

Targeted deletion of *Vegfa* in adult mice induces vision loss

Toshihide Kurihara, Peter D. Westenskow, Stephen Bravo, Edith Aguilar, and Martin Friedlander

Department of Cell Biology, The Scripps Research Institute, La Jolla, California, USA.

Current therapies directed at controlling vascular abnormalities in cancers and neovascular eye diseases target VEGF and can slow the progression of these diseases. While the critical role of VEGF in development has been well described, the function of locally synthesized VEGF in the adult eye is incompletely understood. Here, we show that conditionally knocking out *Vegfa* in adult mouse retinal pigmented epithelial (RPE) cells, which regulate retinal homeostasis, rapidly leads to vision loss and ablation of the choriocapillaris, the major blood supply for the outer retina and photoreceptor cells. This deletion also caused rapid dysfunction of cone photoreceptors, the cells responsible for fine visual acuity and color vision. Furthermore, *Vegfa* deletion showed significant downregulation of multiple angiogenic genes in both physiological and pathological states, whereas the deletion of the upstream regulatory transcriptional factors HIFs did not affect the physiological expressions of angiogenic genes. These results suggest that endogenous VEGF provides critical trophic support necessary for retinal function. Targeting factors upstream of VEGF, such as HIFs, may be therapeutically advantageous compared with more potent and selective VEGF antagonists, which may have more off-target inhibitory trophic effects.

Introduction

Phototransduction by rod and cone photoreceptors in the retina converts light into electrical energy. This process, forming the basis for vision, is facilitated by the vascular plexus (the choriocapillaris) and retinal pigmented epithelial (RPE) cells, both of which underlie and are intimately associated with the photoreceptors. Degeneration of photoreceptors, RPE cells, and the choriocapillaris is associated with age-related macular degeneration (AMD), the leading cause of vision loss in industrialized nations. While VEGF clearly plays a role in physiological vascular development (1-4) and pathological neovascularization (5, 6) in the retina and the choroid, there is increasing evidence that it is also important for the trophic maintenance of neurons (7–9). Inhibition of VEGF has become the mainstay of therapy for treating AMD (10-12), and increasingly potent VEGF antagonists have recently been introduced into the clinics. In order to better understand the trophic activity of VEGF in the adult retina and choriocapillaris and to assess the effect of deleting its expression by RPE cells, we utilized inducible Cre/loxP technology to genetically delete floxed Vegf alleles in vivo with an RPE-specific VMD2-Cre transgenic line (13).

Results and Discussion

Three days after RPE-specific *Vegf* inactivation, we observed complete ablation of the choriocapillaris (Figure 1A). Since almost the entire blood supply to the outer two-thirds of the retina (including the photoreceptors) is provided by the choriocapillaris, we predicted that its loss would induce dramatic secondary effects on the retinal neurons. In fact, following *Vegfa* inactivation in RPE cells, we observed profound retinal dysfunction, especially in cone photoreceptors. Within 3 days after induction, rapid and progressive dysfunction of cone photoreceptors was detected

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using electroretinography (ERG) (Figure 1B) compared with multiple controls, including Vegfaff (no Cre), Vegfaff; VMD2-Cre, Vegfaf/+;VMD2-Cre with doxycycline, and Vegfaf/f;VMD2-Cre without doxycycline (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI65157DS1). Using immunohistochemistry to selectively stain for the cone photopigment opsin, we observed a nearly complete absence of cone photoreceptor outer segments (Figure 1C). Cone dysfunction was observed by at least 7 months after Vegfa deletion in RPE (Supplemental Figure 2). These data demonstrate that RPE-derived VEGF is required for maintenance of the choriocapillaris and for cone photoreceptors, the most metabolically demanding cells in the central nervous system (14, 15). Photoreceptors also express VEGF receptors (9, 16), and paracrine-derived VEGF may have a direct role in their maintenance. In humans, loss of cones results in legal blindness, since fine (or "reading") visual acuity is maintained by cones located in the central retina, or macula.

