

Automated detection of photoreceptors in *in-vivo* retinal images

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Abstract

The inclusion of adaptive optics (AO) into ophthalmic imaging technology has allowed the study of histological elements of retina *in-vivo*, such as photoreceptors, retinal pigment epithelium (RPE) cells, retinal nerve fiber layer and ganglion cells. The high-resolution images obtained with ophthalmic AO imaging devices are rich with information that is difficult and/or tedious to quantify using manual methods. Thus, robust, automated analysis tools that can provide reproducible quantitative information about the tissue under examination are required. Automated algorithms have been developed to detect the position of individual photoreceptor cells and characterize the RPE mosaic. In this work, an algorithm is presented for the detection of photoreceptors. The algorithm has been tested in synthetic and real images acquired with an Adaptive Optics Scanning Laser Ophthalmoscope (AOSLO) and compared with the one developed by Li and Roorda. It is shown that both algorithms have similar performance on synthetic and cones-only images, but the one here proposed shows more accurate measurements when it is used for cones-rods detection in real images.

Keywords: Photoreceptor; adaptive optics; image processing.

Detección automatizada de fotorreceptores en imágenes retinianas *in-vivo*

Resumen

La inclusión de la óptica adaptativa (adaptive optics, AO) en la tecnología de imágenes oftálmicas ha permitido el estudio *in-vivo* de los elementos histológicos de retina, como los fotorreceptores, células del epitelio pigmentario de la retina (retinal pigment epithelium, RPE), la capa de fibras nerviosas de la retina y células ganglionares. Las imágenes de alta resolución obtenidas con dispositivos oftálmicos con AO son ricas en información, que es difícil y/o tediosa de cuantificar por medio de métodos manuales. Por lo tanto, se requieren herramientas de análisis automatizadas robustas que puedan proporcionar información cuantitativa reproducible del tejido bajo examen. Algoritmos automatizados han sido desarrollados para detectar la posición de células individuales fotorreceptoras y caracterizar el mosaico RPE. En este trabajo, se presenta un algoritmo para la detección de los fotorreceptores. El algoritmo ha sido probado en imágenes sintéticas y reales adquiridas con un oftalmoscopio de barrido láser con óptica adaptativa (Adaptive Optics Scanning Laser Ophthalmoscope, AOSLO) y comparado con el desarrollado por Li y Roorda. Se muestra que ambos algoritmos tienen un rendimiento similar en imágenes sintéticas e imágenes con sólo conos, pero el algoritmo propuesto muestra mediciones más precisas cuando se utiliza para la detección de conos-bastones en imágenes reales.

Palabras clave: Fotorreceptores; óptica adaptativa; procesamiento de imágenes.

1. Introduction

Typically, the retina is shown to have several layers: inner limiting membrane, retinal nerve fiber layer (RNFL), retinal ganglion cell layer, inner plexiform layer, inner nuclear layer,

outer plexiform layer, outer nuclear layer, external limiting membrane, photoreceptor layer and retinal pigment epithelium (RPE) layer [1]. The RNFL conducts signals from the eye. The ganglion cell layer contains the cell bodies of ganglion cells and displaced amacrine cells; the ganglion cell

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dendrites are in the plexiform layer, where they receive bipolar cell and amacrine cell inputs, while their axons are in the RNFL. The photoreceptors are responsible for visual transduction, the transformation of light energy into electrical energy, a process called isomerization [2,3]. Finally, the RPE is vital for the maintenance of photoreceptor function, since it contains nutrients and enzymes [4].

In the retina, there are two types of photoreceptors: cones and rods. Structurally, the photoreceptors are different in shape, as their names indicate, and in size, rods being longer than cones, and cones wider than rods; they also have different functional properties that affect our perception. Cones cover all the retina, although higher density is in the fovea, where there are no rods. There are about 120 million rods, and they are in the periphery of the retina [2,4,5].

It is very important to know where the photoreceptors, cones and rods, are located and how many photoreceptors there are in the retina, which could be used to monitor the evolution of therapies for ocular globe diseases. A method previously used for cell detection and their characterization is shown in [6]; it uses adaptive optics scanning light ophthalmoscopy (AOSLO), which has become an important tool for the study of retinal diseases [6-22]. Our interest is primarily in detecting the cones and rods. Since changes in the photoreceptor mosaic may precede cell death, it is possible that morphometric changes can be measured before larger damages are observed, allowing early therapeutic intervention.

