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9-1-2013

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Accepted version. *JAMA Ophthalmology*, Vol. 131, No. 9 (September 2013): 1207-1215. DOI. ©  
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### **Abstract**

#### *Objective*

To characterize outer retinal structure in Best Vitelliform Macular Dystrophy (BVMD), using spectral domain optical coherence tomography (SD-OCT) and adaptive optics scanning light ophthalmoscopy (AOSLO).

#### *Methods*

Four symptomatic members of a family with BVMD with known *BEST1* gene mutation were recruited. Thickness of two outer retinal layers corresponding to photoreceptor inner and outer segments were measured using SD-OCT. Photoreceptor mosaic AOSLO images within and around visible lesions were obtained, and cone density was assessed in two subjects.

#### *Results*

Each subject was at a different stage of BVMD, with photoreceptor disruption evident by AOSLO at all stages. When comparing SD-OCT and AOSLO images from the same location, AOSLO images allowed for direct assessment of photoreceptor structure. A variable degree of retained photoreceptors was seen within all lesions. The photoreceptor mosaic immediately adjacent to visible lesions

appeared contiguous and was of normal density. Fine hyperreflective structures were visualized by AOSLO, and their anatomical orientation and size are consistent with Henle fibers.

### *Conclusions*

AOSLO findings indicate substantial photoreceptor structure persists within active lesions, accounting for good visual acuity in these patients. Despite previous reports of diffuse photoreceptor outer segment abnormalities in BVMD, our data reveal normal photoreceptor structure in areas adjacent to clinical lesions.

### *Clinical Relevance*

This study demonstrates the utility of AOSLO for understanding the spectrum of cellular changes that occur in inherited degenerations such as BVMD. Photoreceptors are often significantly affected at various stages of inherited degenerations, and these changes may not be readily apparent with current clinical imaging instrumentation.

Best Vitelliform Macular Dystrophy (BVMD), also known as Vitelliform Macular Dystrophy type 2 or Best's Disease (OMIM # 607854; BEST1) is an autosomal dominant form of macular degeneration of variable penetrance characterized by varying accumulation of yellowish vitelliform material in the macula.<sup>1,2</sup> Affected individuals also show a reduction in the electrooculogram (EOG) light peak but a normal full-field electroretinogram (ERG).<sup>1,3</sup> Mutations in the *BEST1* gene on chromosome 11q13 encoding bestrophin-1 cause BVMD.<sup>4-6</sup> Bestrophin-1 is an integral membrane protein that has been localized to the basolateral membrane of the retinal pigment epithelium (RPE)<sup>7</sup> and is thought to be a Ca<sup>+2</sup> sensitive Cl<sup>-</sup> channel protein or influences the regulation of Ca<sup>+2</sup> channels.<sup>8</sup>

The clinical appearance of BVMD varies by the stage of the disease.<sup>2</sup> Initially, retinal fundi may appear normal (previtelliform). Characteristically, there is development of macular fluid- and debris-filled retinal detachments forming a yellow yolk-like or vitelliform lesion or lesions. With time, the vitelliform material may become more heterogenous with various layers (pseudohypopyon) and may appear to dissolve, leaving isolated clumps of material at the edges of the

lesion (vitelliruptive). Eventually localized atrophy and fibrosis develops in the location of the vitelliform lesion.<sup>2</sup> Despite presence of vitelliform lesion(s), vision is usually good in earlier stages of the disease, visual acuity of 20/40 or better vision being reported in 76% of individuals less than 40 years of age.<sup>9</sup> It has been shown that normal acuity can be maintained in individuals having substantial photoreceptor degeneration.<sup>10,11</sup> Thus, the good visual acuity in patients with BVMD does not necessarily inform about the degree of photoreceptor degeneration.

Histopathologic findings from BVMD donor eyes are limited but demonstrate abnormal accumulation of lipofuscin granules in the RPE<sup>12-15</sup> and photoreceptor degeneration over areas of intact RPE.<sup>16,17</sup> Recently a knock-in mouse model of BVMD showed increased accumulation of lipofuscin in the RPE and deposition of subretinal debris composed of unphagocytosed photoreceptor outer segments and lipofuscin granules.<sup>18</sup> It is hypothesized that impairment (rather than loss) of RPE to fully degrade phagocytosed outer segments leads to photoreceptor degeneration in BVMD, either alteration of the ionic milieu of the subretinal space due to bestrophin mistargeting or loss of cell-to-cell contact.<sup>13,16</sup>

Optical coherence tomography (OCT) imaging techniques allow for non-invasive assessment of retinal structure, and numerous studies have utilized this imaging approach to assess outer retinal structure in BVMD.<sup>19-23</sup> OCT imaging has shown that the characteristic vitelliform lesions of BVMD are the result of accumulation of material in the subretinal space above the RPE and below the outer segments of the photoreceptors.<sup>20,21,24,25</sup> Also, despite bestrophin-1 being localized to the RPE, OCT has shown significant changes to outer retinal structure are evident at various stages of the disease, and it has been suggested that thickening of the reflective layer corresponding to the photoreceptors may be one of the earliest anatomical changes visible by OCT with BVMD.<sup>20,21,26</sup> However, examples exist where the resolution of existing OCT technology is not sensitive enough to detect pronounced photoreceptor disruption.<sup>27-29</sup> Thus, despite the OCT findings in BVMD, the nature of photoreceptor structure in BVMD remains unclear.

Adaptive optics (AO) imaging systems enable cellular-resolution imaging of the human retina, allowing for direct visualization of cone and rod photoreceptor mosaic.<sup>30,31</sup> To better understand photoreceptor structure across the spectrum of BVMD, we used spectral-domain OCT (SD-OCT) and adaptive optics scanning light ophthalmoscopy (AOSLO) to assess retinal structure in four members of the same family who are at various stages of BVMD and have a known *BEST1* mutation.