In contrast, rod photoreceptor function (responsible for achromatic and night vision) is insensitive to Vegfa inactivation at the same time points when we observe cone degeneration. Immunohistochemical analyses revealed that the layer of rod photoreceptor outer segments was as thick as that of controls (Figure 2A). Since the murine retina is composed largely of rods and the thickness of the entire retina does not change as late as 7 months after Cre recombinase induction (Figure 2B), this suggests that there is no significant rod photoreceptor atrophy. Scotopic (dark adapted) ERGs demonstrate that rod photoreceptors are fully functional, since there was no change in rod photoreceptor light responsiveness in recordings before and after induction in Vegfa mutant mice (Figure 2C). The preservation of rod photoreceptors for several months after gene deletion suggests that these cells are supported by a minimal vasculature not dependent on RPE-derived Vegfa for its survival. Indocyanine green angiography demonstrated 2 blood supply sources for photoreceptors in Vegfa mutants; intraretinal and outer, major choroidal vessels (Figure 2D). Utilizing electron microscopy, we determined that



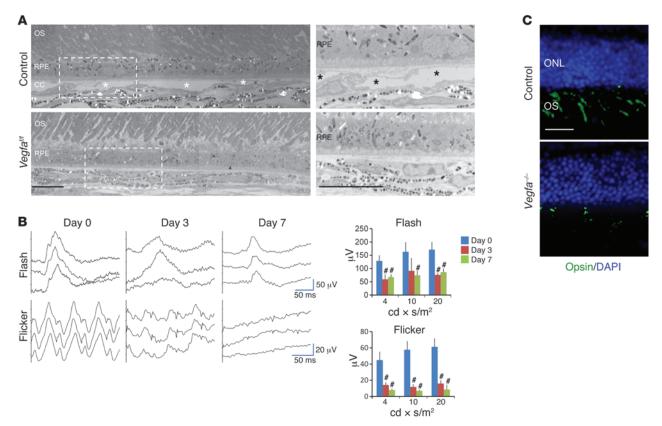


Figure 1
Inducible Vegfa deletion in adult RPE cells promotes rapid choriocapillaris degeneration and vision loss. (**A**) Electron micrographs of control (upper panel, $Vegfa^{iff}$ without Cre) and mutant (lower panels, $Vegfa^{iff}$ with VMD2-Cre) murine retinas 3 days after Vegfa deletion in adult RPE. Right panels are enlargements of the boxed regions of the left panels. Note that Vegfa mutants lack choriocapillaris normally observed in controls (asterisks). (**B**) ERG of Vegfa mutant eyes shows loss of photopic and flicker signals. b-wave amplitudes from flash, or first peaks from flicker, ERG in photopic light-adapted conditions captured in Vegfa mutants are significantly attenuated compared with the same retina prior to Vegfa gene deletion (n = 4 for each time point). This dramatic attenuation is consistent with vision loss. (**C**) Immunohistochemical analyses for cone-opsin in Vegfa mutants and controls 7 days after induction. The absence of cone outer segments is apparent in the Vegfa mutants. Vegfa mutants. Vegfa mutants. Vegfa mutants. Vegfa mutants and controls 7 days after induction. The absence of cone outer segments is apparent in the Vegfa mutants. Vegfa mutants. Vegfa mutants indicate mean Vegfa mutants. Vegfa mutants and controls 7 days after induction. The absence of cone outer segments is apparent in the Vegfa mutants. Vegfa mutants. Vegfa mutants indicate mean Vegfa mutants. Vegfa mutants are significantly attenuated compared with the same retina prior to Vegfa gene deletion (Vegfa mutants and controls 7 days after induction. The absence of cone outer segments is apparent in the Vegfa mutants.

the choriocapillaris remained atrophic through 7 months and choroidal vessels were observed only sporadically (Figure 2E). These findings are consistent with results from a previous study that examined the effects of global deletion of the soluble VEGF isoforms (VEGF^{188/188} mice) (17).

Atrophy and thinning of the choriocapillaris is a known agerelated event in humans (18). To compensate for choriocapillaris thinning, the RPE is known to synthesize proangiogenic factors that maintain the choriocapillaris (19, 20). Our findings suggest that VEGF may act both directly and indirectly to regulate this phenomenon. Using gene-profiling assays, we demonstrated that the loss of Vegfa affects the expression of multiple angiogenesisrelated genes (most are downregulated) in isolated RPE/choroid preparations (Figure 2F and Supplemental Table 1). Vegfa is tightly regulated by HIFs during development and pathologically in disease states (21, 22). To determine the role for HIFs in maintenance of the choriocapillaris, we genetically ablated Hif1a (encoding HIF-1α protein), Epas1 (HIF-2α), and Hif1a/Epas1 using VMD2-Cre in adult RPE. No obvious morphological, functional, or transcriptional differences, including Vegfa mRNA, were observed in naive-state adult HIF mutants (Supplemental Figures 3 and 4, and Supplemental Table 1). Several pathways, such as NF-κB/JunB and PGC1α/ERRα are known to induce *Vegfa* gene expression independently of HIFs (23, 24). These factors, or others, rather may maintain VEGF levels at sufficient levels in healthy retinas. Using laser photocoagulation to induce choroidal neovascularization (CNV), we determined that the extent of CNV was significantly reduced in *Vegfa* and *Epas1* mutants (and partially reduced in *Hif1a* mutants; Figure 3, A and B). The upregulation of angiogenic genes was observed in wild-type C57BL/6 mice, but normal expression of these genes was observed in HIF mutants and expression was attenuated in *Vegfa* mutants compared with naive controls (Figure 3C and Supplemental Table 2). These data strongly suggest that HIFs become activated in pathological states to promote VEGF-mediated neovascularization, but are not required to maintain physiological vasculature.