Accuracy in the detection and classification of patterns in biomedical images is central to detecting and monitoring tissue damage, as well as to quantifying its extent. Hence, it is necessary to have robust and reliable methods for classifying and quantifying retinal structures in in-vivo retinal images. This is the reason why, in recent years, there has been an increased attention to this area [6-8,22-29], and it spurred our interest in developing a more efficient algorithm for the detection of retinal cones and rods. The algorithm that we propose quantifies and locates the photoreceptors using the local maximum value, maximum absolute deviation and binary operations.

This paper is organized as follows. In section 2, the methods used for the development of the algorithm, as well as the generation of synthetic and real AO images, are discussed. In section 3, the comparison between the algorithm proposed by Li and Roorda and the one here presented is undertaken. Finally, the main discussion and conclusions are in section 4.

2. Method

2.1. Algorithm

The aim of the algorithm is to quantify and locate photoreceptors in in-vivo retinal images. Fig. 1 shows the schematic representation of the algorithm.

2.1.1. Adaptive threshold using local maximum values

Let $I(x,y)$ be a bidimensional intensity function (a matrix) where x and y are the spatial coordinates of a pixel; let $I_{max}(x,y)$ be the local maximum value within a window of 3×3 pixels, being:

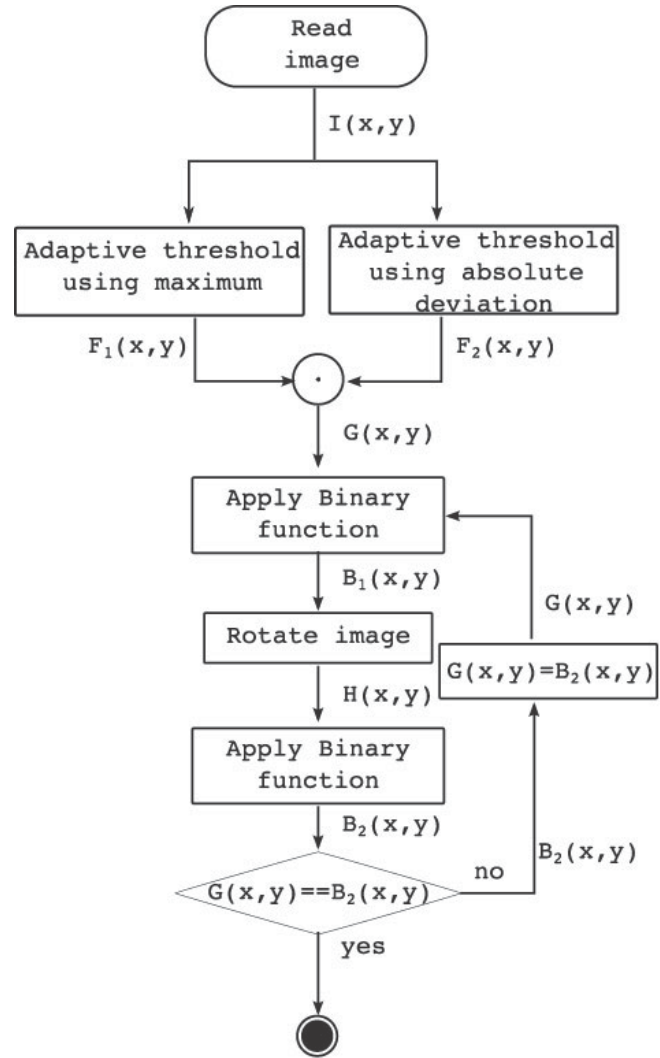


Figure 1. Schematic representation of the algorithm
Source: The authors

$$I_{max}(x,y) = \max\{I(x+s,y+t) | -1 \leq s \leq 1 \text{ and } -1 \leq t \leq 1\} \quad (1)$$

Thereby, $F_1(x,y)$ is defined as follows:

$$F_1(x,y) = \begin{cases} 1 & \text{if } I(x,y) == I_{max}(x,y) \\ 0 & \text{if } I(x,y) \neq I_{max}(x,y) \end{cases} \quad (2)$$

