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Weifang Zhu,† Li Zhang,† Fei Shi,† Dehui Xiang,† Lirong Wang,† Jingyun Guo,† Xiaoling Yang,† Haoyu Chen,‡§, and Xinjian Chen,‡,*

†Soochow University, School of Electronics and Information Engineering, Suzhou, China
‡Shantou University and the Chinese University of Hong Kong, Joint Shantou International Eye Center, Shantou, China
§The Chinese University of Hong Kong, Department of Ophthalmology and Visual Sciences, Hong Kong, China

Abstract. Cystoid macular edema (CME) and macular hole (MH) are the leading causes for visual loss in retinal diseases. The volume of the CMEs can be an accurate predictor for visual prognosis. This paper presents an automatic method to segment the CMEs from the abnormal retina with coexistence of MH in three-dimensional-optical coherence tomography images. The proposed framework consists of preprocessing and CME segmentation. The preprocessing part includes denoising, intraretinal layers segmentation and flattening, and MH and vessel silhouettes exclusion. In the CMEs segmentation, a three-step strategy is applied. First, an AdaBoost classifier trained with 57 features is employed to generate the initial results. Second, an automated shape-constrained graph cut algorithm is applied to obtain the refined results. Finally, cyst area information is used to remove false positives (FPs). The method was evaluated on 19 eyes with coexistence of CMEs and MH from 18 subjects. The true positive volume fraction, FP volume fraction, dice similarity coefficient, and accuracy rate for CMEs segmentation were 81.0% ± 7.8%, 0.80% ± 0.63%, 80.9% ± 5.7%, and 99.7% ± 0.1%, respectively. © 2017 Society of Photo-Optical Instrumentation Engineers (SPIE) [DOI: 10.1117/1.JBO.22.7.076014]

Keywords: optical coherence tomography; cystoid macular edema; macular hole; segmentation; graph cut.

1 Introduction

Optical coherence tomography (OCT) is becoming a mainstay in ophthalmology as a noninvasive imaging technique for human retina.† OCT has been widely used in the diagnosis of ocular diseases, such as macular edema (ME), macular hole (MH), glaucoma, etc. The macula is located near the center of the retina and is responsible for central, high-resolution, and color vision. Some acute maculopathies may lead to the loss of central vision and even blindness.† MH is a full-thickness defect of neurosensory retina involving the anatomic fovea which severely affects central visual acuity.† The overall incidence of MH is ~3.3 cases in 1000 in the United States.† MH often progresses gradually and is common in people of age 60 and over.† A person with MH may notice a slight distortion or blurriness in straight-ahead vision. Cystoid macular edema (CME) occurs in a wide variety of ocular diseases, such as diabetic retinopathy, intraocular inflammation, and central or branch retinal vein occlusion.†,§ If CME is untreated, the final result is the permanent deterioration of the central vision. Figure 1 shows a B-scan of the 3-D OCT image with CMEs and MH. Red curves show the manual segmentation of CMEs and yellow arrow shows the position of the MH. In clinical practice, if the CMEs and MH co-occur, the CMEs usually surround the MH as in Fig. 1. It has been proved that the volume of CMEs in the retina can be an accurate predictor of visual acuity.§ Furthermore, the volume of CMEs can lead to better metrics for making treatment protocols.

Therefore, automated methods for CMEs segmentation in the 3-D OCT images with coexistence of CMEs and MH have urgent needs.