## Methods

Four members of a family with a previously identified mutation, p.Arg218Cys (c.652C->T) (University of California Ophthalmic Molecular Diagnostic Lab, La Jolla, CA), in the *BEST1* gene reported to be causative mutation in BVMD<sup>32</sup> and with clinical findings consistent with BVMD participated (Table 1, eFigure 1). The p.Arg218Cys mutation is predicted to affect the charge of the bestrophin protein, altering its function. Visual acuity was assessed, and a comprehensive eye exam including fundus photography was performed for all four subjects. Each patient was dilated using one drop of phenylephrine (2.5%). Mircoperimerty was performed. Then accommodation was suspended using one drop of tropicamide (1%) for subsequent high-resolution imaging. Axial length was measured using an IOL Master (Carl Zeiss Meditec, Dublin, CA). This prospective study was conducted in accordance with the tenets of the Declaration of Helsinki and with institutional review board approval.

Subject	Age, y	Gender	Mutation in <i>Best1</i> gene	Type of Lesion*	Visual Acuity** (OD, OS)	Axial Length (OD, OS, mm)	Eye imaged with AOSLO
IV-3	16	M	Arg218Cys (c.652C->T)	Early vitelliform	20/20, 20/20	22.57/22.14	OS
IV-2	18	F	Arg218Cys (c.652C->T)	Vitelliform with early vitelliruptive	20/20, 20/20	22.72/22.62	OS
III-5	50	F	Arg218Cys (c.652C->T)	Late vitelliruptive	20/30, 20/100	22.11/21.98	OD
III-4	59	F	Arg218Cys (c.652C->T)	Atrophic	20/200, 20/50	23.14/23.69	OS

## **Table 1** Patient Demographics

\*Same lesion type in both eyes,

\*\*Snellen

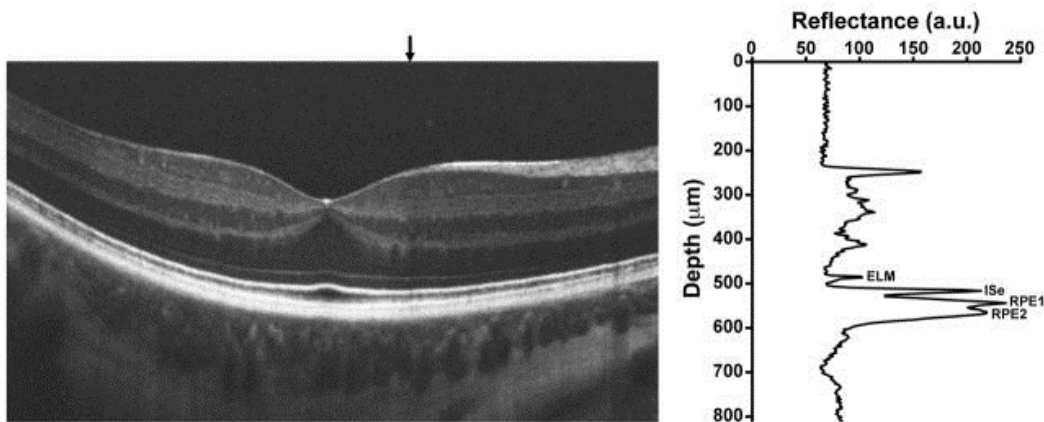
y-years, OD –right, OS –left, M-male, F- female, mm- millimeters

Macular microperimetry was performed using the Spectral OCT/SLO MP system (OPKO instrumentation) after a brief training to allow for familiarization of the test. A Polar 3 standardized grid composed of 28 points arranged in 3 concentric circles (2.3°, 6.6°, and 11° in diameter from fovea, 4 points in innermost circle, 12 in middle and outer circles) was performed using a Goldman III stimulus, a 200ms duration and a test strategy 4-2. Results were compared to previously published normative data<sup>33</sup>.

Volumetric images of the macula were obtained using Cirrus HD-OCT (Carl Zeiss Meditec, Dublin, CA). Volumes were nominally 6 mm × 6 mm and consisted of 128 B-scans (512 A-scans/B-scan). Retinal thickness was assessed using the built-in macular analysis software (software version 5.0), which is automatically generated by calculating the difference between the inner limiting membrane (ILM) and RPE boundaries. The software's "fovea finder" algorithm was used to determine the location of the fovea on the line scanning ophthalmoscope (LSO) image. Additional high-density line scans (1000 A-scans/B-scan, 100 repeated B scans) were acquired through the foveal center in the study eye of each participant using the Bioptigen SD-OCT (Bioptigen, Inc., Research Triangle Park, NC). Line scans were registered and averaged to reduce speckle noise in the image using previously described techniques,<sup>34</sup> and were acquired in both the horizontal and vertical direction. All scans shown in the figures are from the Bioptigen device. Numerous naming conventions exist in the literature for the outer hyperreflective layers in SD-OCT scans, so it is important to define the one used here. Shown in Figure 1 is a horizontal line scan from a normal control, and a corresponding longitudinal reflectivity profile (LRP), showing the identity of the bands analyzed.<sup>35,36</sup> The innermost band corresponds to the external limiting membrane (ELM), the second band corresponds to the inner segment ellipsoid (ISe),<sup>37</sup> the third band corresponds to the outer segment/RPE interface (RPE1), and the fourth band corresponds to the RPE (RPE2). The peak-to-peak distance between the ELM and ISe is taken as the length of the inner segments (IS), while the peak-to-peak distance



between the ISe and RPE1 is taken as the length of the outer segments (OS). While these may not correspond precisely to the absolute IS or OS length, we utilized these same definitions in an extensive previously published normative dataset.<sup>36</sup> We examined the IS and OS length across the horizontal line scan from each subject, sampling the scan at 0.2mm intervals. We excluded the central BVMD-related lesion from further analysis, similar to a previous report.<sup>21</sup> Images of the photoreceptor mosaic were acquired using a previously described AOSLO.<sup>30,38</sup> Images were obtained using an Inphenix 775nm superluminescent diode 12nm FWHM bandwidth with either a, 1.0 or 1.75 degree square field of view. The fovea and surrounding areas affected by pathology were imaged in each patient. Parafoveal images (~0.65 degrees from fixation) were acquired by instructing the patient to fixate on the corners or edges of the raster, while more eccentric images were acquired using an internal fixation target. Intraframe distortions within the AOSLO retinal images were corrected as previously described.<sup>30,39</sup> Registration of frames within a given image sequence was performed using a "strip" registration method, in which the images were registered by dividing the image of interest into strips, aligning each strip to the location in the reference frame that maximizes the normalized cross correlation between them.<sup>39</sup> Once all the frames were registered, the 50 frames with the highest normalized cross correlation to the reference frame were averaged, in order to generate a final image with an increased signal to noise ratio (SNR).



