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VEGF-B Is an Autocrine Gliotrophic Factor for Müller Cells under Pathologic Conditions

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Purpose. Müller glia are important in retinal health and disease and are a major source of retinal VEGF-A. Of the different VEGF family members, the role of VEGF-A in retinal health and disease has been studied extensively. The potential contribution of other VEGF family members to retinal pathophysiology, however, remains poorly defined. This study aimed to understand the role of VEGF-B in Müller cell pathophysiology.

Methods. The expression of different VEGFs and their receptors in human MIO-M1 and mouse QMMuC-1 Müller cell lines and primary murine Müller cells was examined by RT-PCR, ELISA, and Western blot. The effect of recombinant VEGF-B or VEGF-B neutralization on Müller cell viability and survival under normal, hypoxic, and oxidative (4-hydroxynonenal [4-HNE]) conditions was evaluated by Alamar Blue, Yo-Pro uptake, and immunocytochemistry. The expression of glial fibrillary acidic protein, aquaporin-4, inward rectifying K+ channel subtype 4.1, glutamine synthetase, and transient receptor potential vanilloid 4 under different treatment conditions was examined by RT-PCR, immunocytochemistry, and Western blot. Transient receptor potential vanilloid 4 channel activity was assessed using a Fura-2–based calcium assay.

Results. VEGF-B was expressed in Müller cells at the highest levels compared with other members of the VEGF family. VEGF-B neutralization did not affect Müller cell viability or functionality under normal conditions, but enhanced hypoxia– or 4-HNE–induced Müller cell death and decreased inward rectifying K+ channel subtype 4.1 and aquaporin-4 expression. Recombinant VEGF-B restored Müller cell glutamine synthetase expression under hypoxic conditions and protected Müller cells from 4-HNE–induced damage by normalizing transient receptor potential vanilloid 4 channel expression and activity.

Conclusions. Autocrine production of VEGF-B protects Müller cells under pathologic conditions.

Keywords: Mueller cell, VEGF-B, hypoxia, retina, oxidative stress

Müller cells are the principal macroglia in the retina. Their unique position, extending from the ganglion cell layer to the photoreceptor inner segments, facilitates their physical contact with many cell types in the retina. Müller cells have a pivotal role in the maintenance of retinal homeostasis, architecture, and responses to stress conditions. Müller cell functions include blood–retinal barrier regulation, release of trophic factors, neurotransmitter recycling (e.g., glutamate, γ-aminobutyric acid), control of water and ion (mainly K+) homeostasis and functional hyperemia, as well as regulation of retinal innate immunity and the visual cycle. In pathologic conditions such as diabetic retinopathy, Müller cells contribute to neurovascular dysfunction by producing inflammatory and angiogenic factors. In particular, Müller cells are a major source of VEGF-A, in the retina under pathophysiologic conditions. The VEGF family consists of five members (VEGF-A, B, C, D and placental growth factor [PIGF]), which are all structurally related. VEGF ligands bind three different tyrosine kinase receptors (VEGFR1, VEGFR2, and VEGFR3), and two neuropilin coreceptors (NRP1 and NRP2) that enhance the affinity of the ligands to VEGFR receptors, thereby enhancing their physiologic actions. The VEGF family members are distinguished by their receptor-binding pattern and unique functions; for example, human VEGF-C and -D bind to VEGFR2 and VEGFR3, although their function lymphangiogenesis role seems to be mediated predominantly by VEGFR3. Mouse VEGF-D only binds to VEGFR3 but not VEGFR2. Similarly, placental growth factor has been described to participate only in pathologic angiogenesis by binding to VEGFR1, whereas VEGF-A, the best understood member of this family, binds mainly to VEGFR2 to elicit angiogenic activity. VEGF-B, the most elusive member of the VEGF family, binds exclusively to VEGFR1, although its functions remain to be fully elucidated.