2.1.2. Adaptive threshold using absolute deviation

Let $M(x,y)$ be the maximum absolute deviation (Eq. 3), which determines the sufficient variance in intensity, with n being the number of elements, and let $I_{max}(x,y)$ be the local maximum value (Eq. 1). Then, let m be a threshold that determines the sufficient variance in intensity of $I(x,y)$.

$$M(x,y) = \left(\sum_{s=-1}^1 \sum_{t=-1}^1 |I(x+s,y+t) - I_{max}(x,y)| \right) / n \quad (3)$$

Thereby, $F_2(x,y)$ is defined as shown in Eq. 4.

$$F_2(x, y) = \begin{cases} \mathbf{1} & \text{if } M(x, y) \geq m \\ \mathbf{0} & \text{if } M(x, y) < m \end{cases} \quad (4)$$

The next step in this stage is the generation of a binary image $G(x, y)$ using Eq. 5, through the logic operation AND of the results from Eqs. 2 and 4.

$$G(x, y) = F_1(x, y) \cdot F_2(x, y) \quad (5)$$

2.1.3. Binary operations

Eq. 6 is a binary function that works within a window of 3 X 3 pixels; it was designed to reduced over-detection of cells, where over-detection of a cell can be represented as elements a_1, a_2, \dots, a_{33} shown in Figs. 2. Element a_1 from Fig. X can be represented by a binary function $\mathbf{a}_1 = \overline{i_{11}} + i_{12} + i_{13} + i_{21} + \overline{i_{22}} + i_{23} + i_{31} + i_{32} + i_{33}$; where $i_{11} = I_{(x-1, y-1)}$, $i_{12} = I_{(x, y-1)}$, $i_{13} = I_{(x+1, y-1)}$, $i_{21} = I_{(x-1, y)}$, $i_{22} = I_{(x, y)}$, $i_{23} = I_{(x+1, y)}$, $i_{31} = I_{(x-1, y+1)}$, $i_{32} = I_{(x, y+1)}$ and $i_{33} = I_{(x+1, y+1)}$; which means that the central element i_{22} is eliminated to avoid over-detection. Thus, every element from Fig. 2 can be represented by a binary function where the central element i_{22} has been eliminated, and it is based on that fact that every element can be represented by a binary function, and the combination of all these functions yields cell location. This equation was reduced by the Quine-McCluskey method; afterward, Postulates and Theorems of Boolean Algebra were used to minimize it. This binary function is applied onto the image $I(x, y)$. Later, the obtained image is rotated 180 degrees, $H(x, y) = \text{rotate180}(B(x, y))$, with the goal of eliminating the multiple detection in the opposite direction. Then, Eq. 6 is applied onto the rotated image $H(x, y)$. Finally, the result is rotated back to the original orientation. This process is performed until algorithm convergence is reached.

$$B(x, y) = i_{22} \cdot [\overline{i_{11}} + i_{13} + i_{23} + i_{33} + i_{32} + i_{31}] \cdot [i_{11} + \overline{i_{13}} + i_{32} + i_{31} + i_{21} + (i_{33} \cdot \overline{i_{23}})] \cdot [i_{13} + i_{23} + i_{33} + \overline{i_{21}} + (i_{12} \cdot \overline{i_{31}}) + (\overline{i_{11}} \cdot i_{32})] \cdot [\overline{i_{12}} + i_{33} + i_{32} + i_{31} + (\overline{i_{11}} \cdot i_{23}) + (\overline{i_{13}} \cdot i_{21})] \cdot [i_{11} + i_{12} + \overline{i_{23}} + \overline{i_{33}} + i_{31} + i_{21} + (\overline{i_{13}} \cdot i_{32})] \cdot \{i_{11} + i_{12} + i_{13} + \overline{i_{32}} + (i_{23} + \overline{i_{31}}) \cdot [\overline{i_{33}} + i_{21} + (i_{23} \cdot \overline{i_{31}})]\} \quad (6)$$

2.2. Synthetic images

The synthetic images were generated using a pre-determined number of pseudo-photoreceptors, which in turn were generated through the use of a Gaussian function and an elliptical one [30]. The pseudo-photoreceptors varied at random in terms of size and reflectance, and they were symmetrically (in equidistant square and hexagonal arrays) and randomly distributed. Background illumination was added to each of the generated images, which corresponds to the illumination in the real images acquired with the AOSLO utilized at the University of Rochester [18].