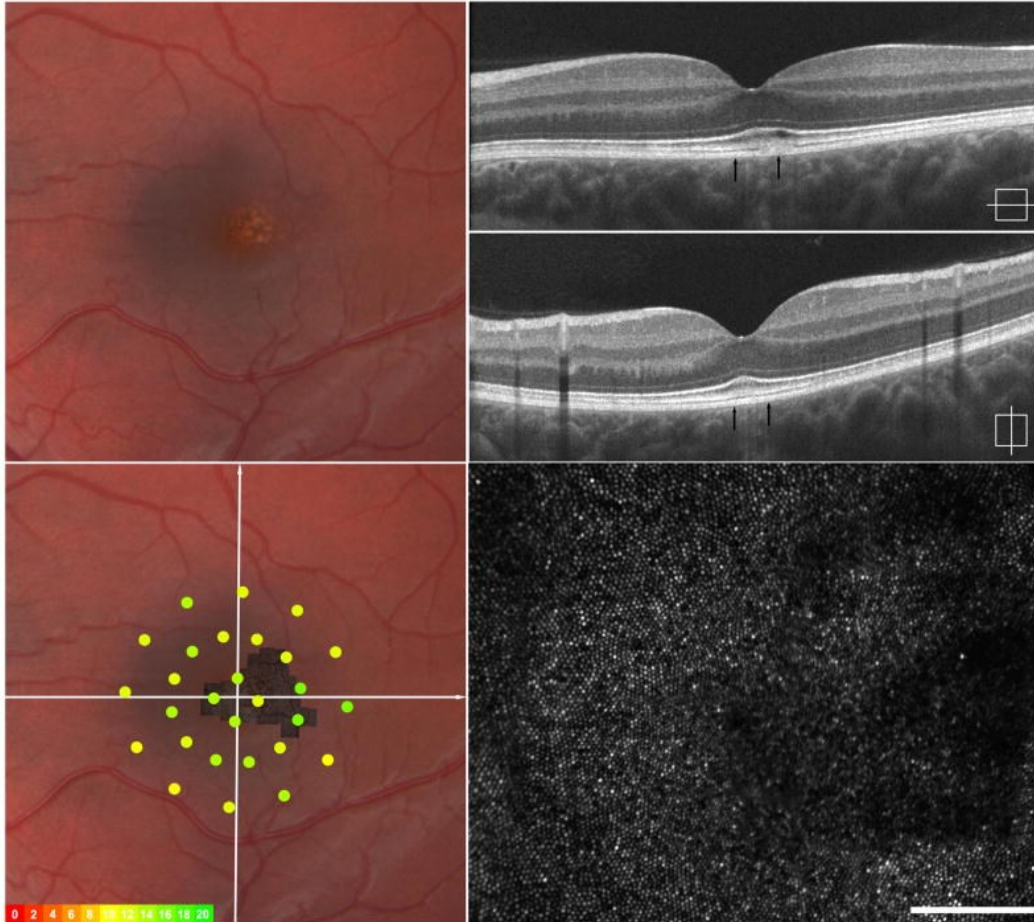
**Figure 1** Assignment of outer retinal bands on SD-OCT. Shown is a horizontal line scan through the fovea of a normal subject. The graph on the right is a longitudinal reflectivity profile (LRP) acquired at the location of the vertical black arrow above the

SD-OCT scan. ELM=external limiting membrane, ISe=inner segment ellipsoid, RPE1=outer segment/RPE interface, RPE2=RPE.

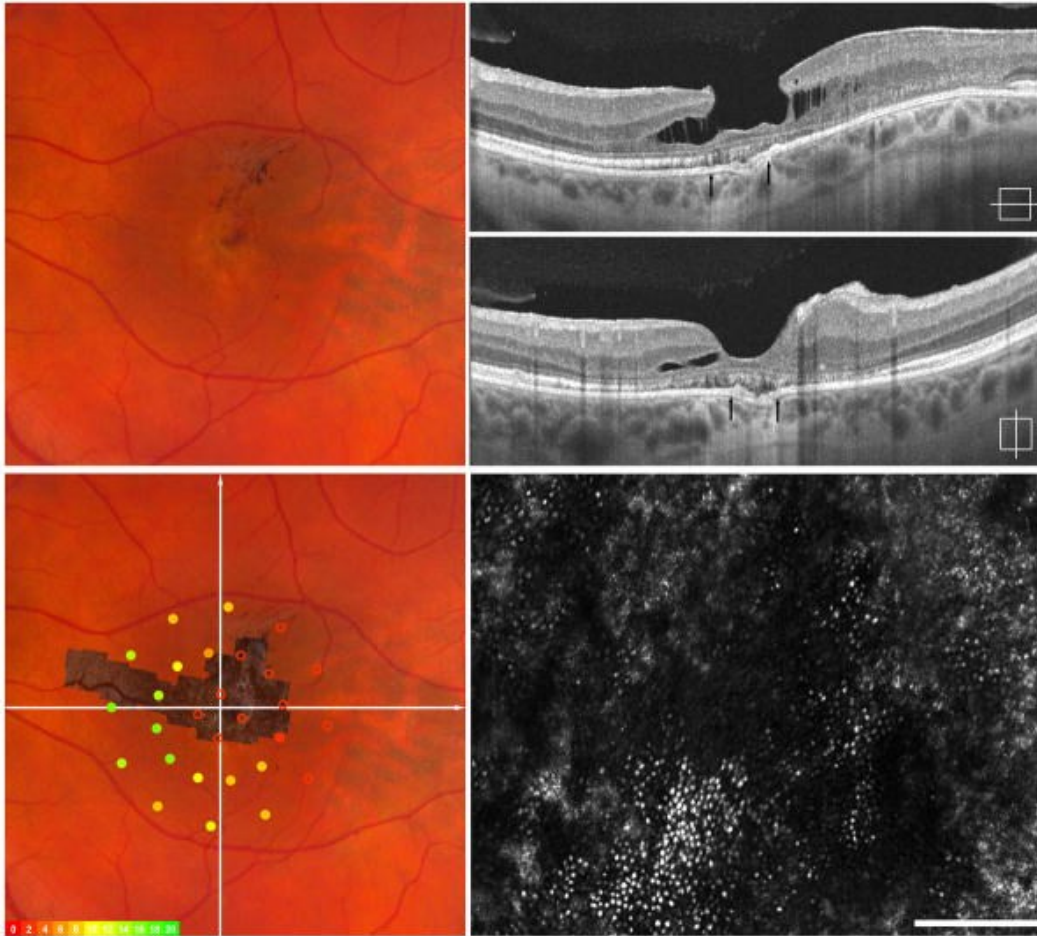
These registered and averaged AOSLO images were then montaged using Adobe Photoshop (Adobe Systems, Inc., CA). The montage was aligned to the color fundus images and the LSO image from the Cirrus HD-CT, which was exported with the location of the foveal pit marked. Scaling of the images was done based on the expected scale of each image and alignment was done manually utilizing blood vessel patterns. Cone density was assessed using 55 $\mu$ m  $\times$  55 $\mu$ m sampling areas adjacent to the visible lesion in two subjects and near the fovea within the active lesion in all four subjects using a previously described semi-automated algorithm.<sup>40</sup> The distance between the sampled area and the foveal pit was measured, enabling comparison of density values to previously published normative values.