Recently, the neuroprotective role of different VEGF family members has received increasing attention. The use of VEGF-A as a neuroprotective therapy, however, has been hindered by its proangiogenic effects. There is some preclinical evidence that VEGF-B is protective in...
pathologies in which oxidative stress is a key pathologic factor, including amyotrophic lateral sclerosis, Parkinson’s disease, retinitis pigmentosa and Alzheimer’s disease. An association between VEGF-B and advanced stages of diabetic retinopathy has also been described, although the cellular source of retinal VEGF-B and exactly how VEGF-B contributes to retinal disease remain elusive.

The current study investigated the role of VEGF family members in Müller glia pathophysiology. We found that, among the different VEGF family members, Müller cells produce more VEGF-B than other VEGF members, and that VEGF-B protects Müller cells under oxidative and hypoxic conditions.

**METHODS**

**Cell Culture and Treatments**

QMMuC-1, an immortalized murine Müller cell line and the human Müller cell line Moorfields Institute of Ophthalmology-Müller 1 (MIO-M1) (courtesy of Dr. G.A. Limb, UCL Institute of Ophthalmology, London, UK) were cultured in Dulbecco’s Modified Eagle Media (Thermo Fisher Scientific, Waltham, MA; Catalog# 41965039) supplemented with 10% fetal calf serum. Primary Müller cells (PMC) were isolated from C57BL/6J neonatal mice (P7) and cultured in Dulbecco’s Modified Eagle Media (Thermo Fisher Scientific, Waltham, MA) supplemented with 10% fetal calf serum. Primary Müller cells (PMC) were used in the study.

To induce different stress conditions and investigate the effect of VEGF-A or VEGF-B and their receptors, Müller cells were treated with the compounds listed in Table 1.

**Quantitative RT-PCR**

Total RNA from murine Müller cells and human MIO-M1 cells was extracted using RNeasy Mini Kit (Qiagen, Hilden, Germany) and TRIzol Reagent (Invitrogen, Carlsbad, CA, USA), respectively, according to the manufacturer’s instructions. cDNA was synthesized using SuperScript II Reverse Transcripase Kit (Invitrogen). Validated TaqMan probes were purchased from Roche (Basel, Switzerland). SYBR Green gene specific primers were designed using NCBI Primer Blast and validated by Integrated DNA Technologies (Coralville, IA) (Table 2).

Real-time RT-PCR was performed using LightCycler 480 (Roche-Diagnostic) in triplicate. Relative gene expression was calculated using the comparative Ct method ($2^{-\Delta\Delta Ct}$) with data normalized to $\beta$-actin or 18s.

**Western Blot**

Samples were homogenized in RIPA buffer containing protease inhibitor cocktail. Protein concentration was determined using a Pierce BCA protein assay kit (Thermo Fisher Scientific, Catalog# 23225) or Bradford Assay (Bio-Rad Laboratories, Hercules, CA). The blot was performed using 15 to 20 μg of protein according to previously described methods. Primary and secondary antibodies used are detailed in Table 3. Membranes were visualized with enhanced chemiluminescence (Clarity Western ECL Blotting Substrates; Bio-Rad Laboratories) and bands detected using Syngene G-Box imaging system (Syngene, Cambridge, UK).

**ELISA**

ELISA kits for VEGF-A and VEGF-B (R&D, DY493 and DY471), NRPI and VEGF-B (Abcam, Cambridge, UK; ab213880 and ab213897) were used according to the manufacturers’ instructions. VEGFR2 indirect ELISA was performed by adding 20 μg protein or supernatants to the wells and incubated overnight. After several washes and 2 hours of blocking (SuperBlock TBS, Thermo Fisher Scientific), anti-VEGFR2 (1 μg/mL, R&D af644) antibody was added and incubated for 2 hours at room temperature. After thorough washes, wells were incubated with a biotinylated horse anti-goat IgG antibody (1:8,000, Agilent DAKO, Santa Clara, CA) for 2 hours, followed by with HRP-conjugated streptavidin (1:900, DAKO, 20 minutes). The signal was developed by adding TMB (20 minutes, 37°C). Absorbance values were measured at 450 nm/570 nm using a POLARstar OMEGA plate reader (BMG LabTech, Aylesbury, UK). All values were collected from at least four independent samples and each experiment repeated in duplicate.

