



VASCULAR BIOLOGY, ATHEROSCLEROSIS, AND ENDOTHELIUM BIOLOGY

Deletion of Endothelial Transforming Growth Factor— β Signaling Leads to Choroidal Neovascularization



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The molecular pathogenesis of choroidal neovascularization (CNV), an angiogenic process that critically contributes to vision loss in age-related macular degeneration, is unclear. Herein, we analyzed the role of transforming growth factor (TGF)- β signaling for CNV formation by generating a series of mutant mouse models with induced conditional deletion of TGF- β signaling in the entire eye, the retinal pigment epithelium (RPE), or the vascular endothelium. Deletion of TGF- β signaling in the eye caused CNV, irrespectively if it was ablated in newborn or 3-week-old mice. Areas of CNV showed photoreceptor degeneration, multilayered RPE, basal lamina deposits, and accumulations of monocytes/macrophages. The changes progressed, leading to marked structural and functional alterations of the retina. Although the specific deletion of TGF- β signaling in the RPE caused no obvious changes, specific deletion in vascular endothelial cells caused CNV and a phenotype similar to that observed after the deletion in the entire eye. We conclude that impairment of TGF- β signaling in the vascular endothelium of the eye is sufficient to trigger CNV formation. Our findings highlight the importance of TGF- β signaling as a key player in the development of ocular neovascularization and indicate a fundamental role of TGF- β signaling in the pathogenesis of age-related macular degeneration. (*Am J Pathol* 2017, 187: 2570–2589; <http://dx.doi.org/10.1016/j.ajpath.2017.06.018>)

The therapeutic options for treating age-related macular degeneration (AMD), the leading cause of vision loss and blindness in industrialized countries,^{1–3} are limited.^{4,5} AMD is a complex disease⁶ whose molecular pathogenesis is not well understood.^{4,5} Vision loss in AMD is caused by choroidal neovascularization (CNV) or geographic atrophy.^{4,5,7} In geographic atrophy (late dry AMD), the progressive atrophy of the retinal pigment epithelium (RPE) and of the choriocapillaris are likely causes of photoreceptor degeneration.⁵ Angiogenic processes that cause immature choroidal vessels to break in and through the RPE into the subretinal space characterize CNV.⁸ The resulting blood and plasma leakage causes fibrous scarring of the retina (wet AMD). The pathogenic mechanisms underlying the onset of CNV in AMD are unclear. Dysfunction of the RPE and a dysregulation of proangiogenic and immune-stimulating molecular factors in the region of the retinal/choroidal interface are considered as contributing factors.^{4,7,9,10}

The choroidal vasculature forms the choriocapillaris, a vascular bed of highly anastomosed capillaries that is essential for nutrition and oxygen supply of both photoreceptors and RPE.¹¹ The capillaries have a fenestrated endothelial layer and are highly permeable,¹² a substantial difference from retinal capillaries. There is evidence from several studies of genetically engineered mouse models indicating that RPE-derived vascular endothelial growth factor (VEGF) is essential for choriocapillaris maintenance. VEGF is expressed in high amounts by RPE cells during embryonic development and in adult life.¹³ If VEGF expression is compromised, formation of the choriocapillaris in development or its maintenance in adult eyes is severely impaired, leading to choriocapillaris ablation.^{14–18}

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Other growth factors that are secreted by RPE cells in high amounts are transforming growth factor (TGF)- β 1 and TGF- β 2,¹⁹ causing a concentration of both factors in the RPE/choroid complex that is >10-fold higher than in the retina.²⁰ A recent study showed that a conditional ocular deletion of TGF- β signaling results in pronounced structural changes of retinal capillaries, including the formation of abundant microaneurysms, leaky capillaries, and retinal hemorrhages. Overall, a phenotype resulted similar to that of diabetic retinopathy in humans.²¹

Because TGF- β s are present at high concentrations in the region of the RPE/choroid interface, we wondered if TGF- β signaling might not only be required for stability of retinal capillaries, but also for that of the choriocapillaris. To this end, we targeted the TGF- β type II receptor (T β RII), which is essential for TGF- β signaling, and generated a series of mutant mouse strains with an induced conditional deletion in the entire eye, the RPE, or vascular endothelial cells. Herein, we provide evidence that impairment of TGF- β signaling in the vascular endothelium of the eye is sufficient for the development of CNV. Our findings highlight the importance of the TGF- β signaling pathway as a key player in the development of ocular neovascularization and indicate a fundamental role of TGF- β signaling in the pathogenesis of AMD.

Materials and Methods

Mice

All experiments were performed in mice of either sex that were tested negatively for the *Rd8* mutation.²² Mice with two floxed alleles of *Tgfb2* (*Tgfb2*^{fl/fl})²³ were crossed with heterozygous *CAGGCre-ER*,²⁴ *VMD2*^{rtTA-Cre},²⁵ or *VECad-Cre-ER*^{T226} mice. Resulting *Tgfb2*^{fl/fl}; *CAGGCre-ER*, *Tgfb2*^{fl/fl}; *VMD2*^{rtTA-Cre}, or *Tgfb2*^{fl/fl}; *VECad-Cre-ER*^{T2} mice were used as experimental mice. For simplicity, *Tgfb2*^{fl/fl}; *CAGGCre-ER* mice will be referred to as *Tgfb2*^{eye}, *Tgfb2*^{fl/fl}; *VMD2*^{rtTA-Cre} mice will be referred to as *Tgfb2*^{ARPE}, and *Tgfb2*^{fl/fl}; *VECad-Cre-ER*^{T2} mice will be referred to as *Tgfb2*^{EC}. *Tgfb2*^{fl/fl} littermates with two unrecombined *Tgfb2* alleles served and are referred to as controls. Genetic backgrounds were 129SV (*Tgfb2*^{fl/fl}, *VMD2*^{rtTA-Cre}, and *VECad-Cre-ER*^{T2}) and C57/Bl6 (*CAGGCre-ER*). All mice were reared in a light/dark cycle of 12 hours (lights on at 7 AM). Genotypes were identified by isolating genomic DNA from tail biopsy specimens, and testing for transgenic Cre sequences and floxed *Tgfb2* sequences, as described previously.^{21,27}

Induction of Cre Recombinase

Cre recombinase in *CAGGCre-ER* and *VECad-Cre-ER*^{T2} mice is tamoxifen dependent. In *CAGGCre-ER* mice, the *Cre-ER* fusion protein is ubiquitously expressed. Cre recombinase is restricted to the cytoplasm and will only

access the nucleus after binding to tamoxifen.^{24,28} *VECad-Cre-ER*^{T2} mice carry the Cre-ER^{T2} expression cassette under regulatory control of the mouse vascular endothelial–cadherin gene promoter region that specifically drives gene expression in the vascular endothelium.²⁶ To activate Cre recombinase in *Tgfb2*^{eye} and *Tgfb2*^{EC} mice, the conditional knockout animals and their respective control littermates were equally treated with tamoxifen containing eye drops from postnatal day (P) 4 to P8, as described previously.²⁹ A second stock of *Tgfb2*^{eye} and control mice was treated from P21 to P25, a time when the development of the retinal vasculature is complete. For simplicity, *Tgfb2*^{eye} mice that were tamoxifen treated from P4 to P8 will be referred to as early induced, and *Tgfb2*^{eye} mice that were tamoxifen treated from P21 to P25 will be referred to as late induced. Cre recombinase in *VMD2*^{rtTA-Cre}; *Tgfb2*^{fl/fl} is doxycycline dependent and under the control of the RPE-specific bestrophin promoter.²⁵ Doxycycline (AppliChem, Darmstadt, Germany) was diluted in phosphate-buffered saline (PBS) to a final concentration of 0.1 g/mL, and doxycycline eye drops (10 μ L) were pipetted onto the eyes of *Tgfb2*^{ARPE} mice and their control littermates from P21 to P25 three times per day. A detailed list of the individual treatment time points and subsequently performed analyses of the eyes is presented in Table 1.

mT/mG and *R26R* Reporter Mice

The efficiency of the Cre recombinase activation in *VECad-Cre-ER*^{T2} mice was confirmed using *mT/mG* reporter mice,³⁰ which express a membrane-targeted green fluorescent protein after Cre-mediated excision. Heterozygous *VECad-Cre-ER*^{T2} mice were crossed with homozygous *mT/mG* mice. The resulting offspring (*VECad-Cre-ER*^{T2+/-}; *mT/mG*^{+/-} and *wildtype*; *mT/mG*^{+/-}) were treated with tamoxifen eye drops from P4 to P8, according to previously published protocols.^{21,29} Mice were euthanized at P10. After enucleation, the eyes were fixed in 4% paraformaldehyde (PFA) for 4 hours, washed extensively in phosphate buffer (PB; 0.1 mol/L, pH 7.4), incubated in 10%, 20%, and 30% sucrose overnight, and shock frozen in tissue-mounting medium (O.C.T. Compound; DiaTec, Bamberg, Germany). Frozen sections were washed three times in PB (5 minutes each), and cell nuclei were counterstained with DAPI (Vectashield; Vector Laboratories, Burlington, CA) 1:10 diluted in fluorescent mounting medium (Serva, Heidelberg, Germany). Successful activation of the Cre recombinase in *CAGGCre-ER* mice was confirmed using Rosa26 reporter (*R26R*) reporter mice.³¹ Heterozygous *CAGGCre-ER* mice were crossed with homozygous *R26R* mice. The resulting offspring were treated with tamoxifen eye drops from P21 to P25, according to previously published protocols.^{27,29} Mice were euthanized at the age of 4 weeks, and the eyes were enucleated and processed for LacZ staining, which was performed as described previously.^{29,32}

Table 1 Treatment Time Points and Performed Analyses

Mouse model	Treatment	Performed analyses	Age
<i>Tgfr2^{Δeye}</i> and controls	Tamoxifen P4-P8	Morphology	2 and 3 months
		FITC-dextran	6 weeks
		3D imaging	6 weeks
		Electron microscopy	2.5 months
		RNA analyses	6 weeks
		Immunohistochemistry	4 and 6 weeks
<i>Tgfr2^{Δeye}</i> and controls	Tamoxifen P21-P25	Morphology	3 and 6 months
		FITC-dextran	2 months
		Angiography	3 and 6 months
		ERG	3 and 6 months
		Electron microscopy	3 months
		RNA analyses	5 weeks
<i>Tgfr2^{ΔEC}</i> and controls	Tamoxifen P4-P8	Immunohistochemistry	3 months
		Morphology	6 weeks
		FITC-dextran	6 weeks
		Electron microscopy	4 and 6 weeks
<i>Tgfr2^{ΔRPE}</i> and controls	Doxycyclin P21-P25	Immunohistochemistry	4 and 6 weeks
		Morphology	6 months
		Angiography	6 months

3D, three dimensional; ERG, electroretinography; FITC, fluorescein isothiocyanate; P, postnatal day.