## Results

Four affected subjects from a family with BVMD with known p.Arg218Cys mutation in *BEST1* gene participated. (eFigure 1) All family members were found to be at different stages of the disease, summarized in Table 1. The SD-OCT and AOSLO imaging findings were unique to each stage (Figures 2–5).. Macular microperimetry performed within a 6 degree radius of the fovea revealed areas of subnormal individual point sensitivities in regions corresponding to clinical retinal lesions (lower left, Figures 2–5) in all but subject IV-3 with early vitelliform findings. In patient IV-2, decreased point sensitivities were seen both in regions surrounding the vitelliform lesion and overlying the lesion itself.

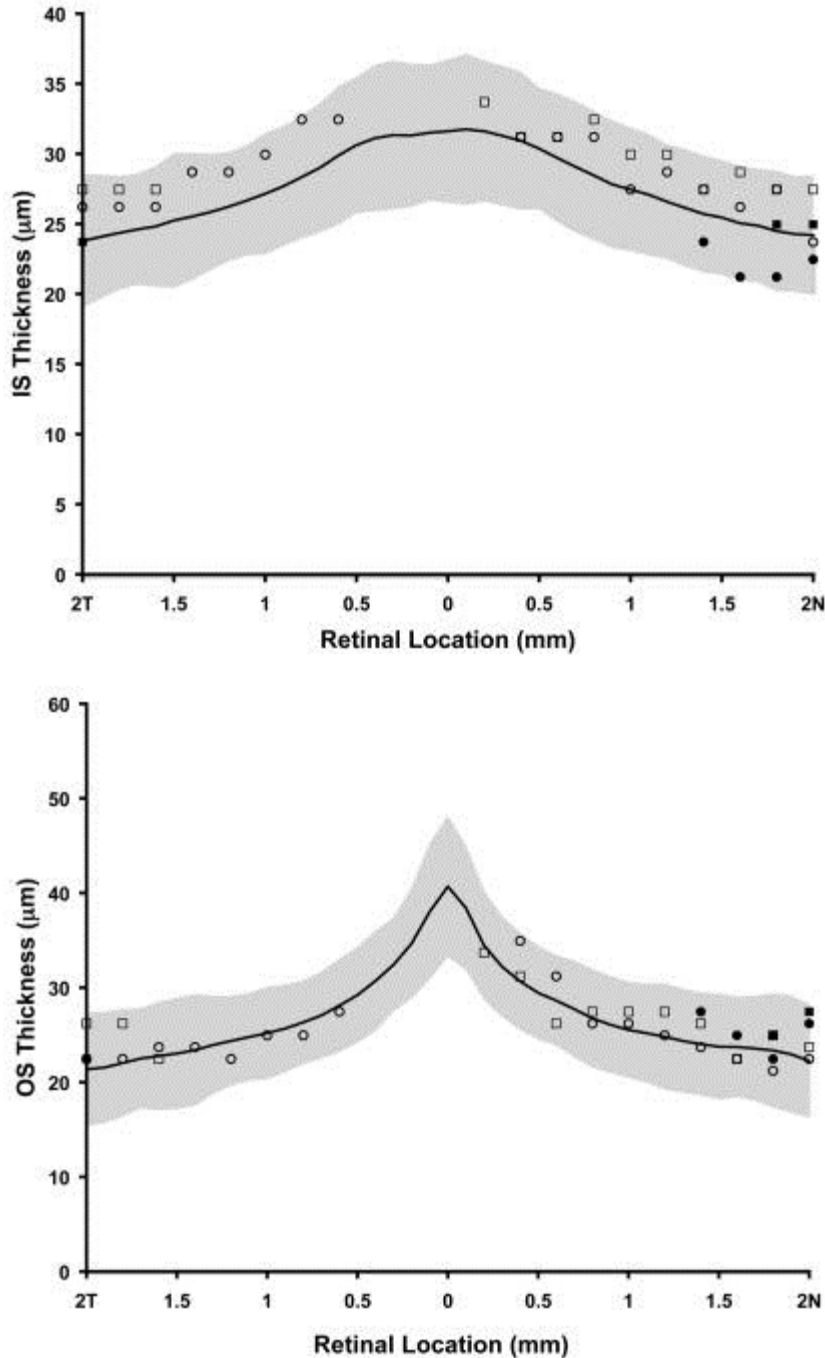


**Figure 2** Imaging of IV-3, left eye – Early vitelliform findings. Fundus exam revealed a focal area of granularity just temporal to the fovea (upper left). SD-OCT horizontal and vertical scans show normal retinal lamination but focal increased hyper-reflectivity in the area of granularity seen clinically (upper right). Macular microperimetry showed normal point sensitivities in the central 12 degrees (overlay, left lower) Adaptive optics imaging of this location (montage registered lower left, area imaged indicated by arrows on SD-OCT upper right) shows focal photoreceptor mosaic disruption around the area of hyper-reflectivity on OCT with photoreceptor mosaic surrounding this area appearing normal (lower right).