**Cell Viability and Apoptosis**

The viability of Müller cells under different conditions was assessed using the AlamarBlue (Thermo Fisher) and YO-PRO uptake assay (YO-PRO-1 Iodide (491/509), Thermo Scientific).
Table 2. Primer Sequences: Genes Whose Expression Was Examined in Müller Cells in Normal or Stress Conditions

<table>
<thead>
<tr>
<th>Genes</th>
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<td>AQP4</td>
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Table 3. Antibodies Used for Western Blotting and Immunofluorescence

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<td>1:500(WB)/1:50 (IF)</td>
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<tr>
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Immunofluorescence Staining

Caspase-3/7 expression was measured using the CellEvent caspase-3/7 Green Detection Reagent (Invitrogen) following the manufacturer's instructions. The number of caspase-3/7 positive cells was analyzed using ImageJ.

Intracellular Ca^{2+} Measurements

Intracellular Ca^{2+} concentration ([Ca^{2+}]_{i}) was measured in QMMuC-1 mouse Müller cells cultured in black 96-well plates (Corning Inc., Corning, NY, Catalog# 3603) using a Fura-2 QBT Calcium kit (Molecular Devices, Fisher) according to the manufacturer's instructions. Fluorescence signal was read in a plate reader (POLARStar Omega, BMG Labtech). After thorough washes, samples were incubated with Alexa Fluor-conjugated goat anti-rabbit IgG for 1 hour at room temperature. For F-actin staining, Alexa Fluor 594 Phalloidin (Thermo Fisher Scientific) was used. Cells were mounted using Vectashield medium containing DAPI (Vector Laboratories, Burlingame, CA) and imaged using a Nikon C1 Eclipse TE200-U (Nikon, Surrey, UK). Images were processed using ImageJ software.
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**Results**

The Expression Profile of Different VEGF Family Members and Their Receptors in Müller Cells

Real-time RT-PCR showed that under normal culture conditions the most abundantly expressed VEGF family members in mouse QMMuC-1 Müller cells (Fig. 1A) and PMCs (Fig. 1B) were VEGF-B and D. VEGF-A was detected at low levels in both QMMuC-1 and PMC. PlGF was absent in QMMuC-1 but expressed at low levels in PMC (Figs. 1A, 1B). In human MIO-M1 Müller cells, VEGF-B expression was also higher than VEGF-A and VEGF-C, VEGF-D and PlGF (Fig. 1C).

Using ELISA, we confirmed that a significantly higher level of VEGF-B was detected in QMMuC-1 Müller cell lysates compared with VEGF-A (Fig. 1D) although the levels of VEGF-B and VEGF-A in the supernatants did not differ (Supplementary Fig. S1A) under normal culture conditions.

In terms of VEGF receptors, mRNA expression of the coreceptor, NRP1, was markedly higher than VEGFR1 and VEGFR2 in all Müller cell types (Figs. 1E–1G). VEGFR1

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**Statistics**

All graphing and statistical analysis was performed using Prism 6 (GraphPad Software, San Diego, CA). An unpaired Student’s *t* test, one-way ANOVA, followed by Dunnett’s or Newman-Keuls or two-way ANOVA with Tukey’s multiple comparison post-test were utilized to determine the statistical outcome; a *P* value of less than 0.05 was considered statistically significant.
mRNA expression was higher than VEGF2 in murine (Figs. 1E, 1F) but not human (Fig. 1G) Müller cells. However, at the protein level, VEGFR1 (38,368 ± 9552 pg/mg) was significantly higher than the coreceptor NRP1 (10,171 ± 830 pg/mg), and the latter significantly higher than VEGFR2 (2107.0 ± 115.9 pg/mg) in QMMuC-1 Müller cells (Fig. 1H).