Paraffin sections were analyzed by light microscopy (Carl Zeiss, Jena, Germany).

Tgfr2 Deletion PCR

Tgfr2 deletion PCR screens its successful genomic deletion after Cre-mediated recombination. DNA samples of the sensory retina and the choroid (including cells of the RPE) served as templates. Primer pairs and protocols are described elsewhere.^{29,33} Actin was used as the loading control.

Morphology and Microscopy

Eyes were enucleated, fixed for 24 hours in 2% PFA/2.5% glutaraldehyde,³⁴ and embedded in Epon (Serva), as described elsewhere.^{35,36} Semithin meridional sections (1 μm thick) were cut through the eyes and stained, according to the method of Richardson et al.³⁷ The sections were analyzed on an Axio Imager Z1 microscope (Carl Zeiss) using Axiovision software version 4.8 (Carl Zeiss). Ultrathin sections were processed according to protocols published previously,^{38,39} stained with uranyl acetate and lead citrate, and analyzed on a transmission electron microscope (Libra; Carl Zeiss).

Immunohistochemistry

Before TβRII, collagen IV, and VEGF-A staining, eyes were fixed for 4 hours in 4% PFA, washed extensively in PB (0.1 mol/L, pH 7.4), and embedded in paraffin, according to standard protocols. Paraffin sections (6 μm thick) were deparaffinized and washed in water. For detection of

TβRII, sections were treated with boiling citrate buffer (1 × 10 minutes, pH 6), washed again in water, and incubated in PB. For detection of collagen IV and VEGF-A, sections were pretreated with 0.05 mol/L Tris-HCl (5 minutes) and covered with Proteinase K [100 μL of Proteinase K in 57 mL of Tris-HCl (0.05 mol/L), 5 minutes], washed in water, incubated in 2N HCl (20 minutes), and washed again in water. Sections were incubated in PB for 5 minutes and then blocked with 2% bovine serum albumin, 0.2% cold water fish gelatine, and 0.1% Triton X (TβRII), 2% bovine serum albumin and 0.1% Triton X (collagen IV), or 5% nonfat dry milk (VEGF-A) at room temperature for 60 minutes. Primary antibodies (Table 2) were diluted in a 1:10 dilution of blocking solution in PB. Before F4/80 and plasmalemma vesicle-associated protein (PLVAP) staining, eyes were fixed for 4 hours in 4% PFA, washed extensively in PB, incubated in 10%, 20%, and 30% sucrose/PBS overnight at 4°C, and shock frozen in tissue-mounting medium. For immunohistochemistry, frozen sections were washed three times in PB or PBS (F4/80) for 5 minutes each and blocked at room temperature with 2% bovine serum albumin in PB for 45 minutes (F4/80: 1% nonfat dry milk and 0.01% Tween in PB). Primary antibodies (Table 2) were diluted in a 1:10 dilution of blocking solution in PB (F4/80: 2% bovine serum albumin, 0.02% NaN₃, and 0.01% Triton X in PBS), and incubated at 4°C overnight. After three washes in PB/PBS (5 minutes each), biotinylated antibodies were applied for 1 hour and diluted in a 1:10 dilution of the blocking solution. Then, appropriate secondary antibodies (Table 2), diluted in a 1:10 dilution of the blocking solution, were applied on the sections for 1 hour. Sections were washed again three times in PB/PBS, and cell nuclei were counterstained with DAPI (Vectashield) diluted (1:10) in

Table 2 Antibodies Used for Immunohistochemistry

Primary antibody (manufacturer)	Secondary antibody (manufacturer)
T β RII-L21 (Santa Cruz Biotechnology, Dallas, TX), 1:20	Anti-rabbit, biotinylated (Vector Laboratories, Burlington, CA), 1:500; streptavidin Alexa 488 (Invitrogen, Carlsbad, CA), 1:1000
Collagen IV (Rockland Immunochemicals Inc., Limerick, PA), 1:100	Anti-rabbit Cy-3 conjugated (Jackson Immuno Research Labs, West Grove, PA), 1:2000
VEGF-A (R&D Systems, Minneapolis, MN), 1:50	Anti-goat, biotinylated (Vector Laboratories), 1:500; streptavidin Alexa 546 (Invitrogen), 1:1000
F4/80 (Acris Antibodies GmbH, Herford, Germany), 1:600	Anti-rat Cy-3 conjugated (Jackson Immuno Research Labs), 1:2000, in PBS
PLVAP (Santa Cruz Biotechnology), 1:50	Anti-rat Cy-3 conjugated (Jackson Immuno Research Labs), 1:2000

In all cases, fixation was performed with 4% paraformaldehyde.

PBS, phosphate-buffered saline; PLVAP, plasmalemma vesicle-associated protein; T β RII, transforming growth factor- β type II receptor.

fluorescent mounting medium (Serva). Sections were investigated on an Axio Imager Z1 fluorescent microscope using appropriate Axiovision software.

Quantification of F4/80-Positive Cells on Immunohistochemical Sections

Experimenters (A.S. and B.M.B.) were blinded regarding the genotype. Antibody incubation times were strictly monitored, and microscope settings (eg, the illumination time and intensity) were identical between individual sections. Only sections in midsagittal orientation were analyzed that stretched through the optic nerve to ensure comparable situations between the individual sections. F4/80-positive cells were counted per hemisphere in the retina and in the choroid/choriocapillaris.

Dextran Perfusion

Before dextran perfusion, mice were deeply anesthetized with ketamine [120 mg/kg body weight (bw)] and xylazine (8 mg/kg bw). Afterward, mice were perfused through the left ventricle with 1 mL of PBS containing 50 mg fluorescein isothiocyanate (FITC)-dextran (mol. wt., 2000 kDa; TdB Consultancy, Uppsala, Sweden). The eyes were enucleated and fixed in 4% PFA for 2 hours (flat mounts)/4 hours (sections) and washed in PB. Eyes were cut into sections, placed on glass slides, and counterstained with DAPI (Vectashield) 1:10 diluted in fluorescent mounting medium. FITC-dextran perfused sections were investigated on an Axio Imager Z1 fluorescent microscope.

Quantification of CNV

After dextran perfusion, flat mounts of the posterior eye segment (containing the retina, RPE, choroid, and sclera) were dissected. Similar to the preparation of retinal flat mounts, the posterior eye segment was flat mounted on glass slides using fluorescent mounting medium. The entire flat mount was analyzed using an Axio Imager Z1 fluorescent microscope. Focusing on the retinal pigment epithelium allowed the quantification of the CNV penetrating through

the RPE. To visualize the CNV, Z-stacks were generated that ranged from the superficial retinal plexus to the optical barrier of the pigmented RPE.

ERG Data

Mice were dark adapted for at least 12 hours before the experiments and anesthetized by s.c. injection of ketamine (65 mg/kg bw) and xylazine (13 mg/kg bw). Pupils were dilated with tropicamide eye drops (Mydriaticum Stulln; Pharma Stulln GmbH, Stulln, Germany). Silver needle electrodes served as reference (forehead) and ground (tail), and gold wire ring electrodes served as active electrodes. Electroretinograms (ERGs) were recorded with a Ganzfeld bowl (Ganzfeld QC450 SCX; Roland Consult, Brandenburg, Germany) and an amplifier/recording unit (RETI-Port; Roland Consult), band pass filtered (1 to 300 Hz), and averaged. Single-flash scotopic (dark-adapted) responses to a series of 10 light-emitting diode flash intensities (range, -3.5 to 1.0 log cd \cdot second/m²) were recorded. After 10 minutes of adaptation to a white background illumination (20 cd/m²), single-flash photopic (light-adapted) responses to three xenon-flash intensities (1, 1.5, and 2 log cd \cdot second/m²) were recorded. All analyses and plotting were performed with R version 3.2.1 (The R Foundation for Statistical Computing, Vienna, Austria) and ggplot2 version 2.1.0.⁴⁰

Fundus Imaging and Angiography

Imaging of the retinal vasculature was performed with a commercially available imaging system (Micron III; Phoenix Research Laboratories, Pleasanton, CA). Light source and imaging path filters (low and high pass at 500 nm) were used for fluorescein angiography. Mice were anesthetized by s.c. injection of ketamine (65 mg/kg bw) and xylazine (13 mg/kg bw), and their pupils were dilated with tropicamide eye drops before image acquisition. Fluorescein angiography was performed with s.c. injection of 75 mg/kg bw fluorescein-sodium (Alcon, Hünenberg, Switzerland).