There are several previous studies focused on the segmentation of CMEs. Both semiautomated and automated methods have been studied. The semiautomated methods rely on manual initialization in each B-scan to quantify the volume of CMEs. Kashani et al.11 used the OCTOR software to quantify the cystoid spaces in eyes. Fernández12 proposed a method that requires the manual initialization of the snake and the fluid regions in 3-D OCT images by thresholding and boundary tracing. Wilkins et al.13 developed a technique that utilized the computerized segmentation combined with the minimal expert interaction to quantify the volume of CMEs. Although the abovementioned methods can detect the volume of CMEs accurately, they are semiautomatic so that they are laborious and time-consuming. Several automated methods have also been proposed which can obtain satisfactory results. Wilkins et al.14 segmented the CMEs in 3-D OCT images by thresholding and boundary tracing. Roychowdhury et al.15 localized cysts in diabetic macular edema (DME) by iterative high-pass filtering with further analysis using solidity, mean, and maximum pixel value as decisive features. Sisternes et al.16 segmented the cysts in 3-D OCT images using lasso regularization-based regression and adaptive thresholding. Esmaili et al.17 proposed a 3-D curvelet transform-based dictionary learning method for automatic segmentation of cysts. Girish et al.18 proposed a marker-controlled watershed transform-based method to segment cysts on 2-D OCT B-scan images. Venhuizen et al.19 proposed a multiscale convolutional...
neural network-based method to predict if an image voxel belongs to a cyst. Wang et al.\textsuperscript{20} proposed a fuzzy level set-based method to automatically segment the intraretinal fluid and subretinal fluid in DME eyes using structural OCT and angiography OCT data. However, these methods may fail when the MH and CMEs coexist. There may be two reasons. First, most of the CMEs segmentation methods in OCT images are based on the intensity differences between CMEs and the other tissues of retina, and the intensity of MH is similar to the one of CMEs, so MH may be very likely segmented as a false positive (FP) region. Second, some of the above methods, e.g., Sisternes et al.\textsuperscript{10} use the position information about the fovea center because CMEs usually surround the fovea. When MH co-occurs, the normal shape of retina is altered and the correct position of the fovea center cannot be detected. This may be another cause of FPs. Therefore, new methods that can fully segment the CMEs in the retina with MH are urgently needed.

The automatic segmentation of the CMEs coexisting with the MH is a challenging problem. The key difficulties lie in two aspects: (1) MH is a great disturbance for CMEs segmentation, and MH cannot be easily segmented because the shapes of the MHs may be irregular and even distorted by the detached tissues, such as posterior vitreous detachment.\textsuperscript{21} (2) The number and size of CMEs are always uncertain, making it difficult to quantify the volume of the cysts.

In this paper, we propose an automated framework for intraretinal CME segmentation in 3-D OCT images with MH. The proposed method consists of two steps: preprocessing and CMEs segmentation. In the preprocessing, the operations including signal-to-noise (SNR)-balancing, 3-D curvature anisotropic diffusion filtering, layer segmentation and flattening, and MH and vessel silhouettes exclusion, are applied to the OCT scans. In the CMEs segmentation, first, an AdaBoost classifier with 57 features is employed for the coarse segmentation. Second, the morphological dilation and erosion operations are used to generate the foreground (CMEs) seeds and background seeds for the following graph cut-based segmentation method. Third, an automatic shape-constrained graph cut algorithm is applied for the fine segmentation. Finally, the FPs are eliminated according to the CMEs area-related information. The flowchart of the proposed method is shown in Fig. 2.

2 Materials and Methods

2.1 Subjects and Data Collection

The collection and analysis of image data were approved by the Institutional Review Board of Joint Shantou International Eye Center (JSIEC), Shantou University and the Chinese University of Hong Kong and adhered to the tenets of the Declaration of Helsinki. Because of its retrospective nature, informed consent was not required from subjects. The medical records and OCT database of JSIEC from December 2008 to 2013 were searched and reviewed.

Totally, 19 eyes with coexistence of CMEs and MH from 18 subjects were included and underwent macular-centered (6 × 6 mm) SD-OCT scan (Topcon 3-D OCT-1000, 512 × 32 × 480 voxels, 11.72 × 187.5 × 3.50 μm\(^3\)), 512 × 64 × 480 voxels, 11.72 × 93.75 × 3.50 μm\(^3\), or 512 × 128 × 480 voxels, 11.72 × 46.88 × 3.50 μm\(^3\)). There were 8 males and 10 females with the mean age of 45.6 ± 19.7 years (range: 9 to 66 years). Subjects with other eye diseases were excluded except for refractive error < ±6 diopter. The raw, uncompressed data were exported from the OCT machine as “.fds” format for analysis.

2.2 Ground Truth Delineation

All the CME regions in the 3-D OCT images were manually marked slice by slice by two observers independently, under the directions of an experienced ophthalmologist. The marked results marked are considered as the ground truth one (GT1) and ground truth two (GT2), respectively.

2.3 Preprocessing

The OCT volume is first segmented into 10 layers by 11 surfaces using the multiscale 3-D graph-search approach.\textsuperscript{22–24} To remove the image deformation caused by eye movement, the 3-D OCT...
volume is flattened based on the retinal pigment epithelium (surface 11). The inner retina is defined as the region between surfaces 1 and 6, and the outer retina refers to the region between surfaces 7 and 11. As the CMEs are mainly located in the inner retina, the inner retina is our initial volume of interest (VOI). So the precise segmentation of surfaces 1 and 6 is necessary. As shown in Fig. 3(b), surface 6 is well detected by the 3-D graph-search-based approach except the region around the CMEs, due to the coexistence of CMEs and MH. Surface 6 is then adjusted by replacing it with the lower boundary of its convex hull. Figure 3(c) shows the results after adjustment.

