]

where $D(.)$ is a diagonal matrix, the operator $\otimes$ is a matrix multiplication and $T$ indicates transpose of a matrix. In the above equation, the parameter $\delta_\varnothing$ is to control the level of contrast and is obtained by using the equation (3). Where, $\varnothing \in \{R, G, B\}$

\[
\begin{bmatrix} \delta_R \\ \delta_G \\ \delta_B \end{bmatrix} = \begin{pmatrix} 9g_R^{\text{min}}(x, y) - f_R^{\text{max}}(x, y) \\ 5f_R^{\text{max}}(x, y) + 5f_R^{\text{min}}(x, y) \\ 7f_G^{\text{max}}(x, y) - f_G^{\text{max}}(x, y) \\ 4f_G^{\text{max}}(x, y) + 4f_G^{\text{min}}(x, y) \\ 9f_B^{\text{min}}(x, y) - f_B^{\text{max}}(x, y) \\ 5f_B^{\text{max}}(x, y) + 5f_B^{\text{min}}(x, y) \end{pmatrix}
\]

The values for $f_\varnothing^{\text{max}}(x, y)$ and $f_\varnothing^{\text{min}}(x, y)$ are defined by.

\[
f_\varnothing^{\text{max}}(x, y) = \max\{f_\varnothing(x, y) : 0 \leq x \leq m - 1, 0 \leq y \leq n - 1\}
f_\varnothing^{\text{min}}(x, y) = \min\{f_\varnothing(x, y) : 0 \leq x \leq m - 1, 0 \leq y \leq n - 1\}
\]

In Assessment of visual quality, Fig. 2 (a) shows the low contrast image from dataset and Fig. 2 (b) shows the contrast enhanced image. By observing the resultant image, we can find that the proposed method yields a better and good contrast image which assists diagnosis. Beside, the histograms for the original image and enhanced images as shown in Fig. 3. The contrast enhancement of various stages i.e., Ring, Gametocyte, Trophozoite of malaria low contrast microscopic images from dataset are shown in Fig. 4. The median filter is used to reduce the noise before
enhancement [5]. In order to evaluate the performance of the proposed method with other existing methods quantitatively by edge-based contrast measure (EBCM) [10–15]. The EBCM for original image \( f(x, y) \) as shown in Eqn. (4).

\[
EBCM[f(x, y)] = \frac{\sum_{x=1}^{m} \sum_{y=1}^{n} C(x, y)}{\sum_{k=0}^{l-1} H(k)} = \frac{1}{mn} \sum_{x=1}^{m} \sum_{y=1}^{n} C(x, y)
\] (4)

Analogous definition can be given for enhanced image \( g(x, y) \).

In this study, the proposed technique is be compared with some other existing contrast enhancement methods, which includes Histogram Equalization (HE) and Contrast Limited Adaptive Histogram Equalization (CLAHE) [6–9]. Table 1 shows EBCM for the tested images using various methods. Table 1 reveals that the EBCM value of the proposed method have higher values than original image and ensure for good and natural enhancement of image when compared to other methods. The contrast enhancement results of the proposed with existing standard methods HE, CLAHE resultant images are shown in Fig. 5. The proposed yields better contrast enhancement when compared to the existing methods. The proposed method works well when an image suffers from artifacts and noise.

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Conflict of Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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| Image ID | Original | HE | CLAHE | Proposed |
|----------|----------|----|-------|----------|
| 1        | 246.06   | 179.82 | 224.41 | 249.71   |
| 2        | 141.05   | 133.78 | 125.22 | 252.43   |
| 3        | 152.34   | 144.17 | 156.29 | 252.85   |
| 4        | 230.61   | 232.26 | 167.43 | 244.78   |
| 5        | 243.28   | 213.02 | 243.211 | 244.67   |

Note: More than original image value indicates better enhancement performance.
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