Table 3 Primers Used for Real-Time PCR Amplification

Gene	Sequence forward	Sequence reverse
<i>GAPDH</i>	5'-TGTCCGTCGTGGATCTGAC-3'	5'-CCTGCTTACCACCTTCTTG-3'
<i>GNB2L</i>	5'-TCTGCAAGTACACGGTCCAG-3'	5'-ACGATGATAGGTTGCTGCT-3'
<i>RPL32</i>	5'-GCTGCCATCTGTTTTACGG-3'	5'-TGACTGGTGCCTGATGAACT-3'
<i>Tgfbβ2</i>	5'-AGAAGCCGCATGAAGTCTG-3'	5'-GGCAAACCGTCTCCAGAGTA-3'
<i>Ccl2</i>	5'-CATCCACGTGTTGGCTCA-3'	5'-GATCATCTTGTGTTGGAATGAGT-3'
<i>Cd68</i>	5'-CTCTCTAAGGCTACAGGCTGCT-3'	5'-TCACGGTTGCAAGAGAAACA-3'
<i>Gfap</i>	5'-TCGAGATCGCCACCTACAG-3'	5'-GTCTGTACAGGAATGGTGATGC-3'
<i>Il-6</i>	5'-GCTACCAAACCTGGATATAATCAGGA-3'	5'-CCAGGTAGCTATGGTACTCCAGAA-3'
<i>iNos</i>	5'-GGGCTGTCACGGAGATCA-3'	5'-CCATGATGGTCACATTCTGC-3'
<i>Tnf-α</i>	5'-TCTTCTCATTCCTGCTTGTGG-3'	5'-GGTCTGGGCCATAGAACTGA-3'
<i>Vegf-a-120</i>	5'-AAAGCCAGCACATAGGAGAG-3'	5'-GGCTTGTCACATTTTCTGG-3'
<i>Vegf-a-164</i>	5'-GAACAAAGCCAGAAAATCACTGTG-3'	5'-CGAGTCTGTGTTTTTGCAGGAAC-3'
<i>Fgf2</i>	5'-CGGCTCTACTGCAAGAACG-3'	5'-TGCTTGGAGTTGTAGTTGACG-3'

RNA Analysis

Total RNA from retinas was extracted with TriFast (Peqlab Biotechnologie GmbH, Erlangen, Germany), and first-strand cDNA synthesis was performed with the iScript cDNA Synthesis Kit (Bio-Rad Laboratories, Inc., Hercules, CA), according to the manufacturer's guidelines. Real-time quantitative RT-PCR analyses were performed with the iQ5 Realtime PCR Detection System (Bio-Rad Laboratories, Inc.). The temperature profile was denaturation at 95°C for 10 seconds and annealing and extension at 60°C for 40 seconds for 40 cycles. All primer pairs were purchased from Invitrogen (Carlsbad, CA) and extended over exon-intron boundaries, except for *Gapdh*. Sequences of primer pairs are shown in Table 3. RNA that was not reverse transcribed served as a negative control for real-time RT-PCR. Before relative quantification, mRNAs from three different potential housekeeping genes were tested: glyceraldehyde 3-phosphate dehydrogenase, guanine nucleotide binding protein, and ribosomal protein L32. Statistical analyses were performed to confirm that none of the housekeeping genes were regulated and then the geometric mean of the crossing thresholds of all three housekeeping genes was used for relative quantification. Quantification was performed using BioRad iQ5 Standard-Edition software version 2.1 (Bio-Rad Laboratories GmbH, Munich, Germany) and the $\Delta\Delta Ct$ method in Excel (Microsoft Corp., Redmond, WA).⁴¹

Deep Tissue Imaging by Three-Dimensional Light-Sheet Fluorescence Microscopy

Mice were deeply anesthetized with ketamine (120 mg/kg bw) and xylazine (8 mg/kg bw), and 130 μ L of *Tomato lectin* (Perkin Elmer, Inc., Waltham, MA) was slowly injected intravenously. Mice were euthanized 15 minutes after injection by cervical dislocation. The eyes were enucleated and fixed using Paxgene Tissue Containers (Qiagen, Hilden, Germany), according to the manufacturer's

recommendations, and thereafter underwent a chemical procedure of optical clearing, as previously described.⁴² Cleared transparent whole mouse eyes were imaged on a light-sheet fluorescence microscope (UltraMicroscope II; LaVision BioTec, Bielefeld, Germany), equipped with an sCMOS camera (Andor Neo, Concord, MA) and a 2 \times /0.5 numerical aperture objective lens (MVPLAPO 2 \times ; Olympus, Hamburg, Germany). The specimens were two sided, illuminated by a planar light sheet using a white light laser (SuperK EXTREME EXW-9; NKT Photonics, Birkerød, Denmark). To visualize the specific TLelectinSense 680 signals for vascularization, a bandpass filter set with an excitation range of 640/30 nm and an emission range of 690/50 nm was used in combination with an additional filter set (excitation, 531/40 nm; emission, 593/40 nm) for detection of autofluorescence (morphology). By moving the specimen chamber vertically stepwise (step size, 4 μ m) through the laser light sheet, optical sections were obtained. To ensure standardized imaging regions for each eye, the scan always covered 600 μ m above to 600 μ m below the optical nerve. Maximum intensity projections were performed by InspectorPro software version 5.0 (LaVision BioTec, Bielefeld, Germany).

Statistical Analysis

All results are expressed as means \pm SEM. Comparisons between the mean variables of two groups were made by a two-tailed *t*-test. Significance of the ERG analyses was evaluated using a one-way analysis of variance test, followed by a Tukey honest significant difference post hoc test. $P \leq 0.05$ was considered statistically significant.

Study Approval

All procedures conformed to the tenets of the NIH's *Guide for the Care and Use of Laboratory Animals*,⁴³ European Union Directive 2010/63/E, and institutional guidelines;

all procedures were approved by the local authority (Regierung der Oberpfalz, AZ 54-2532.1-44/12 and DMS 2532-2-85).

Results

Deletion of Ocular TGF- β Signaling in Mouse Pups Leads to Choroidal Neovascularization

To investigate the role of the TGF- β signaling pathway for the structure of retinal and choroidal vessels, the essential T β R2 receptor in the eyes was removed via tamoxifen-induced conditional deletion. To this end, *Tgfr2^{fl/fl};CAGGCre-ER* mice were generated and treated from P4 to P8 with tamoxifen eye drops. For simplicity, *Tgfr2^{fl/fl};CAGGCre-ER* mice that received tamoxifen by this protocol will be referred to as early-induced *Tgfr2^{Δeye}* mice, and tamoxifen-treated littermates with two unrecombined *Tgfr2^{fl/fl}* alleles are referred to as controls. As shown previously, the treatment results in a significant reduction of T β R2 and its mRNA in the eyes of early-induced *Tgfr2^{Δeye}* mice compared to controls.²¹ The previous study also showed that the deletion of ocular TGF- β signaling leads to the formation of abundant microaneurysms in retinal vessels and to a phenotype that largely mimics that of non-proliferative and proliferative diabetic retinopathy in humans.²¹ Consequently, the retina of 2- and 3-month-old early-induced *Tgfr2^{Δeye}* mice contained vascular microaneurysms in the inner nuclear layer, which were characterized by their large lumen and thick coat of endothelial cells (Figure 1, A–C). In contrast, no changes were seen in control mice (Figure 1, A and B). We now turned our attention to the choroidal/retinal interface that had not been analyzed previously. Herein, we frequently observed focal areas in which the outer segments of photoreceptors were completely missing and the thickness of the outer nuclear layer was substantially reduced (Figure 1, B and C). Accumulations of cells were present between photoreceptors and RPE. The cells frequently surrounded vascular openings filled with erythrocytes, strongly indicating their endothelial origin and an ongoing neoangiogenic process. The RPE was multilayered in those areas, and contained cystic and amorphous inclusions (Figure 1C). When investigating serial sections, we observed vessels beneath and above the RPE, concomitant with type 1 (sub-RPE, occult) and type 2 (subretinal, classic) CNV in human patients.

To characterize the origin of the newly formed vessels in the subretinal space (between the sensory retina and the RPE) of early-induced *Tgfr2^{Δeye}* mice, we perfused the mice with high-mol. wt. FITC-dextran for vascular labeling. Although the retinal and choroidal vasculature of control animals was essentially normal, distinct changes were observed in the subretinal space of early-induced *Tgfr2^{Δeye}* mice. Vessels were observed breaking from the choroid through the Bruch membrane (BM) across the RPE into the subretinal space (Figure 2A). FITC-dextran-perfused flat-mounted posterior eye segments (containing the retina,

RPE, choroid, and sclera) of 4- to 6-week-old animals had 2.86 ± 0.59 CNVs per eye ($n = 7$) (Supplemental Figure S1B). To confirm this observation by an independent method, three-dimensional imaging of lectin-perfused, optical-cleared, and transparent whole eyes was performed (Figure 2B). The choriocapillaris did not resolve into single capillaries but rather appeared as an intense fluorescent line covering the outer surface of the retina, a fact attributed to the immense density and blood flow of this capillary bed.⁴⁴ In control eyes, the three plexuses of the retinal vasculature could be completely visualized and were essentially normal. In contrast, in early-induced *Tgfr2^{Δeye}* mice, vessels that originated from the choriocapillaris entered the retina and continued into the retinal vascular bed, forming obvious anastomoses with retinal vessels (Figure 2B). We now characterized the vascular wall of the newly formed vessels by transmission electron microscopy. As expected, in control mice, the endothelium surrounding retinal vessels was continuous (Figure 2C), whereas that around the choriocapillaris was fenestrated. When the newly formed vessels in the subretinal space of early-induced *Tgfr2^{Δeye}* mice were analyzed, distinct fenestrations covered by a typical diaphragm were regularly observed (Figure 2C). To further support this finding, immunohistochemical staining was performed for PLVAP, a vascular protein, an intrinsic component of the diaphragm of vascular fenestrae, and a marker for fenestrated endothelia.^{45–48} As expected, in control and early-induced *Tgfr2^{Δeye}* animals, the continuous endothelium of the retinal vasculature did not show any PLVAP immunoreactivity (Supplemental Figure S1A), whereas the fenestrated endothelium of the choriocapillaris showed an intense PLVAP signal (Figure 2D). In early-induced *Tgfr2^{Δeye}* mice, intense PLVAP immunoreactivity was observed in the areas of neovascularization in the subretinal space (Figure 2D). Overall, our results strongly indicated that loss of TGF- β signaling in early-induced *Tgfr2^{Δeye}* mice caused CNV.