**Figure 5** Imaging of III-4, left eye - Atrophy and Fibrosis. Fundus exam shows central hypopigmentation with focal pigment mottling and trace epiretinal membrane (ERM) (upper left). SD-OCT horizontal and vertical scans show a lamellar hole, trace ERM and loss of the hyperreflective ISe band (upper right). Macular microperimetry revealed subnormal point sensitivities in areas central and temporal to fovea when fibrosis and atrophy are present clinically (overlay, left lower). Adaptive optics imaging of central fovea (montage registered lower left, area imaged indicated by arrows on SD-OCT upper right) reveal patchy areas of retained photoreceptors between areas of significant photoreceptor loss (lower right).

Measurement of IS and OS retinal thickness was performed using the SD-OCT horizontal line scan in all four subjects. Shown in Figure 6 is the IS and OS thickness profile in areas immediately adjacent to clinical visible lesions for all four subjects compared to data from a previously published normative group.<sup>41</sup> Thickness values were not calculated over the visible lesion. All 4 subjects were found to have IS and OS thickness values within 2 standard deviations of normative data.



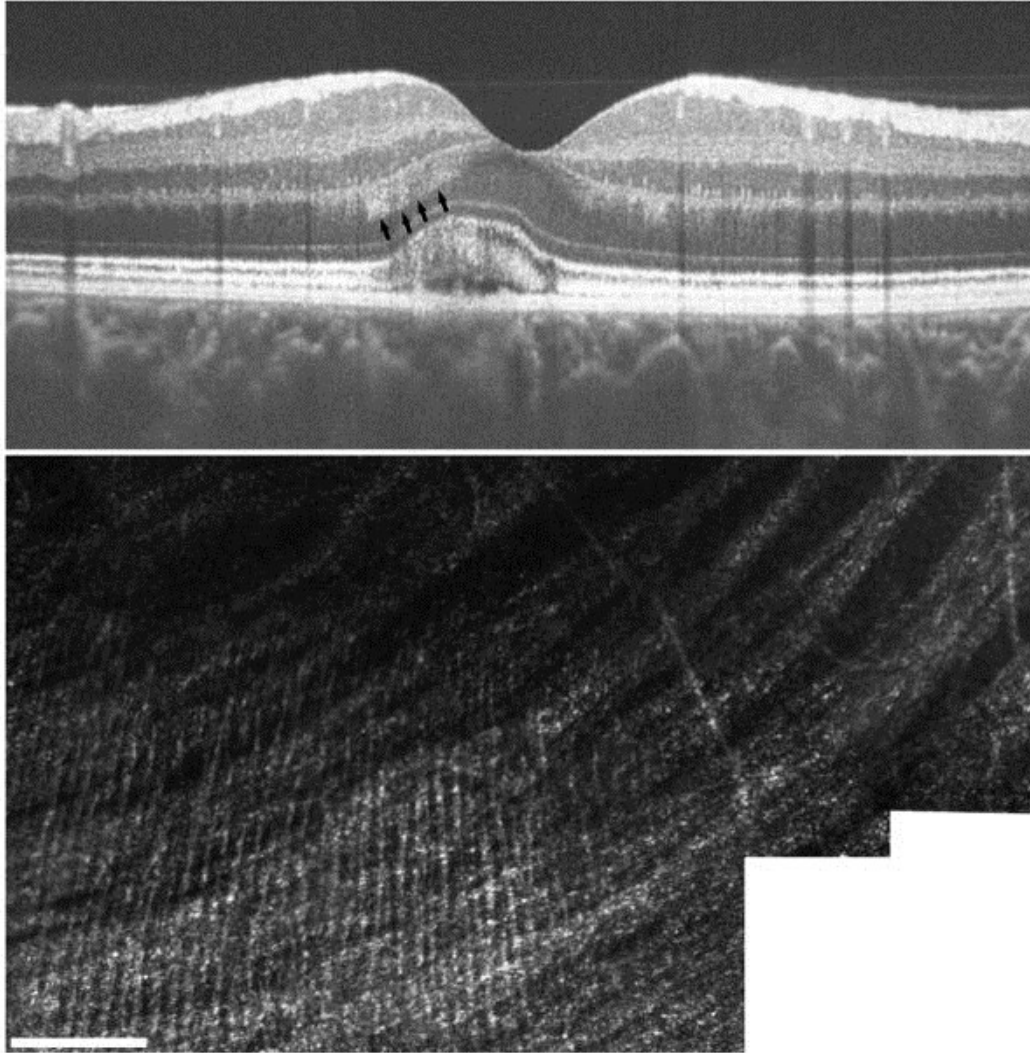
**Figure 6** Assessment of photoreceptor layer thickness in BVMD. Top, plot of inner segments (IS) length as a function of retinal location along the horizontal meridian. Bottom, outer segment (OS) length as a function of retinal location along the horizontal meridian. Black line – normative data from 93 people, average age 25.7 years with standard deviation (stdev) of 8.2 years. Shaded grey area is + 2 stdev. Open squares – subject IV-2, Open circles – subject IV-3, Filled squares – subject III-



5, Filled circles – subject III-4. Thickness values were not calculated over visible lesion(s).