The OCT images acquired from different patients have different intensity profiles, which will affect the CMEs segmentation. In this paper, a SNR balancing is used on each B-scan to obtain uniform intensity profile. The noise level $N$ is taken as the mean intensity within a $20 \times 20$ window in the background. The signal level $S$ is taken as the mean intensity within a $20 \times 20$ window in the region between surfaces 1 and 6. The image data are SNR balanced using the equation $I_{\text{SNR}} = (I_O - N) / (S - N)$, where $I_O$ is the original intensity value and $I_{\text{SNR}}$ is the final intensity value.

Speckle noise is the main quality-degrading factor in OCT scans. The denoising method applied to OCT images should be efficient for speckle noise suppression while maintaining edge-like features. A 3-D curvature anisotropic diffusion filter is adopted in this paper. This filter performs anisotropic diffusion using the modified curvature diffusion equation (MCDE), which can efficiently enhance the contrast of edges. The MCDE equation is given as

$$f_t = |\nabla f| \| \nabla \cdot c(|\nabla f|) \| \nabla f,$$

where $f$ denotes the input image, $c$ is the conductance function. The conductance modified curvature term is $\nabla \cdot c(|\nabla f|) / |\nabla f|$. Figure 4(b) shows a B-scan image after the 3-D curvature anisotropic diffusion filtering, in which the speckle noise is effectively reduced.

### 2.4 Macular Hole Detection and Vessel Silhouettes Exclusion

Because the MH and vessel silhouettes have similar intensity value to the CMEs, it is necessary to isolate the MH and vessel silhouettes before extracting the CMEs volumetric feature. This procedure can reduce the volume of data to be analyzed and reduce the FPs.

The likelihood that a point belongs to an MH footprint can be calculated from the 2-D projection image in $X-Y$ plane. Here, a 2-D projection image is generated by averaging the OCT sub-volume between surfaces 1 and 6 in the $z$-direction, as shown in Fig. 5(a). Then a footprint of MH is defined by thresholding [threshold value = 33, in range (0, 255)]. The initial result is shown in Fig. 5(b). Because the footprint of MH usually locates around the center of the 2-D projection image, the falsely detected regions nearby the border, caused by signal attenuation, are discarded. The final result of the MH footprint is shown in Fig. 5(c).

With the detected MH footprint, an improved region-growing method is applied to delineate the MH in the 3-D OCT image. First, the seeds for region growing are determined based on the following criteria:

1. belonging to the A-scans indicated by the 2-D MH footprint,
2. between surfaces 1 and 6,
3. with intensity smaller than $m_{\text{MH}} + \text{std}_{\text{MH}}$, in which $m_{\text{MH}}$ and $\text{std}_{\text{MH}}$ are the statistical mean intensity and standard deviation of the MH region, respectively.

![Fig. 3 Illustration of layer segmentation and adjustment. (a) One slice from the original 3-D OCT image. (b) Layer segmentation results of surface 1 (red) and surface 6 (purple) by the graph-search method. (c) Layer segmentation after surface 6 adjustment.](https://biomedicaloptics.spiedigitallibrary.org/journals/Journal-of-Biomedical-Optics/22/7/76014-3)

![Fig. 4 SNR balancing results followed by a 3-D curvature anisotropic diffusion filtering. (a) Original B-scan and (b) denoised B-scan.](https://biomedicaloptics.spiedigitallibrary.org/journals/Journal-of-Biomedical-Optics/22/7/76014-3)
Second, with the detected seeds, the region-growing method becomes automatic. The neighbors of the seeds are searched and determined whether they belong to the object region according to intensity.

The CMEs segmentation results may occasionally include the low-intensity regions, such as vessel silhouettes, which are shown in Fig. 6. The arrows indicate vessel silhouettes formed by the vasculature of the retina. The green line in (b) indicates the location of the slice shown in (a) on the projection image of the retina. The vessel silhouettes should be detected and excluded before the CMEs segmentation. Here, the vessel silhouettes are effectively detected using a vessel detector. First, a 2-D projection image [Fig. 6(b)] is calculated by averaging the intensities of pixels in the outer retina (from surfaces 7 to 12) in the z-direction, as the vessel silhouettes have excellent contrast only in the outer retina. Then the vessel silhouettes are segmented from the projection image by a KNN classifier with 31 features. Finally, if the voxels in the inner retina regions have the same x, y locations with the vessel silhouettes, they are excluded from the VOI.

