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An Automated Blood Vessel Segmentation Algorithm Using Histogram Equalization and Automatic Threshold Selection

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This paper focuses on the detection of retinal blood vessels which play a vital role in reducing the proliferative diabetic retinopathy and for preventing the loss of visual capability. The proposed algorithm which takes advantage of the powerful preprocessing techniques such as the contrast enhancement and thresholding offers an automated segmentation procedure for retinal blood vessels. To evaluate the performance of the new algorithm, experiments are conducted on 40 images collected from DRIVE database. The results show that the proposed algorithm performs better than the other known algorithms in terms of accuracy. Furthermore, the proposed algorithm being simple and easy to implement, is best suited for fast processing applications.

KEY WORDS: Diabetic retinopathy, blood vessel segmentation, automatic thresholding, histogram equalization

INTRODUCTION

ue to the rapid development in computing technology and techniques, algorithms that support automated medical diagnosis have been gaining importance. Retinal vasculature has received attention by specialists in different pathologies, where the detection and analysis of retinal vasculature may lead to early diagnosis and prevention of several diseases, such as hypertension, diabetes, arteriosclerosis, cardiovascular disease and stroke.¹ One of the well-known and commonest diseases that need a computer-aided medical diagnosis is diabetic retinopathy (DR), which leads in most cases to partial or even complete loss of visual capability.² The accurate diagnosis of this disease depends upon some features which have to be analyzed in order to quantify the severity level of the disease. Retinal blood vessels are considered as one of the most important features for the detection of DR. As

diabetic retinopathy is a progressive disease, regular screening of the human retina is essential for reducing the proliferative diabetic retinopathy and for preventing the subsequent loss of visual capability. The screening should be done every 6 months, which includes obtaining and analyzing a sequence of fundus images and observing the early changes in blood vessel patterns as well as the presence of microaneurysms.^{3–6}

In the literature, a number of algorithms for automated blood vessel segmentation have been reported. These algorithms fall into several categories, such as matched filter based,^{7,8} tracking based,^{9–12} threshold probing based,¹³ model based,^{14,15} neural network based,¹⁶ and pattern recognition based.^{17,18}

This paper presents a blood vessel segmentation algorithm which takes advantage of simple and powerful image processing techniques. For instance, as the contrast between the blood vessels (foreground) and the retinal tissue (background) is generally poor in the fundus images, an effective technique called contrast-limited adaptive histogram equalization (CLAHE)¹⁹ is utilized for contrast enhancement by limiting the maximum slope in the transformation function. Another technique, called *Isodata*²⁰ that provides an auto-

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matic threshold value for binarization is also employed. Furthermore, the proposed algorithm is characterized by low processing time thus making it suitable for fast processing applications.

The remainder of the paper is organized as follows: "Overview of the Proposed Methodology" presents an overview of the proposed algorithm, "Results and Discussion" presents the results and discussion and "Conclusion" provides conclusions.

OVERVIEW OF THE PROPOSED METHODOLOGY

Database

Our algorithm has been tested over sets of fundus images collected from publicly available database, called DRIVE database.²¹ The DRIVE database contains 40 test images, compressed in JPEG format of size 565×584 pixels obtained from a diabetic retinopathy screening program. The images are acquired using a Canon CR5 non-mydriatic 3CCd camera at 45° field of view.

The 40 images were divided into two sets, a test set and a training set, each containing 20 images. The images have been manually segmented by three observers to be used as references for comparing the computer-generated segmentations. For each image in test set, two manual segmentations (first and second manuals) are available, whereas for an image in the training set, only one manual segmentation is available.

Proposed Methodology

The proposed algorithm is designed for retinal blood vessels segmentation. Input to the system is a color fundus image of human retina acquired by a fundus camera and the output is a binary image which contains only the blood vessels. The main modules of the algorithm are: Color image (RGB) to gray/green conversion, contrast enhancement, background exclusion, and thresholding and postfiltration. Figure 1 shows the block diagram of the proposed algorithm.

RGB to Gray/Green Conversion

Color fundus image is first converted into a gray-scale/green-channel image in order to facili-

tate the blood vessels segmentation and to decrease the computational time. Gray-scale image provides only the luminance information from the color image after eliminating the hue and saturation, while the green-channel image provides maximum local contrast between the background and foreground.^{22,23} The segmentation results are obtained using gray and green-channel images and they are then compared with the results obtained using other known algorithms. The conversion from the color image to gray-scale image is done by forming a weighted sum of the RGB components, as in Eq. (1):

$$g = 0.2989 * R + 0.5870 * G + 0.1140 * B$$
(1)

where R, G, and B represent the red, green, and blue components respectively. Figure 2 shows an RGB image and the extracted gray-scale and green-channel images.