We sought to further assess photoreceptor structure in the retinal area adjacent to the BVMD lesions. In two patients having lesions with a clear boundary, we were able to obtain AOSLO montages that were large enough to encompass the entire lesion (Figure 2, eFigure 2). We assessed cone density just nasal to the lesion boundary in both IV-3 and IV-2, and determined that the areas sampled were 1 degree from the foveal center. The cone mosaic appeared contiguous and cone density was 55,900 cones/mm<sup>2</sup> in IV-3 and 43,700 cones/mm<sup>2</sup> in IV-2. Both values are within the normal range for this retinal eccentricity.<sup>41</sup>

In the SD-OCT scans of one of the subjects (IV-2) we noticed significant hyperreflective material in the outer nuclear layer (ONL). This has been previously reported in BVMD,<sup>42</sup> and is attributed to the physical deformation of the Henle fiber layer by the underlying vitelliform lesion. Inspection of the SD-OCT volume revealed the strongest signal in the inferior retina, just nasal to the fovea. AOSLO images from this same location focused in the inner retina revealed thin hyperreflective structures running perpendicular to the nerve fiber layer (Figure 7). The anatomical location and orientation is consistent with that of Henle fibers, and the diameter of these structures (average =  $2.76 \pm 0.32 \mu\text{m}$ ) is consistent with previous histology reports.<sup>43</sup>

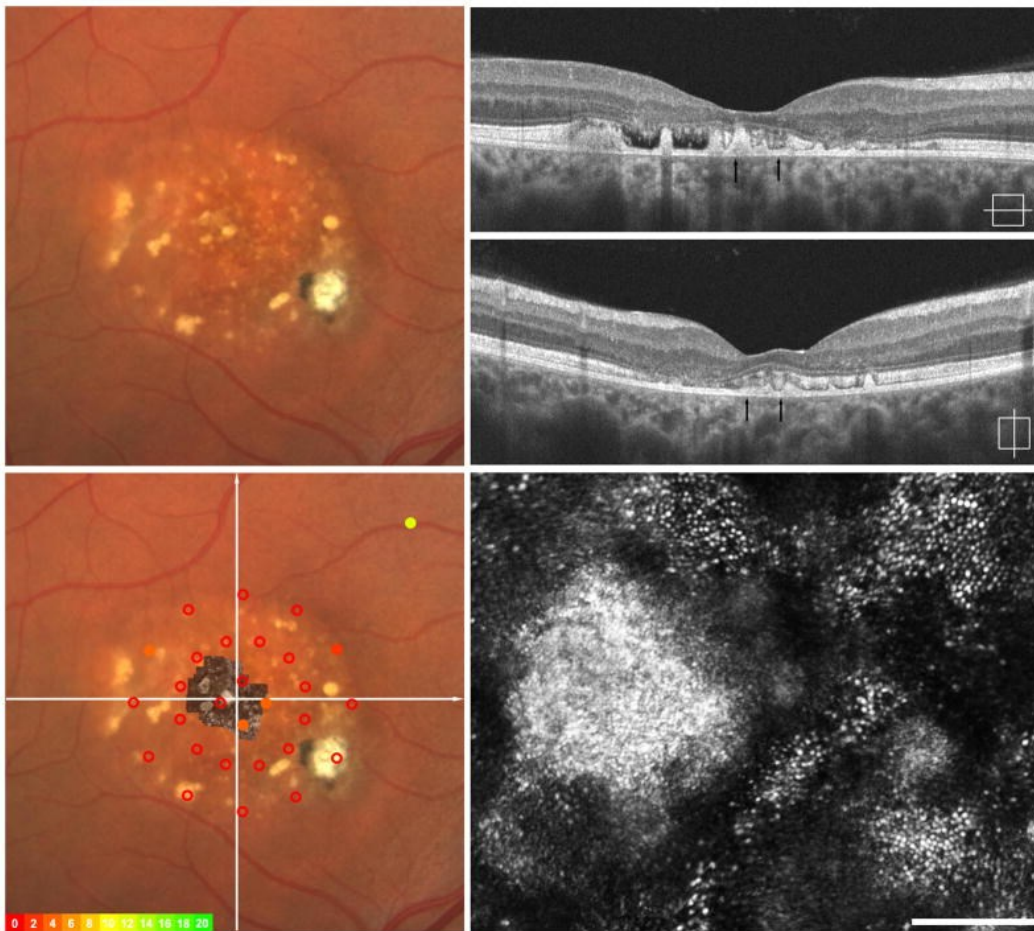


**Figure 7** Visualization of Henle fiber layer in BVMD. Presence of the vitelliform lesion has altered the angle of retina structure, allowing for visualization of Henle fiber layer on SD-OCT (arrows, top). AOSLO imaging at this same location focused at the level of the inner retina revealed thin hyperreflective structures running perpendicular to nerve fiber bundles, consistent with known anatomy of Henle fibers (lower).

## Comment

In our study we used SD-OCT and AOSLO to assess outer retinal structure in four members of a single family harboring a previously reported *BEST1* mutation (p.Arg218Cys). The phenotypes ranged from early vitelliform changes to a central atrophic area. Disruption of the cone mosaic was evident in the AOSLO images at all stages of BVMD presented here, including the patient with the earliest stage of

vitelliform clinical findings (Figure 2), suggesting this is an early finding in patients with BVMD. The degree of this photoreceptor disruption varied by stage of disease, and was often patchy with areas of significant photoreceptor disruption surrounded by areas of a contiguous photoreceptor mosaic, even in the patient with advanced atrophy and fibrosis. (Figure 5) It is important to note that disruption of visualization of cone structure on AOSLO does not necessarily mean the cone cell has been lost. When comparing SD-OCT and AOSLO images from the same location, the AOSLO images allowed for better understanding of the degree of retained photoreceptor structure at that location. This is illustrated in the patient with late vitellirruptive changes (Figure 4). SD-OCT of this individual shows significant disruption of the hyperreflective ISe band in the areas of subretinal nodules, but the AOSLO images reveal islands of contiguous cone mosaic adjacent to areas of significant disruption.





**Figure 4** Imaging of III-5, right eye – Vitelliruptive. Fundus exam reveals an ovoid area of hypopigmentation containing several fibrotic nodules (upper left). SD-OCT horizontal and vertical scans show outer retinal atrophy and several focal deposits of debris in the subretinal space, some separated by trace subretinal fluid. Patchy disruption of the hyperreflective ISe band is evident in some areas (upper right). Macular microperimetry revealed subnormal point sensitivities in all areas of central 6 degrees (overlay, left lower). Adaptive optics imaging of central fovea (montage registered lower left, area imaged indicated by arrows on SD-OCT upper right) reveals significant photoreceptor mosaic disruption overlying these nodules, but relative preservation of the photoreceptor mosaic surrounding these lesions (lower right).