At last, the subvolume of the retina between surfaces 1 and 6 without the MH and the vessel silhouettes is defined as our real VOI.

2.5 AdaBoost Classifier-Based Coarse Cystoid Macular Edemas Segmentation

After preprocessing, 57 features are extracted for each voxel in the VOI, including textural and structural information of the image. Table 1 shows the list of features. Features 1 to 8 describe the local texture of the image. The first 6 features were defined in our previous research. The skewness and kurtosis are used to describe the statistical distribution characteristics of the intensity values in the VOI. Features 9 to 57 describe the local image structure. In order to speed up the feature extraction phase, the training images are sampled which yielded images of 256 × 240 × 32 voxels. Then, an AdaBoost classifier is applied for the coarse segmentation of CMEs in the VOI. AdaBoost is an iterative boosting algorithm constructing a strong classifier as a linear combination of weak classifiers. The algorithm takes the set of \((x_1, y_1), (x_2, y_2), \ldots, (x_n, y_n)\) as the input, where \(x_i (i = 1, 2, \ldots, n)\) is a \(m\)-dimensional feature, and \(y_i (i = 1, 2, \ldots, n)\) is the corresponding label, set as 1 or 0 for cyst or noncyst voxels. In this paper, \(m = 57\).

The coarsely detected CME regions are processed by the morphological dilation and erosion operations, to generate the object and background seeds for the following graph cut-based fine CMEs segmentation.

2.6 Constrained Graph Cut-Based Fine Cystoid Macular Edemas Segmentation

The graph cut algorithm has been widely used for the image segmentation in recent years. The energy function of traditional graph cut includes a regional term and boundary term. Obtaining seeds is an important step of the graph cut algorithm.
Table 1 List of the features used in the AdaBoost classifier.

<table>
<thead>
<tr>
<th>Feature number</th>
<th>Feature description</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Max–min normalized intensity</td>
</tr>
<tr>
<td>2 to 3</td>
<td>Block mean intensity and standard deviation (block size: 3×3×3)</td>
</tr>
<tr>
<td>4 to 6</td>
<td>Gradient in three directions</td>
</tr>
<tr>
<td>7 to 8</td>
<td>Skewness and kurtosis</td>
</tr>
<tr>
<td>9 to 13</td>
<td>First eigenvalues of the Hessian matrix at scales = 1, 3, 6, 9, 14</td>
</tr>
<tr>
<td>14 to 18</td>
<td>Second eigenvalues of the Hessian matrix at scales = 1, 3, 6, 9, 14</td>
</tr>
<tr>
<td>19 to 23</td>
<td>Third eigenvalues of the Hessian matrix at scales = 1, 3, 6, 9, 14</td>
</tr>
<tr>
<td>24 to 53</td>
<td>Gaussian filter banks of zero, first, and second orders with derivatives at scales = 2, 4, 8</td>
</tr>
<tr>
<td>54 to 57</td>
<td>Occurrence matrix contrast, correlation, energy, and homogeneity</td>
</tr>
</tbody>
</table>

With the given seeds generated in the coarse segmentation step, an automated shape-constrained graph cut method is applied for the final fine segmentation.\(^{41}\) In the graph cut algorithm, the overall problem can be formulated as an energy minimization problem.\(^{39}\) The energy function of the shape-constrained graph cut is defined as

\[
E(l) = \sum_{p \in P} [\alpha D_p(l_p) + \beta S_p(l_p)] + \sum_{p \in P, q \in N_p} \gamma B_{p,q}(l_p, l_q),
\]

where \(\alpha, \beta, \gamma\) denote the weights for the data term, shape term, and boundary term, respectively. They satisfy \(\alpha + \beta + \gamma = 1\). The data term is defined as a probability of the voxel to be the object or not. The boundary term is defined on the gradient of the intensity between the adjacent voxels. They are defined as follows:

\[
D_p(l_p) = \begin{cases} 
-\ln P(l_p|O), & \text{if } l_p \text{ is object label} \\
-\ln P(l_p|B), & \text{if } l_p \text{ is background label}
\end{cases}
\]