Contrast Enhancement

Low contrast images could occur often due to several reasons, such as poor or non-uniform lighting condition, nonlinearity or small dynamic range of the imaging sensor, i.e., illumination is distributed non-uniformly within the image. Therefore, it is necessary to deepen the contrast of these images to provide a better transform representation for subsequent image analysis steps.^{24,25} The contrast stretching process applied to the grayscale/green-channel image is illustrated in Figure 3, where T(r) represents the transformation function controlled by the locations of points (r_1, s_1) and (r_2, s_2) . A wide variety of techniques is employed to improve the contrast by stretching the range of intensity values of the image so that the full dynamic range of the image is covered. Techniques such as decorrelation stretch transform, unsharp mask, histogram equalization, adaptive histogram equalization (AHE), contrast-limited adaptive histogram equalization (CLAHE) are used for enhancing the image contrast. In the proposed algorithm, CLAHE technique is adopted to perform the contrast enhancement.

CLAHE is a widely used contrast enhancement technique which has proved itself to be very effective for medical images. This technique enhances the contrast adaptively across the image



Fig. 1. Block diagram of the proposed algorithm.

by limiting the maximum slope in the transformation function. Instead of applying the histogram equalization on the entire image, it is applied only on small non-overlapping regions in the image. Then, the neighboring tiles are combined using bilinear interpolation to reduce induced boundaries.^{26,27} In our implementation, the size of each region was 8×8 pixels with 128 bins for the histogram. Figure 4 shows the results of applying the CLAHE on the gray-scale and green-channel images.

Background Exclusion

The main purpose of this step is eliminating background variations in illumination from an image so that the foreground objects may be more easily analyzed. In the proposed algorithm, the background exclusion is performed by subtracting the original intensity image from the averagefiltered image. Average filter is one of the simplest local operations over an image, which is also called as "neighborhood average method". The essential idea of a standard moving average filter is to replace the value of the center pixel $\hat{g}(x, y)$ by the average value of a predefined number of neighboring pixels $g_i(x, y)$ as shown in Eq. (2).^{28,29}

$$\widehat{g}(x,y) = \frac{1}{N \times M} \sum_{i=1}^{N \times M} g_i$$
(2)

In the proposed algorithm, a window size $(M \times N)$ of 9×9 pixels is used for implementing the average filter. Figure 5 shows the results of applying the average filter on the gray-scale and green-channel images.

The original image g is subtracted from the average-filtered image \hat{g} and the result is shown in Figure 6. Mathematically, the difference image h(x, y) between two images \hat{g} and g, is generated by computing the difference 't' between all pairs of



Fig. 2. a Color retinal image, b gray-scale image, c green-channel image.



Fig. 3. Explanatory illustration of contrast stretching transformation.

corresponding pixels in $\hat{g}(x, y)$ and g(x, y),³⁰ as shown in Eq. (3).

$$h(x,y) = \begin{cases} t & \text{if } \widehat{g}(x,y) - g(x,y) > 0\\ 0 & \text{Otherwise} \end{cases} \quad 0 < t \le 255 \end{cases}$$
(3)

Thresholding and Post-Filtration

The aim of this module is to produce a binary image in which the value of each pixel is either 1 (blood vessel) or 0 (background). Unfortunately, there exists no thresholding technique for determining a unique threshold value which will provide perfect results in all cases. However, in the proposed algorithm, we made use of the socalled *Isodata* technique which provides an automatic threshold value for producing a binary image *B*. This technique divides the histogram into two parts, P_1 and P_2 using an initial threshold value T_0 . Subsequently, the mean values μ_1 and μ_2 of both the parts are calculated, and a new threshold value is determined which represents the average of μ_1 and μ_2 , as shown in Eqs. (4) and (5). This process is repeated iteratively until the threshold values T_k and T_{k-1} , converge (i.e., $T_k \approx T_{k-1}$), where:

$$T_0 = (\min(h) + \max(h))/2$$
 (4)

$$T_k = \frac{\mu_1 + \mu_2}{2}$$
(5)

The gray-scale/green-channel images h(x, y) are then converted to binary images based on the threshold values T_k using Eq. (6). The resulting binary images are shown in Figure 7a, b, respectively.

$$B(x,y) = \begin{cases} 1 & \text{if } h(x,y) \ge T_k \\ 0 & \text{Otherwise} \end{cases}$$
(6)

As a result of the thresholding, some unwanted pixels would appear as noise (false positive) in the resultant binary image, and therefore, some post-processing should be adopted for refining the image and retaining the desired objects. For this purpose, a morphological operation (opening)²⁶ is employed to remove the undesired objects that



Fig. 4. Results of contrast enhancement a Gray-scale image, b Green-channel.



Fig. 5. Average-filtered images with a 9×9 pixels a Gray-scale image, b Green-channel.

have fewer than 35 pixels. The bright circle corresponding to the edge of retina can also be removed by subtracting a mask image I_{mask} which is shown in Figure 7c from the binary image, as illustrated in Eqs. 7–10.