Previous studies have suggested that loss of photoreceptors in BVMD could be widespread and not necessarily confined to the clinically apparent lesions, and support for this comes from the fact that bestrophin, the RPE membrane protein encoded by the *BEST1* gene, is found throughout the retina in individuals unaffected by BVMD.<sup>17</sup> Kay *et al.* recently showed increased photoreceptor thickness on SD-OCT in patients with BVMD when compared to normal controls within the macular region.<sup>21</sup> Based on their findings, they conclude that the primary anatomical impact is at the photoreceptor level. Certainly, our finding that the photoreceptor mosaic is disrupted in the earliest stage of clinical vitelliform findings would be consistent with this proposed etiology, but our finding of normal IS and OS thickness and normal cone density in retinal areas adjacent to visible lesions argues against a diffuse structural deficit in BVMD. One possible explanation is that the previous study did not correct the lateral scale of their SD-OCT scans for individual differences in axial length, meaning that different extents of retina contributed to the analysis in each retina. Moreover, since the previous analysis averaged the thickness measurements across the scan, it is unclear if the retina was indeed uniformly affected or if a small retinal area was severely abnormal.<sup>21</sup> Nevertheless, while our findings do not support diffuse disruption of the cone mosaic outside the lesion, it is possible that these cells may not be functioning normally.

Interestingly, macular microperimetry revealed areas of subnormal point sensitivities in areas surrounding the vitelliform lesion in subject IV-2. Both SD-OCT and AOSLO showed normal outer retinal anatomy within these regions. These reduced point sensitivities may be result of eye movements reducing the specificity of registration to the fundus. However, it may also be possible that functional loss of

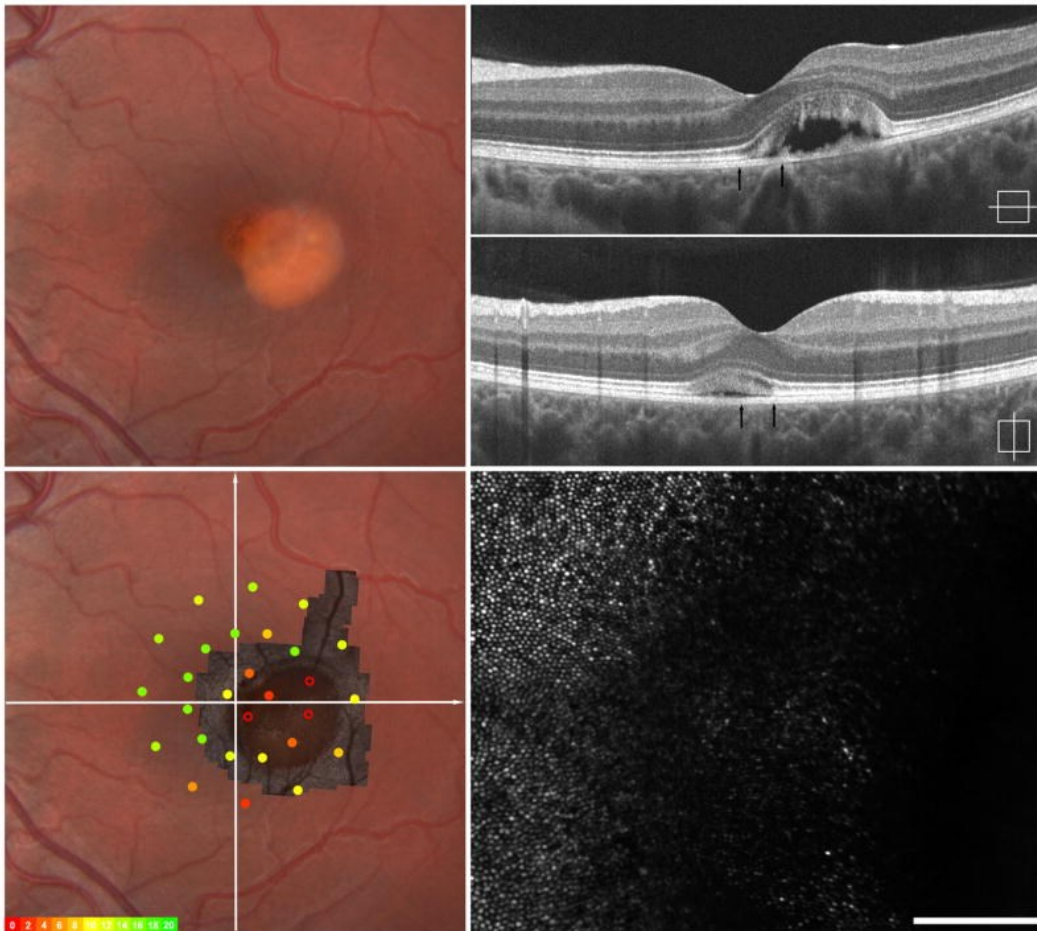
vision precedes anatomical outer retinal structural loss. High resolution microperimetric assessment using adaptive optics technology has been described.<sup>44,45</sup> To better understand and correlate functional vision to photoreceptor mosaic structure pathology, future studies would benefit from AOSLO based microperimetry allowing for functional assessment at resolutions on par to those used to assess retinal structure.

It is becoming appreciated that outer retinal pathology can affect the appearance of the overlying retina on SD-OCT. For example, presence of a vitelliform lesion, large drusen, or pigment epithelial detachment alters the orientation of the fibers of Henle as they traverse the lesion, altering their reflectivity on SD-OCT.<sup>46,47</sup> We also observed this effect in one of our subjects (IV-2), however we also observed the presence of fine hyperreflective structures running perpendicular to the nerve fiber bundles in the AOSLO images at the same retinal location (Figure 7). Their anatomical location, orientation, and size are consistent with that of Henle fibers. As seen with SD-OCT, this demonstrates that when imaged with AOSLO, outer retinal disruptions can alter the appearance of the inner retina, and this should be taken into consideration when analyzing such images.

A potential limitation of the current study is that all four subjects have the same genetic mutation in the *BEST1* gene. While our data reveal a spectrum of clinical and subclinical findings associated with this particular mutation, it is not possible to extend our findings on the integrity of the cone mosaic to other mutations. Future investigations should include high-resolution imaging of other individuals with different mutations in *BEST1* gene to investigate possible genotype-dependent differences in photoreceptor structure.