\(B_{p,q}(l_p, l_q) = \exp \left[-\frac{(l_p - l_q)^2}{2\sigma^2}\right] \frac{1}{d(p, q)} \delta(l_p, l_q),\)

and

\[
\delta(l_p, l_q) = \begin{cases} 
0, & \text{if } l_p \neq l_q \\
1, & \text{otherwise}
\end{cases}
\]

where \(l_p\) is the intensity of the voxel \(p\) and \(l_p\) is the label assigned to the voxel \(p\). \(P(l_p|O)\) and \(P(l_p|B)\) represent the probabilities that the voxel \(p\) belongs to CME and non-CME, respectively. They can be calculated from the histogram of the CME and non-CME voxels during the training phase. \(d(p, q)\) is the distance between voxel \(p\) and \(q\). \(\sigma\) is the standard deviation of the voxel intensity.

The shape term is defined as

\[
S_p(l_p) = 1 - \exp \left[-\frac{d(p, \text{CME})}{r}\right],
\]

where \(d(p, \text{CME})\) is the distance from voxel \(p\) to the coarse CMEs segmentation result in the AdaBoost classification. \(r\) is the radius of a circle that can just enclose the CMEs. The shape-constrained graph cut is implemented on the VOI and the CMEs region can be refined.

2.7 False Positive Rejection

The CMEs FPVs may be caused by the noise and the retinal region with low intensity value. The connected component detection based on morphological operations is used to reject the FPVs. If the area of the detected connected region is smaller than 6 pixels, it will be discarded as an FP.

3 Results

Leave-one-out cross-validation strategy was used for training and testing the AdaBoost classifier and the parameters used in the graph cut. To objectively evaluate the CME volume segmentation results, the true positive volume fraction (TPVF), FP volume fraction (FPVF), accuracy rate (ACC), and dice similarity coefficient (DSC) are used in the experiment.\(^{33,44}\) TPVF indicates the rate of correctly detected volume compared with the reference standard. FPVF denotes the fraction of incorrectly detected volume in the true negative volume. ACC indicates the detection accuracy. DSC is used for comparing the similarity between the automated segmentation results and the ground truth. The definitions are defined as follows:

\[
\text{TPVF} = \frac{|V_{TP}|}{|V_{GT}|},
\]

\[
\text{FPVF} = \frac{|V_{FP}|}{|V| - |V_{GT}|},
\]

\[
\text{ACC} = \frac{|V_{TP}| + |V_{TN}|}{|V|},
\]

\[
\text{DSC} = 2 \times \frac{|V_{TP}|}{|V_{TP}| + |V_{FP}| + |V_{GT}|}.
\]

Table 2 Mean ± standard deviation comparison of the CMEs segmentation results.

<table>
<thead>
<tr>
<th>Method</th>
<th>TPVF (%)</th>
<th>FPVF (%)</th>
<th>ACC (%)</th>
<th>DSC (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ref. 14</td>
<td>72.5 ± 17.6</td>
<td>8.3 ± 5.6</td>
<td>82.5 ± 3.2</td>
<td>71.2 ± 6.8</td>
</tr>
<tr>
<td>Ref. 16</td>
<td>90.5 ± 12.3</td>
<td>8.3 ± 5.9</td>
<td>91.9 ± 5.0</td>
<td>53.5 ± 11.0</td>
</tr>
<tr>
<td>AdaBoost-only</td>
<td>79.5 ± 8.7</td>
<td>0.80 ± 0.42</td>
<td>98.1 ± 0.6</td>
<td>79.1 ± 7.2</td>
</tr>
<tr>
<td>Proposed method</td>
<td>81.0 ± 7.8</td>
<td>0.80 ± 0.63</td>
<td>99.7 ± 0.1</td>
<td>80.9 ± 5.7</td>
</tr>
</tbody>
</table>
where $| \cdot |$ denotes volume, $V_{TP}$ denotes the true positive set, $V_{FP}$ denotes the FP set, $V_{TN}$ denotes the true negative set, and $V_{FN}$ denotes the false negative set. $V_{GT}$ denotes the ground truth, and $V$ denotes the VOI.