TGF- β Signaling Is Required to Prevent CNV in Late-Induced *Tgfr2^{Δeye}* Mice

Next, we wondered whether the formation of CNV after T β R2 deletion might be supported by the facts that retinal vascular development is not completed in mouse pups and that signaling processes that promote angiogenesis are continuously ongoing. To learn if TGF- β signaling is also important for maintenance of the adult choriocapillaris when retinal vascular development is completed, we deleted TGF- β signaling by an essentially comparable approach in 3-week-old mice with completely developed retinal vasculature. Mice were treated with tamoxifen eye drops from P21 to P25 and are further referred to as late-induced *Tgfr2^{Δeye}* mice. To confirm successful tamoxifen-induced recombination, we crossed the mice with Cre reporter mice carrying the *R26R* allele. Tamoxifen-treated eyes were stained for β -galactosidase, and an intense staining was seen

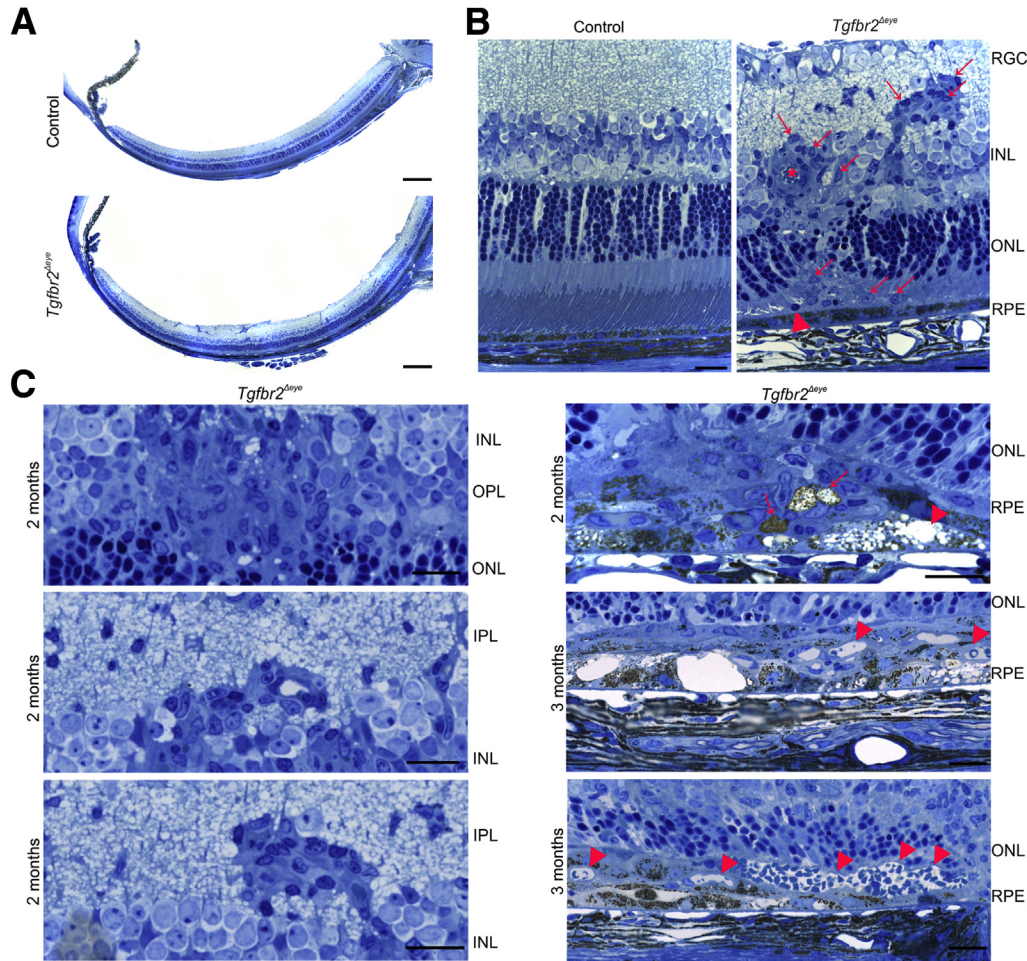


Figure 1 Early-induced deletion of transforming growth factor- β (TGF- β) signaling in the eye causes structural changes in the subretinal space. Stained semithin sections (1 μ m thick) of an early-induced *Tgfb2 Δ eye* mouse and its control littermate, according to the method of Richardson et al.³⁷ **A:** Retinal hemispheres of 2-month-old control and *Tgfb2 Δ eye* animals. **B:** Detailed magnification of the retina. The control mouse shows regular structure. In the retina of the *Tgfb2 Δ eye* mouse, a vessel (**asterisk**) filled with erythrocytes is seen, which disrupts the inner nuclear layer (INL) and outer nuclear layer (ONL) and is surrounded by an accumulation of cells that continue into the subretinal space (**arrows**). In the subretinal space, mononuclear cells (**arrowhead**) are present, whereas photoreceptor outer segments are degenerated and the choroid is thickened. **C, left panels:** In the inner retina of 2-month-old *Tgfb2 Δ eye* mice, accumulations of endothelial cells are seen in both the ONL/outer plexiform layer (OPL) and in the INL/inner plexiform layer (IPL). **C, right panels:** At the retinal/choroidal interface of 2- and 3-month-old *Tgfb2 Δ eye* mice, the retinal pigment epithelium (RPE) is multilayered (**arrows in top panel**) and contains cystic and amorphous inclusions (**arrowhead in top panel**). At the age of 3 months, erythrocyte-containing vessels fill the subretinal space (**arrowheads in middle and bottom panels**). Scale bars: 200 μ m (**A**); 20 μ m (**B** and **C**). RGC, retinal ganglion cell.

throughout the eye of *CAGGCre-ER;R26R* mice, whereas ocular tissues of *R26R* littermates, which did not carry the *CAGGCre-ER* transgene, were completely unstained (**Supplemental Figure S2A**). Furthermore, the significant deletion of retinal *Tgfb2* was confirmed by real-time RT-PCR analyses (**Supplemental Figure S2B**). Next, we analyzed the ocular morphology of late-induced *Tgfb2 Δ eye* mice and controls at the age of 3 (**Figure 3, A and B**) and 6 (**Figure 4, A and C**) months. Retinal structure was essentially normal in control animals (**Figure 3, A and B, and Figure 4, A and C**), and no obvious changes were observed in the inner retina of 3-month-old late-induced *Tgfb2 Δ eye* animals (**Figure 3B**). However, similar to our observations in early-induced *Tgfb2 Δ eye* animals, all of the late-induced

Tgfb2 Δ eye mice showed structural changes in the subretinal space that were essentially comparable to that observed in early-induced *Tgfb2 Δ eye* mice (**Figure 3, A and D, and Figure 4, A–C**). In focal areas in which photoreceptor outer segments were shortened or completely missing, vessels were observed in the subretinal space. By transmission electron microscopy, a fenestrated endothelial layer was identified surrounding the vessels (**Figure 5B**). The RPE in those areas was multilayered and frequently contained cystic and amorphous inclusions. In 6-month-old late-induced *Tgfb2 Δ eye* mice, the choroid was thickened and photoreceptor outer segments were completely missing (**Figure 4, A–C**). The RPE was reduced to a flat layer or formed focal accumulations in those areas in which subretinal vessels