Our data suggest that the VEGF-B and its receptors, VEGFR1 and NRP1, are highly expressed at the mRNA and protein levels by Müller cells and hence may play a role in Müller cell physiology.

Expression of VEGF-B and Its Receptors Is Affected by Hypoxia and Oxidative Stress

To understand how the expression of VEGF-B and its cognate receptors is altered in Müller cells under conditions resembling retinal disease, QMMuC-1 Müller cells were exposed to hypoxia, hyperglycemia, the inflammatory IL-1β, or oxidative stress. To mimic hyperglycemia that occurs in diabetes, an additional 25 mM of D-glucose was added to the media for 72 hours as previously reported.25,26 This concentration and length did not induce any change in MIO-M1 cell viability (data not shown). A sustained exposure to 1% O2 for 72 hours was used to recapitulate hypoxic conditions.27,28 24 hours treatment with 10 μM 4-hydroxynonenal (4-HNE) was used to induce oxidative stress without reduction in viability (Supplementary Fig. S1B) and 20 ng/mL of IL-1β was used based on previous studies in literature.29,30

Hypoxia significantly upregulated VEGF-B and VEGFR1 mRNA expression (Figs. 2A and 2F) and increased the soluble forms of VEGF-B (Fig. 2C) and VEGFR1 (sFlt-1) (Fig. 2H) proteins. NRP1 expression, however, was decreased under hypoxic conditions (Fig. 2E). These results were also confirmed in MIO-M1 cells (Supplementary Fig. S1F).

The 4-HNE increased VEGF-B and its receptors NRP1 and VEGFR1 mRNA expression in QMMuC-1 cells (Figs. 2A, 2D, and 2F). However, at the protein level, only VEGF-B (Fig. 2B) was increased by 4-HNE treatment. Regarding other VEGF family members (Supplementary Figs. S1C-D), only VEGF-A mRNA was augmented under hypoxia and after the IL-1β exposure (Supplementary Fig. S1C), as previously reported.31
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FIGURE 3. The effect of VEGF-B and its receptors on Müller cell viability under oxidative stress and hypoxic conditions. (A) Effect of the neutralization of VEGF-A and its receptors (VEGFR2 and NRP1) and VEGF-B and its receptors (VEGFR1 and NRP1) on QMMuC-1 Müller cell viability under hypoxic conditions (1% O2 for 72 hours). *P < .05; **P < .01 compared with control (hypoxia only), one-way ANOVA followed by Dunnett’s multiple comparison test. (B) Effect of VEGF-A or VEGF-B neutralization on 4-HNE–induced QMMuC-1 Müller cell viability. C, Effect of the neutralization of VEGFR1 and NRP1 on QMMuC-1 and their receptor (VEGFR1 and NRP1) on 4-HNE–induced QMMuC-1 damage. (D) The effect of recombinant VEGF-B and VEGF-A on 4-HNE–induced cell death in QMMuC-1 and PMC Müller cells. The viability of cells was determined by Alamar Blue methods. (A) n = 5–9, (B and C) n = 4–6, (D) n = 5–10 per group. *P < 0.05; **P < 0.01; ***P < 0.005 and ****P < 0.001, one-way ANOVA followed by Newmann-Keuls post hoc test. Antibody, AB. A and B are VEGF-A and VEGF-B respectively. R1 and R2 are VEGFR1 and VEGFR2. Anti-VEGF-A, VEGFR2 500 ng/mL and 10 μg/mL, respectively; NRP1 antibody, 30 μg/mL; VEGF-B and VEGFR1, 500 ng/mL and 10 μg/mL, respectively; recombinant VEGF-A and VEGF-B, 100 ng/mL; 4-HNE 20 μM.

Blocking VEGF-B Does Not Alter Müller Cell Homeostasis Under Normal Culture Conditions

To ascertain the role of VEGF-B in normal Müller cell function, VEGF-B was neutralized in QMMuC-1 cultures using a neutralizing antibody. The expression of proteins responsible for water and ion homeostasis (inward rectifying K+ channel subtype 4.1 [Kir4.1], aquaporin-4 [AQP4]), glutamate metabolism (glutamine synthetase [GS]) and Müller cell activation (vimentin and glial fibrillary acidic protein [GFAP]) was assessed. Our results showed that blockade of VEGF-B for 24 hours (Supplementary Figs. S2A-D) or 72 hours (Supplementary Figs. S2E–S2H) had no effect on the expression of those proteins.