In summary, we provide evidence from cellular imaging with AOSLO that photoreceptor structure can be retained within active BVMD lesions, even in apparently atrophic lesions. This photoreceptor structure is capable of supporting rather good visual acuity, as visual acuity in the eyes imaged here ranged from 20/20 to 20/50. In addition, our SD-OCT and AOSLO data show normal photoreceptor structure in retinal areas outside the clinically visible lesion, in contrast to previous reports<sup>21</sup>, but consistent with previous findings with AOSLO.<sup>48</sup> This may represent a specific feature of the mutation studied here, or be due to different imaging and measurement procedures.

Regardless, our study highlights the utility of AOSLO imaging in directly delineating the degree of retained photoreceptor structure in diseases like BVMD. In particular, combining information from SD-OCT with that from AOSLO gives a complementary view of outer retinal structure and provides a more sensitive approach for measuring photoreceptor structure than either alone.

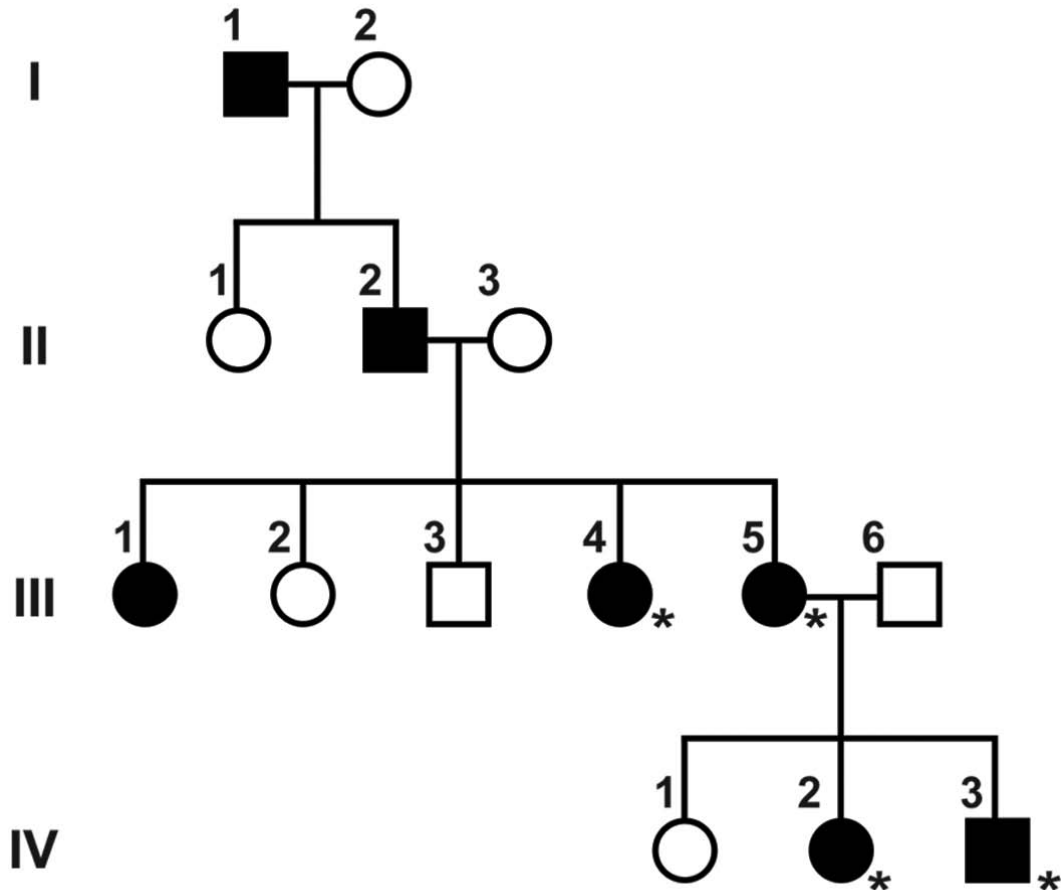


**Figure 3** Imaging of IV-2, left eye – Vitelliform lesion with early vitellirruptive changes. Fundus exam reveals single heterogeneous vitelliform lesion centered just temporal to fovea (upper left). SD-OCT horizontal and vertical scans show the vitelliform lesion contains fluid and debris within the subretinal space. There is patchy disruption of the hyperreflective ISe band over the lesion (upper right). Macular microperimetry revealed subnormal point sensitivities in areas overlying the vitelliform lesion and immediately surrounding it. (overlay, left lower). Adaptive optics imaging of the vitelliform lesion and area immediately surrounding this (montage registered lower left, area imaged indicated by arrows on SD-OCT upper right) reveals disrupted

photoreceptor mosaic over the lesion with normal mosaic seen immediately adjacent to the lesion.

## Supplementary Material

### Supp Figure 1



**eFigure 1:** Pedigree of family with Best Vitelliform Macular Dystrophy caused by Arg218Cys (c.652C>T) mutation in *BEST1* gene. Individuals marked with "\*" were imaged in this study. (will be moved from here...)

## Acknowledgments

This publication was supported by the National Center for Research Resources and the National Center for Advancing Translational Sciences, National Institutes of Health, through Grant Number 8UL1TR000055. Its contents are solely the responsibility of

the authors and do not necessarily represent the official views of the NIH. Additional support from the Clinical and Translational Science Institute and the Biotechnology Innovation Center, Medical College of WI, CTSA Grant UL1 RR 031973, the Thomas M. Aaberg Sr. Retina Research Fund, the Gene and Ruth Posner Foundation, the RD and Linda Peters Foundation, Research to Prevent Blindness, and NIH (R01EY017607, P30EY001931, T32EY014537). P. Godara is supported by a research award from the VitreoRetinal Surgery Foundation. J. Carroll and A. Dubra are recipients of Career Development Awards from RPB. A. Dubra holds a Career Award at the Scientific Interface from the Burroughs Wellcome This investigation was conducted in a facility constructed with support from the Research Facilities Improvement Program, grant number C06 RR016511, from the National Center for Research Resources, National Institutes of Health.

The authors would like to acknowledge Phyllis Summerfelt for her technical/administrative assistance in figure preparation. Dr. Stepien had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

## Footnotes

Presented in part at the Association for Research in Vision and Ophthalmology annual meeting 2012, Fort Lauderdale, FL, poster session, May 9<sup>th</sup>, 2012.

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