Let:
$$I_{rgb} = R + G + B$$
 (7)

$$I_{bw} = \operatorname{inv}(I_{rgb} > 100) \tag{8}$$

$$I_{\text{mask}} = \delta_{SE}(I_{bw}) \quad (SE = 3 \times 3 \text{ pixels})$$
 (9)

$$BW(x,y) = \begin{cases} 1 & \text{if } B - I_{\text{mask}} > 0\\ 0 & \text{Otherwise} \end{cases}$$
(10)

The resulting gray-scale and green-channel images after post-filtration are shown in Figures 7d, e, respectively.

RESULTS AND DISCUSSION

The experiments are carried out using the same values for all of the parameters mentioned in the different steps of the proposed algorithm. All the 40 images of the DRIVE database (20 training images and 20 test images) are used to evaluate the performance of the proposed algorithm. A number of criteria, namely, accuracy (Acc.),³¹ true-positive fraction (TPF), false-positive fraction (FPF), area under the ROC curve (A_z) ,³² and Kappa statistics $(k)^{33}$ are used to evaluate the performance of the proposed algorithm. The accuracy is computed by the ratio of the total number of correctly classified



Fig. 6. The resultant image of background exclusion a Gray-scale image, b Green-channel.



Fig. 7. a Gray-scale binary image after thresholding b green-channel binary image after thresholding c mask image *l*_{mask} d gray-scale binary image after post-filtration e green-channel binary image after post-filtration.

points to the number of points in the image. TPF denotes the fraction of pixels correctly classified as blood vessel pixels, while FPF denotes the fraction of pixels erroneously classified as blood vessel pixels. TPF and FPF are calculated using Eqs. 11 and 12, respectively.

$$TPF = \frac{TP}{TP + FN}$$
(11)

$$FPF = \frac{FP}{FP + TN}$$
(12)

where true positive (TP) refers to positive pixels correctly labeled as positive. False positive (FP) refers to negative pixels incorrectly labeled as positive. False negative (FN) refers to positive pixels incorrectly labeled as negative. Finally, True Negative (TN) refers to negative pixels correctly labeled as negative. Area under the ROC curve is obtained by plotting the TPF against the FPF³². The Kappa coefficient (k) is used to estimate the agreement between the automated- and manually segmented blood vessels which is calculated using Eq. (13).³³

$$k = \frac{\Pr(a) - \Pr(e)}{1 - \Pr(e)} \tag{13}$$

where Pr(a) and Pr(e) represent the observed and chance agreements respectively. The average values of the Acc., TPF, FPF, A_z and k are obtained for the proposed algorithm using DRIVE database images. Table 1 compares the results obtained using the proposed algorithm with those obtained by other known algorithms.

Based on the experimental results presented in Tables 1, it is clear that the proposed algorithm yields superior results compared with other known algorithms with respect to the accuracy and TPF values. The FPF values obtained by the proposed algorithm are comparable to the values

Table 1. Fertilinance of Different Algorithms for blood Vesser organization of Differe Database					
Method	Acc.	TPF	FPF	A_z	k
2nd Human observer ²⁰	0.9473	0.7761	0.0275	-	-
Mendonca (gray-scale) ³¹	0.9463	0.7315	0.0219	-	_
Mendonca (green- channel) ³¹	0.9452	0.7344	0.0236	_	-
Staal ^{20,34}	0.9442	0.7194	0.0227	-	-
Niemeijer ^{20,35}	0.9417	0.6898	0.0304	_	-
RGB-Q ³⁶	_	0.7704	0.0693	_	-
G-Q ³⁶	_	0.7500	0.0732	-	_
Proposed algorithm (gray-scale)	0.9554	0.8303	0.0308	0.8865	0.7298
Proposed algorithm (green-channel)	0.9630	0.8423	0.0342	0.8987	0.7419

Table 1. Performance of Different Algorithms for Blood Vessel Segmentation on DRIVE Database

obtained by other methods. Figure 8 shows some sample results obtained using the proposed algorithm.

Since the proposed algorithm makes use of only simple computing techniques, the processing time for this algorithm will be significantly less compared with the algorithms which make use of computationally intensive techniques such as neural networks and region-growing. Hence this algorithm is best suited for fast-processing screening applications. Using MATLAB 7.6 with a P. IV/CPU 2.80 GHz/2 GB of RAM, the proposed algorithm took an average time of approximately 0.3 s per image.



Fig. 8. Random samples of original and segmented images: a and b Green-channel, c and d gray-scale.

CONCLUSION

In this paper, a simple and computationally efficient algorithm for retinal blood vessel segmentation has been presented. The proposed algorithm has employed modules such as contrast enhancement, background exclusion and thresholding. Experimental results obtained by using gray-scale as well as green-channel images have been presented. For binarization, an automatic thresholding technique, called Isodata has been employed. The performance of the proposed algorithm has been tested using DRIVE database images. From the experimental results, it is found that the proposed algorithm yields superior results compared with other known algorithms with respect to the accuracy and TPF values. Also, since the proposed algorithm makes use of only simple and computationally less intensive processing steps, it is best suited for fast processing applications.

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