We also examined QMMuC-1 cell viability after blockade of VEGF-B, NRP1, and VEGFR1, or both VEGFR1 and NRP1 for 72 hours. None of these treatments affected QMMuC-1 cell viability (Supplementary Fig. S2I), indicating that the VEGF-B pathway is not essential for Müller cell survival under normal culture conditions.

VEGF-B Neutralization Compromises Müller Cell Viability and Survival Under Pathologic Conditions

To assess whether VEGF-B plays a role in Müller cell responses to stress, VEGF-B or its receptors were neutralized or inhibited in QMMuC-1 cells cultured in hypoxia or in the presence of 20 μM 4-HNE (dose inducing approximately 15%–25% decrease of viability, Supplementary Fig. S1B) for 72 or 24 hours, respectively, and cell viability measured. Concentration of anti-VEGF-A and -B antibodies, and VEGFR1, VEGFR2, and NRP1 was chosen according to the company’s recommendation and in line with others in the same way as the selected the doses of recombinant VEGF-A and -B.35,34,37,38
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Under hypoxic conditions (Fig. 3A), neutralization of VEGF-B (but not VEGF-A), significantly decreased Müller cell survival. Blocking VEGFR1, VEGFR2, or NRP1 alone or the combination of VEGFR2 and NRP1 did not affect Müller cell survival (Fig. 3A). However, when both VEGFR1 and NRP1 were blocked, cell viability was significantly reduced to a level similar to that observed with VEGF-B neutralization.

Treatment with 4-HNE decreased QMMuC-1 cell viability by 15% and blocking VEGF-A or VEGF-B further decreased cell viability by 9% and 25%, respectively (Fig. 3B), and similar levels were reached when blocking both VEGFR1 and NRP1 together (Fig. 3C). The results were confirmed in PMCs (Supplementary Fig. S3A and S3B). Yo-Pro uptake assay showed that VEGF-B, but not VEGF-A neutralization, increased 4-HNE-induced apoptosis in QMMuC-1 cells (Supplementary Fig. S3C) via the VEGFR1 and NRP1 pathways (Supplementary Fig. S3D).

Next, the cytoprotective effect of VEGF-B was evaluated (Fig. 3D). Recombinant VEGF-B, but not VEGF-A, significantly increased the viability of Müller cell line QMMuC-1 and PMCs treated with 4-HNE (Fig. 3D).

Our data suggest that the VEGF-B/VEGFR1/NRP1 signaling axis is critically involved in Müller cell survival under hypoxic or oxidative conditions.

VEGF-B Neutralization Enhances Caspase-3 Expression in Müller Cells Under Oxidative Stress Conditions

To determine the mechanism underlying the decrease in Müller cell viability after VEGF-B pathway blockade, proapoptotic caspase-3 levels were analyzed. As shown in Figure 4A, hypoxia treatment did not affect caspase-3 mRNA expression in QMMuC-1 cells, even after VEGF-B neutralization (Fig. 4A). We found that 4-HNE increased caspase-3 expression and neutralization of VEGF-B further increased caspase-3 mRNA expression and its cleavage.
FIGURE 5. The effect of VEGF-B on GS, Kir4.1 and AQP4 expression in Müller cells under hypoxia and oxidative stress. (A) QMMuC-1 cells were treated with recombinant VEGF-A or VEGF-B and the expression of GS was measured by Western blotting. (B) Western Blot analysis of GS expression in QMMuC-1 cells cultured in hypoxic conditions for 72 hours with or without recombinant VEGF-A or VEGF-B. (C and D) Kir4.1 mRNA (C) and protein (D) expression in hypoxia (1% O2 for 72 hours) treated QMMuC-1 (C, D, left) and PMCs (D, right) with or without VEGF-B neutralizing antibody. (E and F) AQP4 mRNA (E) and protein (F) expression in QMMuC-1 cells in hypoxia condition.
with or without VEGF-B neutralization. (G) GS expression in QMMuC-1 cells under oxidative conditions (4-HNE) with/without VEGF-B neutralization. (H) and D) AQP4 mRNA (H) and protein (I) expression in 4-HNE treated QMMuC-1 with or without VEGF-B neutralization. VEGF-B antibody, 500 ng/mL. 4-HNE, 10 μM. Data normalized to control values. n = 3–5/group in qPCR; n = 4–10 in Western Blot. *P < 0.05; **P < 0.01; ***P < 0.005; ****P < 0.001. One-way ANOVA followed by Newman-Keuls post hoc test. AB, antibody. Recomb, recombinant.