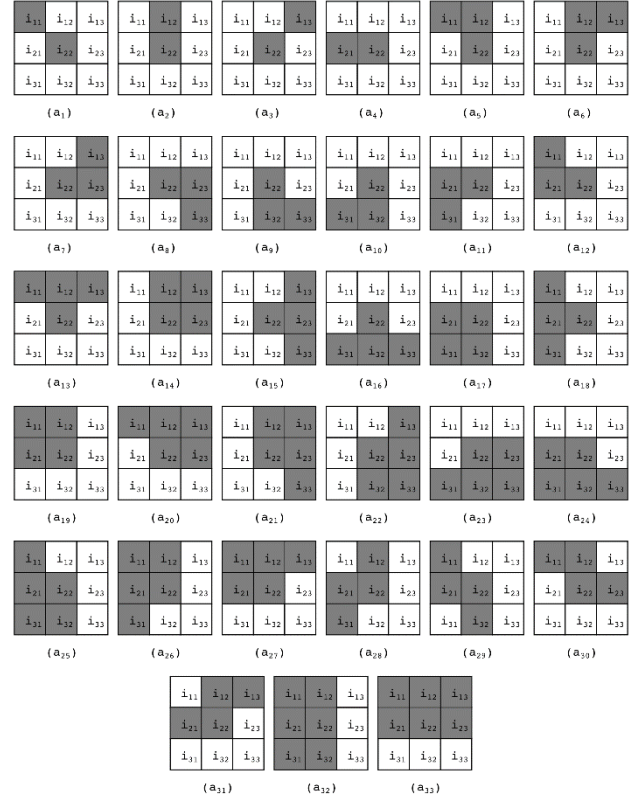


Figure 2. Masks used to obtain the binary function $B(x, y)$, Eq. 6. Source: The authors

2.3. Real AO images

We used AOSLO images obtained for current and previous experiments at the Center for Visual Science of the University of Rochester to test the algorithm. Data shown from human participants are from experiments that were approved by the Research Subjects Review Board at the University of Rochester and adhered to the tenets of the Declaration of Helsinki. Participants gave informed written consent after the nature of the experiments and any possible risks were explained both verbally and in writing.

3. Results

The algorithm developed by Li and Roorda [6] (Algorithm 1) and the algorithm here proposed (Algorithm 2) have been tested on synthetic and real images, Fig. 3 shows the results obtained with both algorithms on synthetic images. In the caption, we show the color code that was used to identify where a pseudo-cell was positioned and the results obtained by each algorithm.

For each function-distribution combination, 1000 synthetic images of 100x100 pixels in size were generated. To those images, we applied algorithms 1 and 2, and the results obtained were later evaluated. The outcome of such evaluation is shown in Tables 1 and 2. Table 1 shows the relative error obtained for each of the combinations, whereas Table 2 shows the number of cells that were correctly detected as corresponding exactly with the place where the center of the pseudo-cells had been positioned.

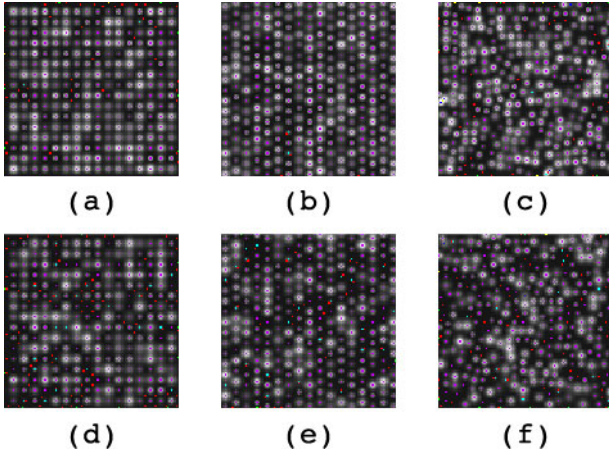


Figure 3. Synthetic images results. (a) Gaussian-Square, (b) Gaussian-Hexagonal, (c) Gaussian-Random (d) Elliptical-Square, (e) Elliptical-Hexagonal and (f) Elliptical-Random. Purple points represent correct detection for both algorithms; yellow points represent where both algorithms fault detecting a cell; cyan points represent where only algorithm 2 detected an actual result; magenta points represent where only algorithm 1 detected correctly; red points represent a wrong detection with algorithm 1; green points represent a wrong detection with algorithm 2; blue points represent non detected cells.