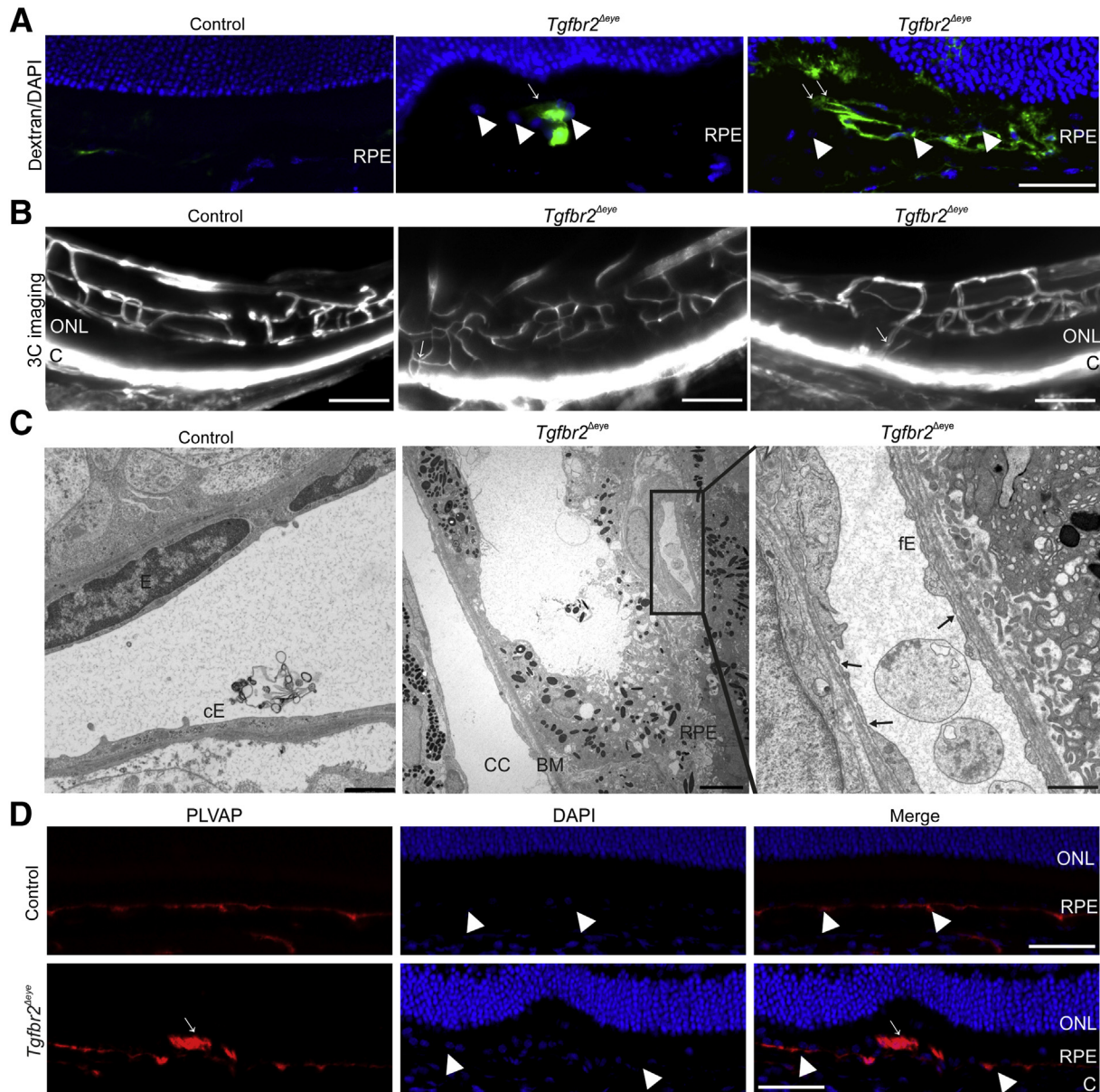


Figure 2 Subretinal neovascularization with fenestrated endothelium (fE) after deletion of transforming growth factor- β (TGF- β) signaling in the eyes of early-induced *Tgfb2 Δ eye* mice. **A:** Fluorescein isothiocyanate–dextran (green)—perfused retinal meridional section of a control and *Tgfb2 Δ eye* mouse at 6 weeks of age. **White arrows** point toward choroidal vessels breaking through the Bruch membrane (BM) and retinal pigment epithelium (RPE; nuclei of the RPE marked by **white arrowheads**) into the subretinal space. Nuclei are DAPI stained (blue). **B:** Light-sheet fluorescence microscopy of transparent eyes of 6-week-old lectin-injected *Tgfb2 Δ eye* mice and a control littermate. The control mouse shows an essentially regular arborized retinal vasculature. The *Tgfb2 Δ eye* mice have an irregular arrangement of the retinal plexus and form anastomoses between retinal and choroidal vessels (**arrows**). **C:** Transmission electron microscopy of an intraretinal vessel outlined with a continuous endothelium (cE) in a 2.5-month-old control animal. In contrast, the subretinal neovascularization in the *Tgfb2 Δ eye* littermate has a fE (**black arrows, right panel**). The **right panel** shows the **boxed region** of the **middle panel** in higher magnification. **D:** Immunoreactivity for plasmalemma vesicle-associated protein (PLVAP; red) in the retina at 4 weeks of age. The control and the *Tgfb2 Δ eye* animal show a thin, one-layered PLVAP signal in the choriocapillaris (CC; **white arrowheads**). In addition, the *Tgfb2 Δ eye* mouse displays PLVAP-positive signals (**white arrows**) in the RPE and subretinal space. Nuclei are DAPI stained (blue). Scale bars: 50 μ m (**A** and **D**); 120 μ m (**B**); 1000 nm (**C, left and right panels**); 5000 nm (**C, middle panel**). 3D, three dimensional; C, choroid; ONL, outer nuclear layer.

were present. *In vivo* imaging and fluorescein *in vivo* angiography of 3-month-old animals showed a regular fundus and retinal vasculature in both control and late-induced *Tgfb2 Δ eye* mice (Figure 3C). To investigate the

retinal vasculature in more detail, we additionally perfused 2-month-old, late-induced *Tgfb2 Δ eye* mice with high-mol. wt. FITC-dextran and analyzed the vasculature on meridional sections. The retinal vasculature had a regular

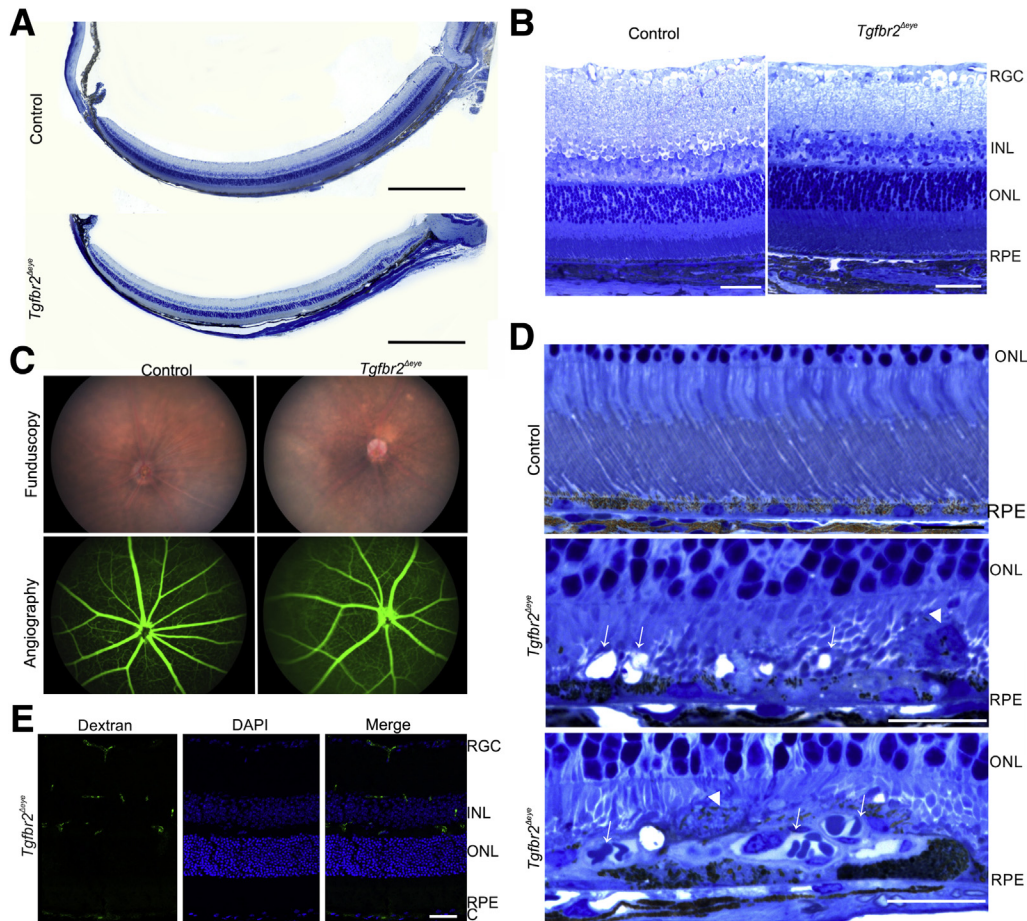


Figure 3 Subretinal neovascularization after deletion of transforming growth factor- β (TGF- β) signaling in the eyes of 3-month-old late-induced *Tgfb2* ^{Δ eye} mice. **A:** Stained semithin sections (1 μ m thick) of the retinal hemispheres of a 3-month-old, late-induced *Tgfb2* ^{Δ eye} mouse and its control littermate, according to the method of Richardson et al.³⁷ **B:** Higher magnification of the retinal sections. **C:** *In vivo* funduscopy and fluorescein angiography of a 3-month-old, late-induced *Tgfb2* ^{Δ eye} mouse and its control littermate. The retinal vasculature and the fundus appear essentially normal. Diameters of the image sections refer to 2 mm retina. **D:** Detailed magnification of stained semithin sections, according to the method of Richardson et al,³⁷ of the retinal/choroidal interface of 3-month-old *Tgfb2* ^{Δ eye} mice and a control littermate. The control mouse shows regular structure. *Tgfb2* ^{Δ eye} mice show focal areas in which the photoreceptor outer segments are shortened or completely missing, with vessels in the subretinal space. In these regions, the retinal pigment epithelium (RPE) is multilayered (**white arrowhead, middle and bottom panels**) and frequently contains cystic and amorphous inclusions (**white arrows, middle panel**). Mononuclear cells (**white arrowhead, middle panel**) are visible in the subretinal space. **E:** Fluorescein isothiocyanate-dextran (green)—perfused retinal meridional section of a 2-month-old *Tgfb2* ^{Δ eye} mouse. The three retinal vascular plexuses are present in their correct localization. Nuclei are DAPI stained (blue). Scale bars: 500 μ m (**A**); 50 μ m (**B** and **E**); 20 μ m (**D**). C, choroid; INL, inner nuclear layer; ONL, outer nuclear layer; RGC, retinal ganglion cell.

appearance, and the vessels were in their anatomically correct localization. Furthermore, we did not observe FITC-dextran or fluorescein leaking into the vitreous, the subretinal space, or the surrounding tissue (Figure 3, C and E). As we had observed a significant reduction of the pericyte marker neural/glia antigen 2 in the retina of early-induced *Tgfb2* ^{Δ eye} animals,²¹ we now focused on the coating of the choroidal vessels with neural/glia antigen 2-positive pericytes in 4-week-old early-induced and 6-week-old late-induced *Tgfb2* ^{Δ eye} animals. We did not observe a difference in pericyte coverage compared to controls (data not shown). When the eyes at the age of 6 months were analyzed by *in vivo* fundus imaging and fluorescein angiography, control mice showed no pathological changes, in neither the fundus nor the retinal vasculature (Figure 4D). In

contrast, in late-induced *Tgfb2* ^{Δ eye} mice, the fundus was hyperpigmented, indicating RPE proliferation or dedifferentiation; in some animals, the retina had detached (Figure 4D). In parallel to the structural changes, marked functional changes by ERG that reflected the observed gradual degeneration of the retina from 3 to 6 months of age in late-induced *Tgfb2* ^{Δ eye} mice were detected (Supplemental Figure S3, A–G). Although, in 3-month-old *Tgfb2* ^{Δ eye} mice, the scotopic responses showed a similar waveform but slightly lower amplitudes compared to control mice, the photopic waveform was unchanged in amplitude and implicit time. However, the difference was not statistically significant. In 6-month-old *Tgfb2* ^{Δ eye} mice, both the scotopic and photopic responses could not be distinguished from noise in almost all eyes.