(Figs. 4B, 4C). Simultaneous blocking of VEGFR1 and NRP1 also significantly increased caspase 3/7 expression in 4-HNE exposed cells (Figs. 4D and 4E).

Role of VEGF-B in Müller Cell Gliosis and Ion/Water Channel Expression Under Stress Conditions

Next, we assessed whether VEGF-B can modulate Müller cell pathologic responses to hypoxia and oxidative stress. Hypoxia did not affect GFAP mRNA and protein expression in Müller cells (Supplementary Figs. S4A and S4B). Decreased GS expression in hypoxia was not affected by VEGF-B neutralization (Supplementary Fig. S4C); however, the addition of recombinant VEGF-B completely prevented hypoxia-induced downregulation of GS (Fig. 5B). This increase in GS with VEGF-B was only observed in hypoxia, because the addition of VEGF-A and VEGF-B recombinant proteins under normal culture conditions did not elicit any changes in GS expression (Fig. 5A). Kir4.1 and AQP4 were not affected by hypoxia treatment but were downregulated after VEGF-B neutralization (Figs. 5C–F). Our results suggest that autocrine VEGF-B is important for Müller cells to maintain the expression of important proteins involved in ion/water homeostasis, whereas exogenous VEGF-B may help to maintain Müller cell glutamate metabolism under hypoxic conditions.

VEGF-B blockade did not modify the 4-HNE–induced increased mRNA or protein GFAP expression (Supplementary Fig. S4D and S4E). GS was upregulated by 4-HNE and this was abrogated by VEGF-B neutralization (Fig. 5G). Downregulated protein expression of Kir4.1 after 4-HNE was not further modified by VEGF-B neutralization (Supplementary Fig. S4G). AQP4 was not affected by 4-HNE when VEGF-B was present, but was downregulated when VEGF-B was neutralized (Figs. 5H and 5I). Our findings suggest that autocrine VEGF-B is important for Müller cells to maintain ion/water homeostasis and glutamate recycling under oxidative stress conditions.

Role of VEGF-B in Müller Cell TRPV4 Expression and Function Under Pathologic Conditions

Apart from AQP4 and Kir4.1, the osmosensor, TRPV4, is also critically involved in Müller cell ion/water homeostasis.55–65 Treatment with 4-HNE did not affect the expression level of TRPV4 protein in Müller cells (Fig. 6C). The treatment, however, altered the cellular distribution of TRPV4 in Müller cells (Figs. 6A and 6B). Under normal culture conditions, QMMuC-1 cells spread out on the culture dish and TRPV4 was distributed uniformly across most of the cells (Fig. 6A). Aggregation of TRPV4 proteins was occasionally observed under normal culture conditions (Fig. 6A). After 4-HNE treatment, many QMMuC-1 cells displayed contracted cell bodies with long-handed shaped actin filaments and TRPV4 channel aggregation (arrows in Fig. 6B). VEGF-B-protected Müller cells from these changes (Figs. 6A and 6B).