Source: The authors

Table 1.

Relative error obtained from synthetic images

Function	Distribution	Pseudo-cells (mean)	Relative error (%)	
			Algorithm 1	Algorithm 2
Gaussian	Square	256.00	15.18	6.10
	Hexagonal	289.00	0.41	0.42
	Random	288.14	9.36	3.15
Elliptical	Square	256.00	21.77	5.8
	Hexagonal	289.00	3.12	3.73
	Random	288.45	14.09	4.62

Source: The authors

Table 2.

Correctly detected from synthetic images

Function	Distribution	Pseudo-cells (mean)	Correctly detected (%)	
			Algorithm 1	Algorithm 2
Gaussian	Square	256.00	99.16	100
	Hexagonal	289.00	99.42	100
	Random	288.14	96.70	97.71
Elliptical	Square	256.00	91.54	100
	Hexagonal	289.00	93.59	100
	Random	288.45	92.16	97.70

Source: The authors

4. Discussion and conclusions

For all the images, both real and synthetic, a threshold $m = 2.0$ was used to calculate the maximum absolute deviation M_{ij} , and it was determined empirically; it is based on different tests that we conducted. If $m > 2.0$, we have a very prominent peak, but if $m < 2.0$, there is a tendency to flatness. The manual count presented in Table 3 was carried out on the basis of the authors' appreciation. For algorithm [6], $cut_off_frequency = 0.61\pi$ and $threshold = 0$ were used to test the all the images.

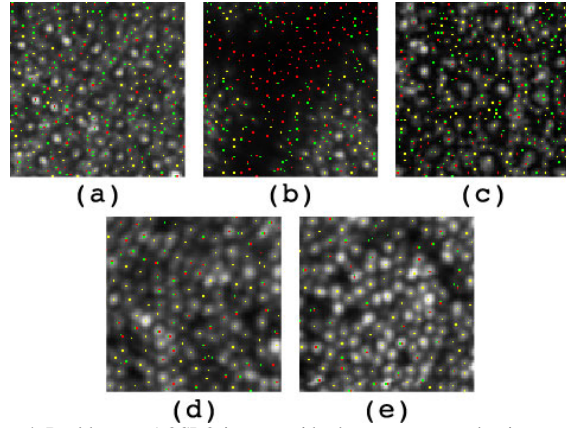


Figure 4. Real human AOSLO images with photoreceptors. red points represent the cell location detected with algorithm 1; green points represent the cell location detected with algorithm 2; and yellow points represent the result of algorithm 1 and 2.

Source: The authors

Table 3.

Cells detected in real AOSLO images

Figure	Manual	Algorithm 1	Algorithm 2
3(a)	305	219	287
3(b)	168	240	187
3(c)	279	247	295
3(d)	151	140	144
3(e)	152	144	145

Source: The authors

Fig. 4 shows the results obtained on real images. In the caption, we describe the color code used to identify the outcome of the algorithm that was utilized in each case. Table 3 shows the number of cells that were manually detected by the authors and those detected by each algorithm.

As shown in Tables 1 and 2 and in Fig. 3, both algorithms exhibited similar performances on synthetic images.

When the algorithm was applied to real images that included cones, rods and vascular tissue, Fig. 4 (a - c), the algorithm developed by Li and Roorda [6] showed detection of cells in the vascular tissue, as well as poor detection of rods. It is evident that there are photoreceptors in these zones, but since the vessels cause the photoreceptor layer to be below or above the blood vessel, it cannot be stated that photoreceptor detection in this zone is adequate.

Both algorithms exhibited similar performances on real images that included only cones, Fig. 4 (d, e). It is worth mentioning that Li and Roorda's algorithm does not show multiple detection for a single photoreceptor, whereas ours does.

As aforementioned, it is very important and useful to know where photoreceptors are located and how many photoreceptors there are in the retina, because that information could be used for the analysis of the distribution and orientation of cones and rods in-vivo as well as to monitor the evolution of therapies for retinal diseases.

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