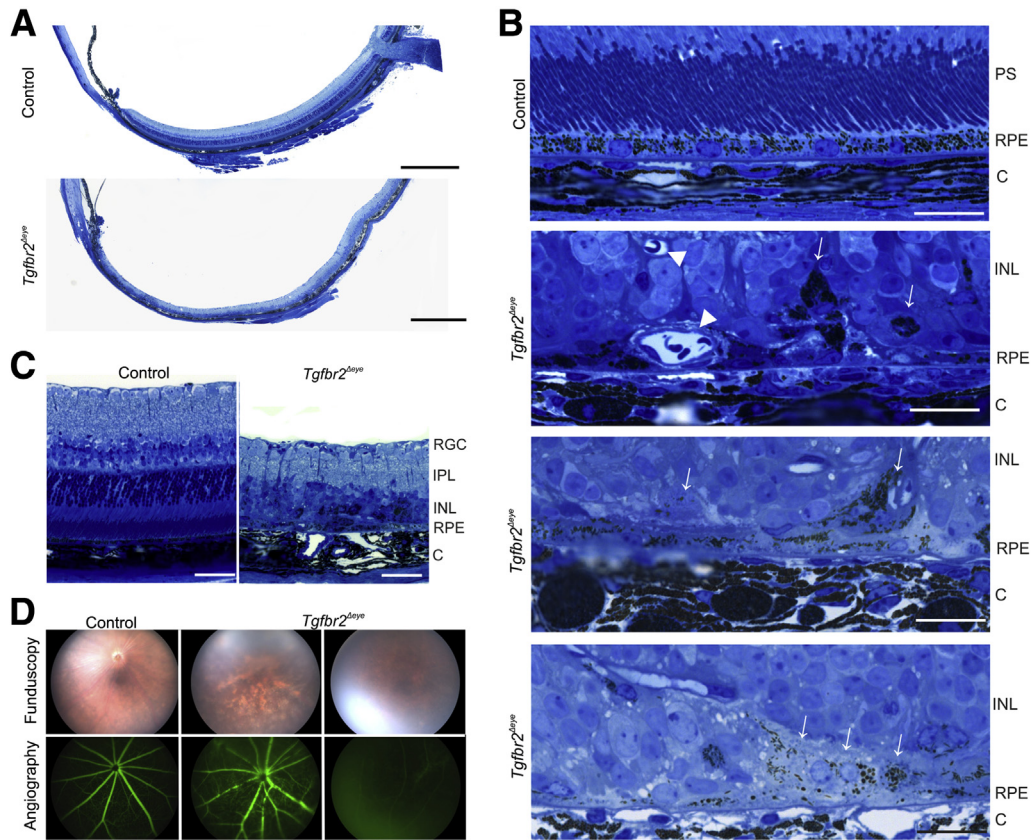


Figure 4 Subretinal neovascularization and retinal degeneration after deletion of transforming growth factor- β (TGF- β) signaling in the eyes of 6-month-old late-induced *Tgfb2 Δ eye* mice. **A:** Stained semithin sections (1 μ m thick) of the retinal hemispheres of 6-month-old animals, according to the method of Richardson et al.³⁷ The control animal shows regular retinal morphology. In contrast, in the *Tgfb2 Δ eye* littermate, photoreceptor outer segments are completely degenerated. **B:** Detailed magnification of stained semithin sections of the outer retina and the choroid (C) of 6-month-old *Tgfb2 Δ eye* mice and a control littermate, according to the method of Richardson et al.³⁷ The morphology of the control mouse is normal. In contrast, the *Tgfb2 Δ eye* mouse has erythrocyte-filled vessels in the retinal pigment epithelium (RPE; *Tgfb2 Δ eye* top panel, arrowheads), areas with a thickened (*Tgfb2 Δ eye* middle panel) and thinned (*Tgfb2 Δ eye* bottom panel) RPE, and an accumulation of pigmented cells in the sensory retina (*Tgfb2 Δ eye* top, middle, and bottom panels, arrows). **C:** Detailed magnification of the retina. In the *Tgfb2 Δ eye* mouse, the retina is degenerated, with a complete loss of photoreceptors. The RPE is disorganized, pigmented cells accumulate in the sensory retina, and the choroid is thickened. **D:** *In vivo* funduscopy and fluorescein angiography of the control animal are normal. *Tgfb2 Δ eye* mice show a hyperpigmented fundus (**middle panels**) or retinal detachment (**right panels**). Diameters of the image sections refer to 2 mm retina. Scale bars: 500 μ m (**A**); 20 μ m (**B**); 50 μ m (**C**). INL, inner nuclear layer; IPL, inner plexiform layer; PS, photoreceptor segment; RGC, retinal ganglion cell.

Expression of Angiogenic Factors and Immune Modulating Cytokines in *Tgfb2 Δ eye* Mice

The immunovascular axis, a cross talk of the RPE with immune and vascular cells, is considered a potent driver of CNV formation.⁴ To clarify the mechanisms behind CNV induced by T β RII deletion, we consequently analyzed the mRNA expression levels of immune-modulating cytokines and markers for the reactivity of Müller glia and microglia cells, respectively. Early-induced *Tgfb2 Δ eye* mice showed a significant increase in retinal mRNA expression levels of glial fibrillary acidic protein, Cd68, inducible nitric oxide synthase, Il-6, tumor necrosis factor- α , and monocyte chemoattractant protein-1 [chemokine (C-C motif) ligand 2] at the age of 6 weeks compared to control littermates (Figure 6A). However, in 3-month-old late-induced *Tgfb2 Δ eye* mice, the retinal mRNA expression levels were not significantly enhanced compared to controls; in fact,

glial fibrillary acidic protein was even significantly down-regulated (Figure 6B). Next, we used an antibody against F4/80 to label macrophages and microglia cells.^{49–51} The number of F4/80-positive cells was significantly ($P \leq 0.05$) increased in the choroid of early-induced *Tgfb2 Δ eye* animals (142.67 ± 17.13 , $n = 3$) when compared with controls (95.0 ± 8.40 , $n = 4$) (Figure 6C). The F4/80-positive cells were detected in the subretinal space in close association with the CNV (Figure 6C). Furthermore, the number of F4/80-positive cells in the retina was significantly increased ($P \leq 0.001$) in early-induced *Tgfb2 Δ eye* animals (145.75 ± 9.73 , $n = 4$) compared to controls (22.5 ± 1.71 , $n = 4$). Similarly, 3-month-old late-induced *Tgfb2 Δ eye* animals also presented a distinct accumulation of F4/80-positive cells in the choroid when compared to controls (Figure 6D). Subsequently, we focused on molecular factors that influence vascular proliferation. In our previously published study, we already showed that early-induced