To further explore the role of TRPV4 in 4-HNE–induced Müller cell damage, intracellular Ca2+ recording was conducted. Under normal culture conditions, the TRPV4 agonist GSK101 did not affect [Ca2+]i in QMMuC-1 cells (Figs. 6D and 6E). Treatment with 4-HNE alone had no effect on [Ca2+]i. However, preincubation of the cells with this aldehyde facilitated [Ca2+]i response to GSK101 (Fig. 6D), which was fully blocked by the TRPV4 inhibitor, HC06, confirming the involvement of these channels (Figs. 6D and 6E). VEGF-B completely abrogated TRPV4-dependent [Ca2+]i signals in 4-HNE treated Müller cells (Figs. 6D and 6E). Our results suggest that 4-HNE can facilitate TRPV4 channel activity in Müller cells and that this effect can be prevented by VEGF-B.

DISCUSSION

In this study, we demonstrate for the first time that Müller cells produce more VEGF-B than VEGF-A, -C, -D, or Pigf. Blocking VEGF-B did not alter Müller cell viability or expression of key functional proteins, indicating that VEGF-B might not be required for normal Müller cell function. These findings concur with a previous report that was unable to identify any obvious function for VEGF-B under physiologic conditions.53 However, it is also possible that VEGF-B may support Müller cells through an autocrine pathway in normal physiologic conditions as the levels of VEGF-B in Müller cell lysates were significantly higher than those in the supernatants. The autocrine support of VEGF-B to Müller cells may be supplemented significantly by the paracrine pathway in stress conditions, evidenced by the protective effects of recombinant VEGF-B and detrimental effects of blocking antibodies. Nevertheless, our results suggest that VEGF-B plays an important protective role in Müller cells under hypoxic or oxidative conditions.

Oxidative stress and hypoxia are known to be involved in several retinal diseases such as glaucoma, AMD, diabetic retinopathy, and retinal detachment to name but a few.1–3,44–46 Müller cells are one of the key responding cells of the retina to oxidative stress and hypoxia.1–3,46 Under hypoxic conditions, such as retinal detachment, the expression of GS was decreased and the TRPV4 activity was increased.44 Similarly, oxidative stress and hypoxia are implicated in the development of diabetic retinopathy.47,48 In this condition, an impaired Müller cell GS and dysregulation K+ and water transportation have been reported; ultimately, this process may lead to glutamate excitotoxicity, Müller cell swelling, and degeneration of the inner retina.2,47,48 Diabetic retinopathy also increases the production of neurotrophic factors such as VEGF-A, and was found to be one of the few pathologies that affects Müller cell viability.1,2

In this study, VEGF-B production in Müller cells was increased under both oxidative stress and hypoxic conditions. Blockade of VEGF-B or its cognate receptors, VEGFR1 and NRP1, compromised Müller cell viability in hypoxic and oxidative conditions most likely through induction of caspase-3 activation. Furthermore, treatment with recombi-
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Figure 6. Effect of VEGF-B on TRPV4 expression and functionality under oxidative stress. (A–C) QMMuC-1 cells were treated with 4-HNE (10 μM) with or without recombinant VEGF-B (100 ng/mL) for 24 hours. The expression of TRPV4 was examined by immunofluorescence staining (A, B) and Western Blot (C). (A) Representative image of TRPV4 (green) and phalloidin (red) in untreated control, 4-HNE treated and 4-HNE + recombinant VEGF-B (100 ng/mL) treated QMMuC-1 cells. Arrow in control indicates cluster aggregation of TRPV4. Arrows in 4-HNE-treated group indicates contracted cell body. (B) Quantitative analysis of different types of TRPV4-expressing QMMuC-1 cells (A, spread cell with evenly distributed TRPV4; B, spread cell with aggregated or uneven TRPV4 expression; C, contracted cell with aggregated TRPV4 expression). ****P < 0.001; n = 4 wells per experiment. Two-way ANOVA followed by Bonferroni post hoc test. (C) Western Blot of TRPV4 expression on 4-HNE–treated QMMuC-1 cells and its representative plots (D). Intracellular Ca2+ levels were measured using the FlexStation. The y-axis was the 340 nm/380 nm fluorescence ratio (ΔR340 nm/380 nm). (D and E) TRPV4 agonist GSK101 (100 nM) alone did not elicit a measurable calcium response in QMMuC-1 cells. A 24 hours 4-HNE (10 μM) treatment significantly increased the GSK101-evoked response, which was fully blocked by the TRPV4 antagonist HC06 (10 μM) and by recombinant VEGF-B (100 ng/mL). Mean ± SD, n = 4–8 ****P < 0.001; one-way ANOVA followed by Newmann-Keuls post hoc test. All experiments were performed at least twice.
nant VEGF-B attenuated 4-HNE–induced Müller cell death. This finding is in line with previous observations that showed that the survival of motor neurons, peripheral neurons, vascular cells, and retinal ganglion cells was promoted by VEGF-B.