VEGF-mediated neovascularization is associated with vision loss in diseases such as AMD (10–12). The majority of patients receiving currently approved anti-VEGF drugs to treat AMD experience no adverse events. However, long-term safety studies of the effects of VEGF antagonism in the eye have not yet been completed, and there are isolated clinical reports showing progressive RPE/photoreceptor atrophy in patients on anti-VEGF therapies (25). Due to the differences in rod and cone photoreceptor distribution in



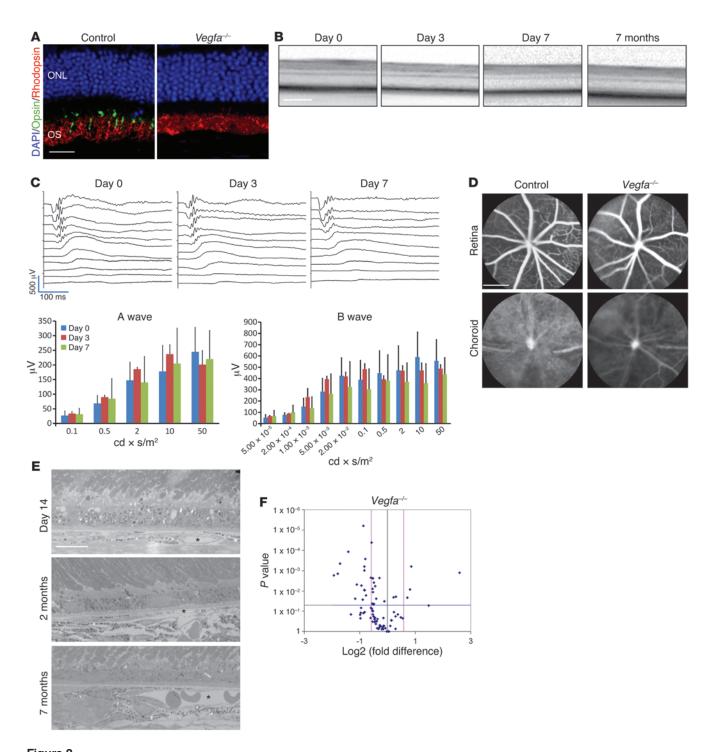
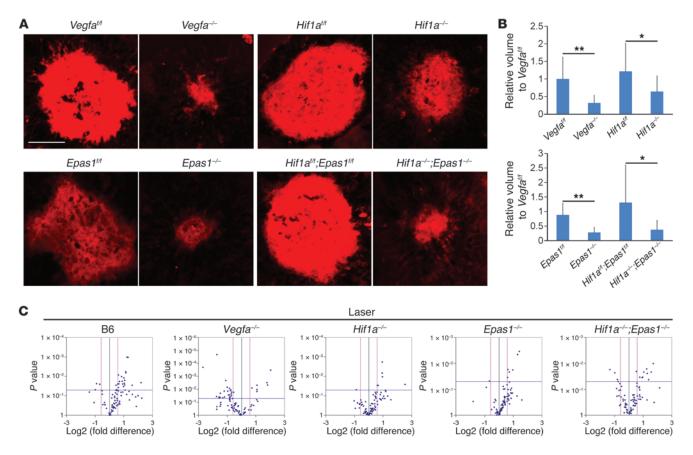