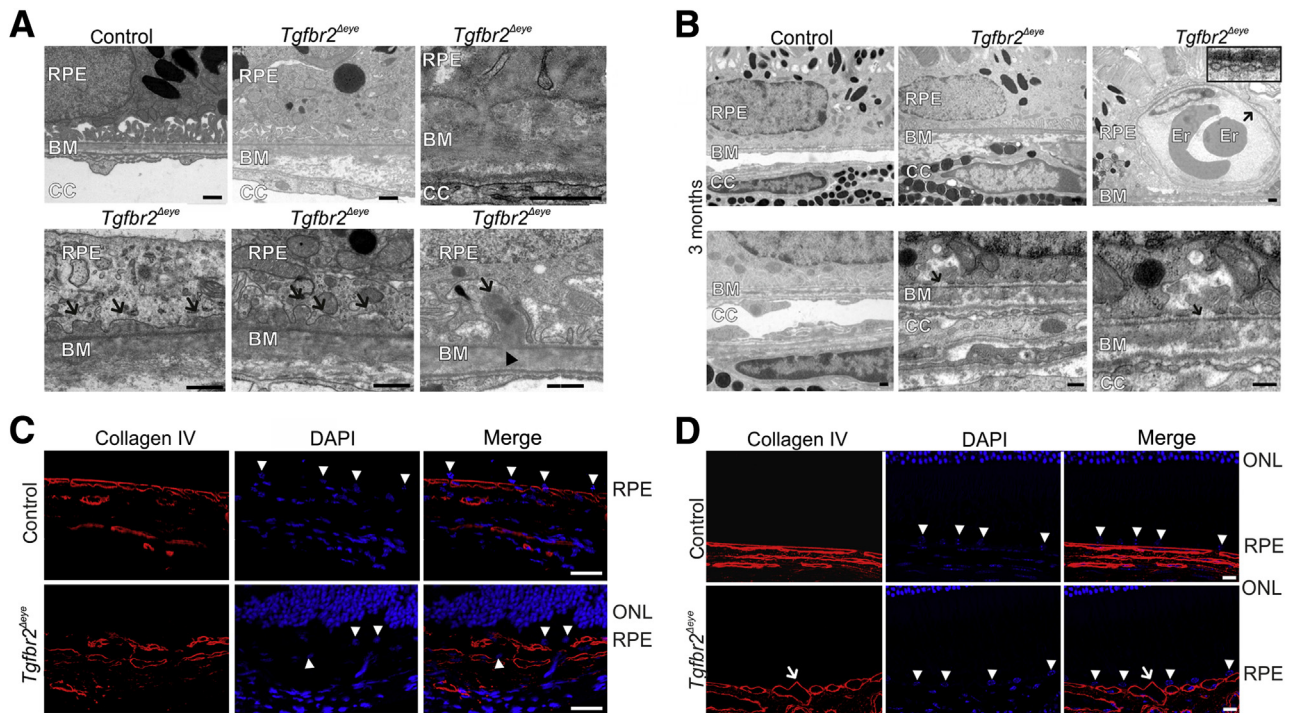


Figure 5 Structural changes of the Bruch membrane (BM) in early- and late-induced *Tgfb2^{Δeye}* mice. **A:** Transmission electron microscopy (TEM) of retinal pigment epithelium (RPE)/BM in early-induced 2.5-month-old mice. **Top panels:** The BM of the control is of regular structure, but thickened in *Tgfb2^{Δeye}* mice as a result of an accumulation of collagen fibers and fine fibrillar extracellular material between the basal lamina of the choriocapillaris (CC) and the elastic layer of the BM. In some areas (**right panels**), the RPE basal lamina is replaced by polymorphous electron-dense material that reaches from the elastic layer of the BM to the RPE basal infoldings. **Bottom panels:** In other RPE/BM areas of early-induced *Tgfb2^{Δeye}* mice, irregular nodules arise from the RPE basal lamina (**arrows**) and extend between the basal infoldings of the RPE. In places (**arrowhead, right panel**), the RPE basal lamina is interrupted where nodules arise. **B:** TEM of RPE/BM in 3-month-old late-induced *Tgfb2^{Δeye}* mice. RPE/BM are of regular structure in control mice, but thickened in *Tgfb2^{Δeye}* mice. Erythrocyte (Er)-filled vessels with fenestrated endothelium (**arrow**) are seen between RPE and photoreceptor outer segments (**top right panel, inset**). In places, the RPE basal lamina is interrupted (**arrow**) and electron-dense nodules arise to extend between the basolateral RPE infoldings (**bottom middle panel**). Higher magnification is given (**bottom right panel**). **C:** Immunoreactivity for collagen IV (red) of the retinal/choroidal interface in early-induced, 4-week-old *Tgfb2^{Δeye}* mice. In the control, the basal laminae of RPE and choriocapillaris endothelium are labeled. In *Tgfb2^{Δeye}* mice, staining is seen in the basal lamina of the choriocapillaris. Staining is irregular and patchy in the region of the RPE. Nuclei are DAPI stained (blue). RPE nuclei are marked by **arrowheads**. **D:** Immunoreactivity for collagen IV (red) in the retinal/choroidal interface in late-induced, 3-month-old *Tgfb2^{Δeye}* mice. In the control, the basal laminae of RPE and choriocapillaris endothelium are continuously labeled. In *Tgfb2^{Δeye}* mice, labeling is seen in the basal lamina surrounding choriocapillaris vessels, but is incomplete underneath the RPE. In places, the vascular basal lamina continues between RPE cells (**arrows**), indicating areas of choroidal neovascularization. Nuclei are DAPI stained (blue). RPE nuclei are marked by **arrowheads**. Scale bars: 500 nm (**A and B, top panels**); 250 nm (**B, bottom panels**); 50 μ m (**C and D**). ONL, outer nuclear layer.

4-week-old *Tgfb2^{Δeye}* mice had significantly elevated mRNA expression levels for the angiogenic factors Vegf-a 120, Vegf-a 164, fibroblast growth factor-2, insulin growth factor-1, angiopoietin 2, and platelet-derived growth factor-b compared to control littermates.²¹ We now performed immunohistochemistry for VEGF-A to localize the increased levels in the retina of early-induced *Tgfb2^{Δeye}* mice (**Figure 6E**). Although VEGF-A immunoreactivity was not detectable in control eyes, there was distinct and intense staining of the ganglion cell layer of early-induced *Tgfb2^{Δeye}* mice. When we screened the retinas of late-induced *Tgfb2^{Δeye}* mice and their control littermates for the expression levels of the Vegf-a isoforms 120 and 164, the expression of Vegf-a 120 was not altered, whereas the expression of Vegf-a 164 was significantly reduced in 5-week-old late-induced *Tgfb2^{Δeye}* mice compared to controls (**Figure 6F**). However, at the age of 6 months, the

retinal expression levels of both Vegf-a 120 and 164 were significantly higher in late-induced *Tgfb2^{Δeye}* mice compared to controls. Overall, angiogenic and immune reactivities were much higher in early-induced than in late-induced *Tgfb2^{Δeye}* mice.

Formation of Basal Lamina Deposits in *Tgfb2^{Δeye}* Mice

Because CNVs need to find their way across BM and through the RPE barrier, we used transmission electron microscopy to analyze the structure of both. Control mice showed the regular, five-layered architecture of the BM (**Figure 5, A and B**). In early- and late-induced *Tgfb2^{Δeye}* mice, the BM had thickened as the result of an accumulation of collagen fibers and fine fibrillar extracellular material was detected between the basal lamina of the choriocapillaris and the elastic layer of the BM (**Figure 5, A and B**).