Müller cells respond to retinal stress by initiating reactive gliosis, whereby they express high levels of gliosis markers such as GFAP and vimentin. Acute and/or sustained gliosis hinders retinal function via inflammatory cytokine induction, which accelerates disease progression. Interestingly, VEGF-B did not affect Müller cell gliosis under hypoxic or oxidative stress conditions. It did, however, regulate other Müller cell responses under pathologic conditions.

One vital function of Müller cells is glutamate transport and recycling in the retina. GS regulates extracellular glutamate levels by converting internalized glutamate molecules into nontoxic glutamine. Expression of GS is decreased in retinal diseases, leading to glutamate accumulation and toxicity. We found that hypoxia decreased GS expression in Müller cells and that this effect was completely prevented by recombinant VEGF-B, but not VEGF-A.

Müller cells also maintain K+ and water homeostasis, via inward rectifying potassium channels Kir4.1 and the water channel AQP4. Dysregulation in K+ and water homeostasis has been linked with a myriad of retinal diseases, such as diabetic retinopathy, retinal detachment, and diabetic macular edema. Kir4.1, AQP4 dysfunction can result in K+ dysregulation, impaired synaptic activity, Müller cell osmotic swelling, and retinal excitotoxicity and neurodegeneration. We found that VEGF-B blockade downregulated the expression of Kir4.1 and AQP4 when Müller cells were subjected to hypoxia and oxidative stress. Therefore, therapies like aflibercept, which targets different VEGF family members in the retina, could be problematic as they may exacerbate ongoing retinal disease via inhibition of the essential potassium and water homeostasis mechanisms.

TRPV4, member of the TRP channels superfamily, can function as an osmosensor and/or mechanosensor. TRPV4 colocalizes with AQP4 in the end feet of Müller glial cells. Activation of TRPV4 in swollen Müller cells has been reported to contribute significantly to the development of different retinal diseases, such as retinal detachment, glaucomatous retinopathy, and photoreceptor death. In this study, although 4-HNE treatment did not alter TRPV4 expression in Müller cells, it significantly disrupted the distribution of actin filaments and altered TRPV4 localization patterns. This finding is in line with a previous report on the interaction between TRPV4 and the microtubule cytoskeleton. In this study, 4-HNE facilitated TRPV4–dependent Ca2+ influx in Müller cells and this effect was blocked by VEGF-B. The restorative effect of VEGF-B on TRPV4 responses may occur via normalization of the cytoskeleton.

Owing to their prominent role in retinal homeostasis, Müller cells are an important therapeutic target in pathologies. We have demonstrated that VEGF-B is a potent gliotropic factor for Müller cells under pathologic conditions, and that the neutralization of this factor may exacerbate diseases processes. Because VEGF-B does not have a significant angiogenic effect, and its overexpression does not elicit any adverse effects, VEGF-B is, therefore, a promising protective molecule and may be a valuable therapy for oxidative stress- and hypoxia-related retinal diseases.

References

15. Le YZ. VEGF production and signaling in Müller glia are critical to modulating vascular function and neuronal integrity in diabetic retinopathy and hypoxic retinal vaso...
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