Figure 2

Cone but not rod photoreceptor dysfunction in RPE-specific Vegfa mutants. (**A**) Immunohistochemistry for rhodopsin (rods) and cone opsin (cones) 45 days after induction. Note that rhodopsin expression in Vegfa mutants is comparable to that observed in controls. (**B**) Optical coherence tomographic analysis indicates that retinal thicknesses remain relatively unchanged at all stages through 7 months. (**C**) Scotopic dark-adapted ERG captured in Vegfa mutants 3 or 7 days after induction. Note that no significant reduction in the a- and b-wave amplitudes is observed (n = 4). (**D**) Indocyanine green angiography for Vegfa mutants 45 days after induction. Note that choroidal circulation is significantly diminished in the mutants, while the retinal circulation is normal. (**E**) Electron micrograph of a cross-sectioned Vegfa mutant retina 14 days, 2 months, and 7 months after induction. Note that a few deep choroidal vessels are observed (asterisk). (**F**) mRNA array for angiogenic genes in Vegfa mutant RPE/choroids compared with controls 3 days after injection (n = 3). Fold-change (x axis) and y value (y axis) of gene expression compared with controls are shown. Note that 32 of 84 genes are significantly downregulated. Error bars indicate mean y Scale bars: 2,000 y m (**B**); 20 y m (**B**); 20 y m (**C**); 200 y





Attenuated angiogenic gene regulation in pathological states of Vegfa conditional mutants. (**A**) Representative Z projections of laser-CNV lesions (stained with GS lectin). (**B**) Quantified lesion volumes of laser-CNV (n = 8-21). Note that laser-CNV is only partially reduced in the Hif1a mutants, while it is completely inhibited in the Epas1 and Hif1a; Epas1 mutants (comparable to that observed in the Vegfa mutants). (**C**) mRNA array for angiogenic genes in Hif1a, Epas1, and Hif1a; Epas1 mutant RPE/choroids compared with controls in laser-irradiated C57BL/6 wild-type, Vegfa, Hif1a, Epas1, and Hif1a; Epas1 mutants compared with naive state controls (n = 3). Fold-change (x axis) and P value (y axis) of gene expression compared with controls. Note that few or no significantly dysregulated genes are observed in naive state Hif1a, Epas1, or Hif1a; Epas1 mutants. Twenty-two upregulated and 3 downregulated angiogenic genes are observed in laser-irradiated B6 wild-type mice, whereas we observed 7 (up) and 23 (down) in Vegfa mutants, 6 (up) and 1 (down) in Vegfa mutants, and 10 (up) and 4 (down) in Vegfa mutants. Vegfa mutants, V

humans and mice, it is difficult to speculate on what the effect of near-complete VEGF antagonism may be in human subjects. However, if a certain threshold of VEGF could be neutralized, the fovea would be very susceptible and the loss of macular function could result in the devastating loss of central vision. These observations, coupled with reported renal, gastrointestinal (GI), and mucosal complications in patients on systemic VEGF inhibition, contribute to the concern that VEGF antagonism may not be completely effective or safe (26, 27). The data in this study reinforce this concept and suggest that HIF-targeted or combination therapies targeting multiple angiogenic pathways may be safer and more effective than increased VEGF suppression therapy. Newer VEGF antagonists that bind VEGF with higher affinities and less rapid clearing from the eye are coming into use to treat a variety of neovascular diseases (28). Our results suggest that "off-target" effects of potent, persistent VEGF antagonism in the eye may exacerbate degeneration of the choriocapillaris, RPE, and photoreceptors due to the diminished trophic effect of RPE-produced VEGF.

Methods

Animals. Transgenic mice carrying the human vitelliform macular dystrophy-2 (VMD2) promoter-directed reverse tetracycline-dependent transactivator (rtTA) and the tetracycline-responsive element-directed (TRE-directed) Cre recombinase (VMD2-Cre mice) (13) (provided by Y.Z. Le of University of Oklahoma Health, Sciences Center, Oklahoma City, Oklahoma, USA) were mated with VegfaVf mice (29), Hif1aVf mice (30), or Epas Vf mice (31) (provided by R.S. Johnson of University of Cambridge, Cambridge, United Kingdom).

Laboratory measures. Electron microscopy, ganzfeld ERG, immunohistochemistry, in vivo imaging, and laser-induced CNV were performed as previously described (32–34). mRNA PCR arrays for 84 angiogenic genes (Mouse Angiogenesis RT² Profiler PCR Array, PAMM-024; QIAGEN) were performed according to the manufacturer's instructions. Please see Supplemental Methods for more details.

Statistics. Comparisons between the mean variables of 2 groups were performed by a 2-tailed Student's t test. $P \le 0.05$ was considered significant. Please see Supplemental Methods for more details.



Study approval. All procedures involving animals were approved by the Scripps Research Institute Animal Care Committee, which ensures that all federal animal experimentation guidelines are met.

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Address correspondence to: Martin Friedlander, The Scripps Research Institute, MB 28, 10550 North Torrey Pines Road, La Jolla, California 92037, USA. Phone: 858.784.9138; Fax: 858.784.9135; E-mail: friedlan@scripps.edu.

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