As shown in Table 2, the mean of TPVF, FPVF, ACC, and DSC for the proposed method was 81.0%, 0.80%, 99.7%, and 80.9%, respectively. The proposed method was compared with the coarse CMEs segmentation based only on the AdaBoost
algorithm, the 2-D thresholding segmentation method proposed in Ref. 14, and the lasso regularization-based regression method proposed in Ref. 16. Compared to the results by the AdaBoost algorithm, the DSC of the proposed method increased slightly (with a $p$-value of 0.061 in the paired $t$-test) and the corresponding ACC showed a significant improvement (with a $p$-value <0.0001). The results of the proposed 3-D method showed a more significant improvement compared to the results of the 2-D thresholding segmentation proposed in Ref. 14. Though the average TPVF of the regression method proposed in Ref. 16 is higher than the proposed method, its FPVF, ACC, and DSC are much poorer than the ones of the proposed method, which is caused by large FP regions.

Figure 7 shows five examples of CMEs segmentation results. The first column shows the original image, the second column shows the coarse segmentation results of the AdaBoost classifier, the third column shows the final fine segmentation results, the fourth column shows the segmentation results by 2-D thresholding segmentation method proposed in Ref. 14, the fifth column shows the segmentation results by lasso regularization-based regression method proposed in Ref. 16, and the last column shows the GT1. Note the improvements between columns 2 and 3. There are FPs caused by the presence of MH in columns 3 and 4.

Figure 8 shows the linear regression analysis results for the GT1 versus the GT2. The correlation $R^2$ between the proposed method and GT1, GT2, and the average of GT1 and GT2 were
0.9794, 0.9668, and 0.9812, respectively. The correlation coefficient between GT1 and GT2 was 0.9677, which indicated the two ground truths are very consistent.

Figure 9 shows the CMEs volume of 19 subjects by three methods. Bland and Altman plots were evaluated to propose the agreement between two different methods. The Bland–Altman plots for the proposed method (auto) versus the GT1 and the GT2 are shown in Fig. 10. The points were mostly located in the 95% limits of agreement (±1.96 standard deviation of the difference), which indicated good agreement between the proposed method and GT1 and GT2.

4 Discussion and Conclusions
Automated segmentation of CMEs in 3-D OCT images is a challenging task due to the variation of CMEs’ shape and size and the textural similarity between the foreground and background. The experiment results showed the good performance of the proposed method. The FPVF of the proposed method was low, which may be due to several facts: (1) compared with the VOI region between surfaces 1 and 6, the volume of the CMEs is very small; (2) the size for FP exclusion in the postprocessing is very effective. The CMEs can be effectively segmented by the proposed method. With coexistence of MH, both the 2-D thresholding method proposed in Ref. 14 and the lasso regularization-based regression method proposed in Ref. 16 fail to exclude the MH effectively, because the intensity value is similar between MH and CMEs and neither of these two methods uses the spatial and shape information about the MH.

Our contributions are summarized as follows:

- This is a method for the automated 3-D CMEs volume segmentation in the retina with coexistence of MH.
- The proposed method can effectively detect CMEs with unpredictable locations and ambiguous boundaries (as shown in Fig. 11).
- An improved efficient region-growing method is proposed to delineate the MH in 3-D OCT image.
- The seeds used in the shape-constrained graph cut algorithm are generated automatically and the proposed method is a fully automated framework.

However, there are still some limitations in our method.
(1) The proposed method can accurately detect a big cyst, but it is not sensitive to very small cysts. This will be improved in the near future.
(2) The final CMEs detection results largely rely on the preprocessed results. If the MH exclusion result is incorrect, the final segmentation results may include falsely detected regions in MH. A better algorithm for MH detection is needed.
(3) A larger dataset will be explored in the near future for further validation of the proposed method.

In conclusion, an effective approach is proposed to segment the CMEs in the 3-D OCT images with coexistence of MH. This may provide a clinically useful tool to help ophthalmologists with disease diagnosis.

Disclosures
Li Zhang has a financial interest in Oracle Inc., which, however, did not support this work. No conflicts of interest, financial or otherwise, are declared by the other authors.

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References


Welfang Zhu is an associate professor at Soochow University, Suzhou, China. She received her PhD in electrical engineering from Soochow University, China, in 2013. She has coauthored over 15 papers in internationally recognized journals and conferences. Her current research is focused on medical image processing and analysis.

Xinjian Chen is a distinguished professor at Soochow University, Suzhou, China. He received a PhD from the Chinese Academy of Sciences in 2006. He conducted postdoctoral research in the University of Pennsylvania, National Institute of Health, and University of Iowa, USA from 2008 to 2012. He has published over 80 top international journals and conference papers. He has also been awarded six patents.

Biographies for the other authors are not available.