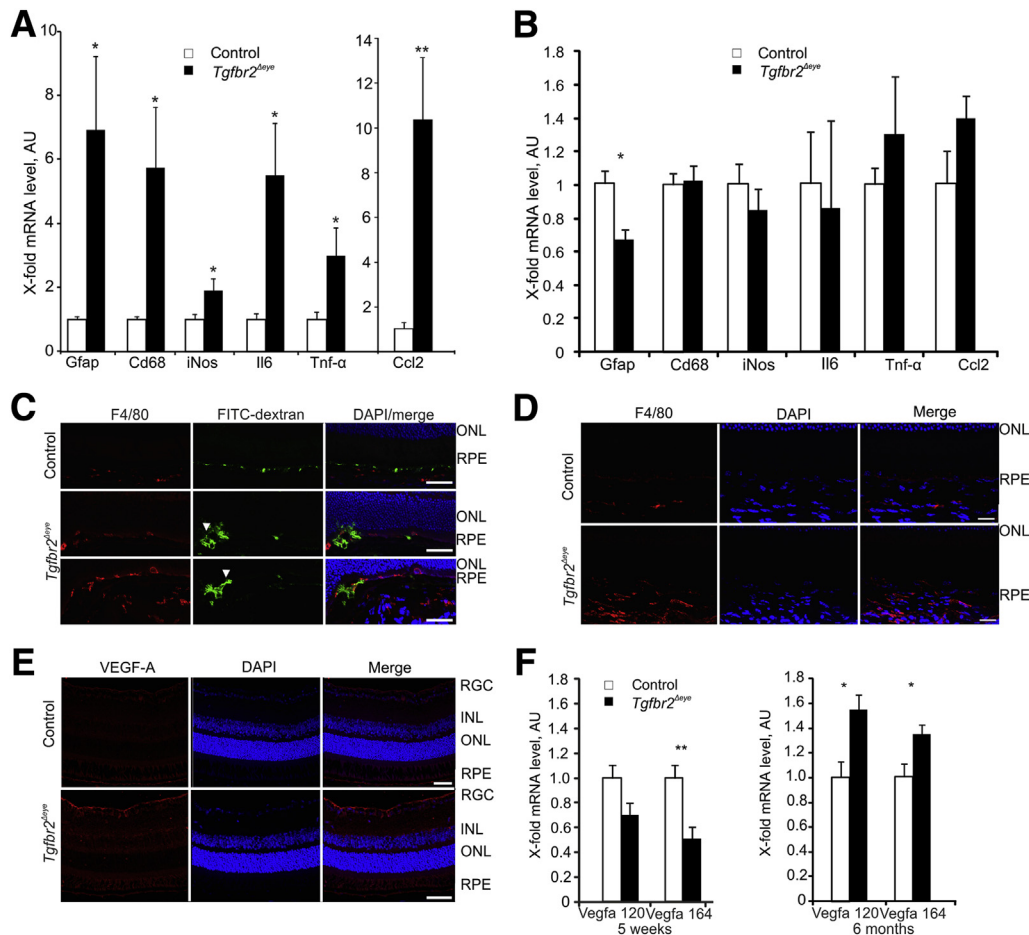


Figure 6 Expression of angiogenic and immune-modulating molecules in early- and late-induced *Tgfb2^{Δeye}* mice. **A** and **B**: Real-time RT-PCR for mRNA of glial fibrillary acidic protein (Gfap), Cd68, inducible nitric oxide synthase (iNos), chemokine (C-C motif) ligand 2 (Ccl2), IL-6, and tumor necrosis factor- α (Tnf- α) in 6-week-old early-induced (**A**) and 5-week-old late-induced (**B**) *Tgfb2^{Δeye}* mice and controls. **C**: Fluorescein isothiocyanate (FITC)-dextran (green)—perfused and F4/80 (red)—immunostained sections of a 6-week-old early-induced *Tgfb2^{Δeye}* mouse and control. The control mouse shows weak immunoreactivity for F4/80 (red)—positive cells in the choroid. In *Tgfb2^{Δeye}* mice, the choroidal immunoreactivity for F4/80 (red) is more pronounced, and numerous F4/80-positive cells accumulate in the subretinal space and in close association to choroidal vessels penetrating the retinal pigment epithelium (RPE; **arrowhead**). Nuclei are DAPI stained (blue). **D**: F4/80 (red)—immunostained sections of a 3-month-old, late-induced *Tgfb2^{Δeye}* mouse and control. A weak signal for F4/80 is visible in the choroid of the control mouse. In contrast, the *Tgfb2^{Δeye}* mouse shows numerous F4/80-positive cells in the choroid. Nuclei are DAPI stained (blue). **E**: Immunoreactivity for vascular endothelial growth factor-A (VEGF-A; red) in early-induced 4-week-old mice. The control mouse shows a faint immunoreactivity for VEGF-A, which is increased in the *Tgfb2^{Δeye}* mouse. Nuclei are DAPI stained (blue). **F**: Real-time RT-PCR for mRNA of Vegfa 120 and Vegfa 164 in late-induced 5-week-old mice (**left panel**) and 6-month-old mice (**right panel**). Data are expressed as means \pm SEM (**A**, **B**, and **F**). $n \geq 7$ early-induced mice, control and *Tgfb2^{Δeye}* (**A**); $n \geq 6$ late-induced mice (5 weeks old), control (**B** and **F**); $n \geq 2$ late-induced mice (5 weeks old), *Tgfb2^{Δeye}* (**B** and **F**); $n = 9$ late-induced mice (6 months old), control (**F**); $n = 7$ late-induced mice (6 months old), *Tgfb2^{Δeye}* (**F**). * $P \leq 0.05$, ** $P \leq 0.01$ (*t*-test). Scale bars: 50 μ m (**C** and **E**); 20 μ m (**D**). AU, arbitrary unit; INL, inner nuclear layer; ONL, outer nuclear layer; RGC, retinal ganglion cell.

Moreover, in some areas, the RPE basal lamina had been replaced by polymorphous electron-dense material that was localized between the elastic layer of the BM and the RPE basal infoldings (**Figure 5A**). In other areas of the same eye, irregular nodules arising from the RPE basal lamina and with comparable electron density were found between the basal infoldings of the RPE (**Figure 5A**). In addition, the basal lamina was frequently found to be interrupted where nodules arised (**Figure 5, A and B**). Nodules and RPE basal lamina interruptions were found frequently in early-induced mice and more rarely in late-induced mice. The changes were essentially similar in structure to basal lamina deposits typically found in CNV in humans patients with AMD. In

addition, we found interruptions of the RPE basal lamina with an associated accumulation of electron-dense material in the adjacent RPE infoldings (**Figure 5A**). Next, we labeled the basal laminae of RPE and choriocapillaris by collagen type IV immunohistochemistry. In control mice, the basal laminae of both RPE and choriocapillaris endothelium were continuously labeled (**Figure 5, C and D**). In early-induced *Tgfb2^{Δeye}* mice, continuous staining was only seen in the basal lamina of the choriocapillaris. Staining was irregular and patchy in the region of the RPE (**Figure 5C**). In late-induced *Tgfb2^{Δeye}* mice, continuous labeling was seen in the basal lamina surrounding choriocapillaris vessels, but was incomplete underneath the RPE

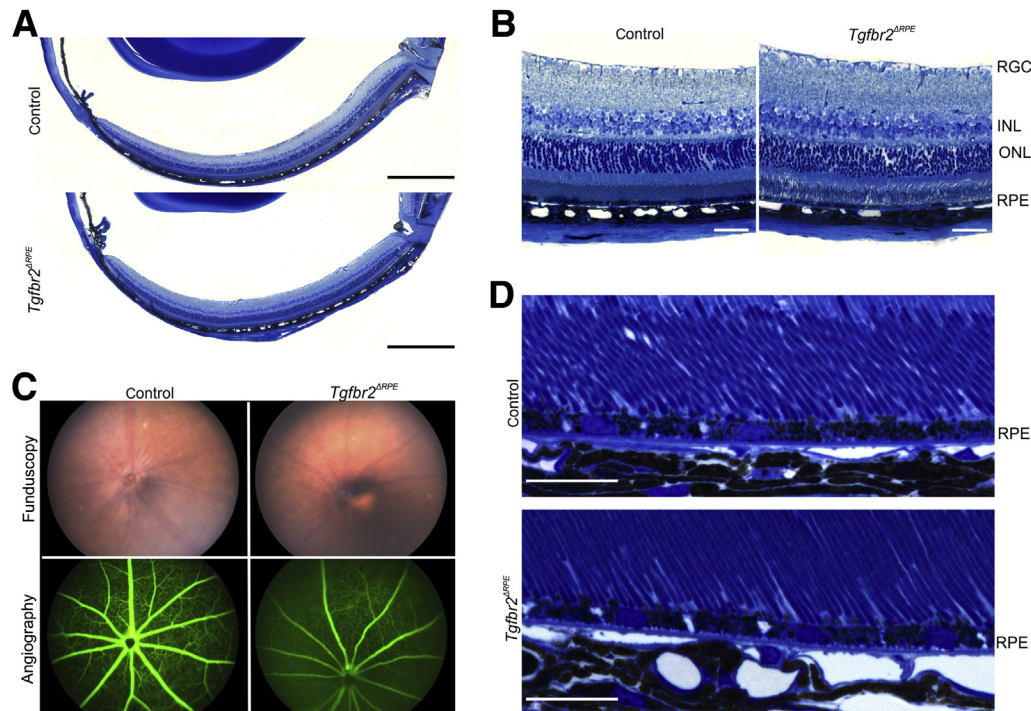


Figure 7 Deletion of transforming growth factor- β (TGF- β) signaling in the retinal pigment epithelium (RPE). Stained semithin sections, according to the method of Richardson et al.³⁷ **A:** Retinal hemispheres of 6-month-old animals. The control mouse and the *Tgfb2* ^{Δ RPE} mouse do not obviously differ in structure. **B:** Detailed magnification of retina/choroid in a *Tgfb2* ^{Δ RPE} mouse and its control littermate, showing an essentially normal morphology. **C:** *In vivo* funduscopy and fluorescein angiography show no obvious alterations in the *Tgfb2* ^{Δ RPE} mouse compared to the control. Diameters of the image sections refer to 2 mm retina. **D:** Detailed magnifications of the interface of the outer retina, RPE, and choroid, again showing a regular morphology in the *Tgfb2* ^{Δ RPE} mice and controls. Scale bars: 500 μ m (**A**); 50 μ m (**B**); 20 μ m (**D**). INL, inner nuclear layer; ONL, outer nuclear layer; RGC, retinal ganglion cell.

(Figure 5D). In places, the vascular basal lamina reached between RPE cells, indicating areas of CNV formation.

Cell-Specific Conditional Deletion of *Tgfb2* in the RPE and Vascular Endothelium

Next, we aimed at identifying the specific cell type that is responsible for CNV in *Tgfb2* ^{Δ eye} mice. We focused on the RPE and the vascular endothelium, as endothelial proliferation and breakdown of the RPE barrier are essential requirements for CNV formation. To this end, we generated *Tgfb2* ^{Δ RPE} mice with a *Tgfb2* deletion specifically in the RPE via doxycycline-driven RPE-specific expression of Cre recombinase. Successful RPE-specific T β RII deletion was confirmed by immunohistochemistry. Control animals showed immunoreactivity for T β RII in photoreceptor outer segments, RPE, and choroid. In contrast, immunoreactivity for T β RII in the RPE was not detectable in the RPE of *Tgfb2* ^{Δ RPE} mice (Supplemental Figure S2C). In addition, we confirmed recombination by PCR (Supplemental Figure S2D). The ocular morphology of *Tgfb2* ^{Δ RPE} mice was analyzed in detail by using essentially similar methods as used for *Tgfb2* ^{Δ eye} mice, but neither choroidal CNV nor other obvious structural changes were detected (Figure 7, A–D). For generation of *Tgfb2* ^{Δ EC} mice with specific deletion of T β RII in vascular endothelial cells, *VeCad-Cre-*

ER^{T2} mice were used with endothelial-specific tamoxifen-inducible Cre expression. Specific recombination in retinal and choroidal vessels was confirmed by green fluorescent protein immunoreactivity in *VeCad-Cre-ER*^{T2} crossed with *mT/mG* reporter mice (Supplemental Figure S2E). When analyzing the structure of 6-week-old *Tgfb2* ^{Δ EC} mice, the pronounced structural changes that we had reported²¹ for early-induced *Tgfb2* ^{Δ eye} mice, such as retinal and vitreal neovascularization, retinal detachment, and vitreal hemorrhages, were completely absent (Figure 8, A and B). Still, similar to our results seen in early-induced *Tgfb2* ^{Δ eye} mice,²¹ dilated vessels were frequently observed in the inner nuclear layer surrounded by electron-dense, extravascular material (Figure 8, C and E). Intriguingly, at the retinal/choroidal interface, focal areas were frequently observed in which photoreceptor outer segments were degenerated or absent, the RPE was multilayered, and vascular endothelial cells, together with extravasated erythrocytes, had accumulated in the subretinal space (Figure 8D). FITC-dextran-perfused ocular sections of *Tgfb2* ^{Δ EC} mice showed tracer leakage from the choriocapillaris toward the RPE and choroidal vessels penetrating the RPE, indicating a breakdown of the outer blood-retinal barrier and the formation of CNV (Figure 9A). In 4-week-old *Tgfb2* ^{Δ EC} animals and controls, the choroidal vessels were covered by neural/glial antigen 2-positive pericytes (data not shown). In contrast to

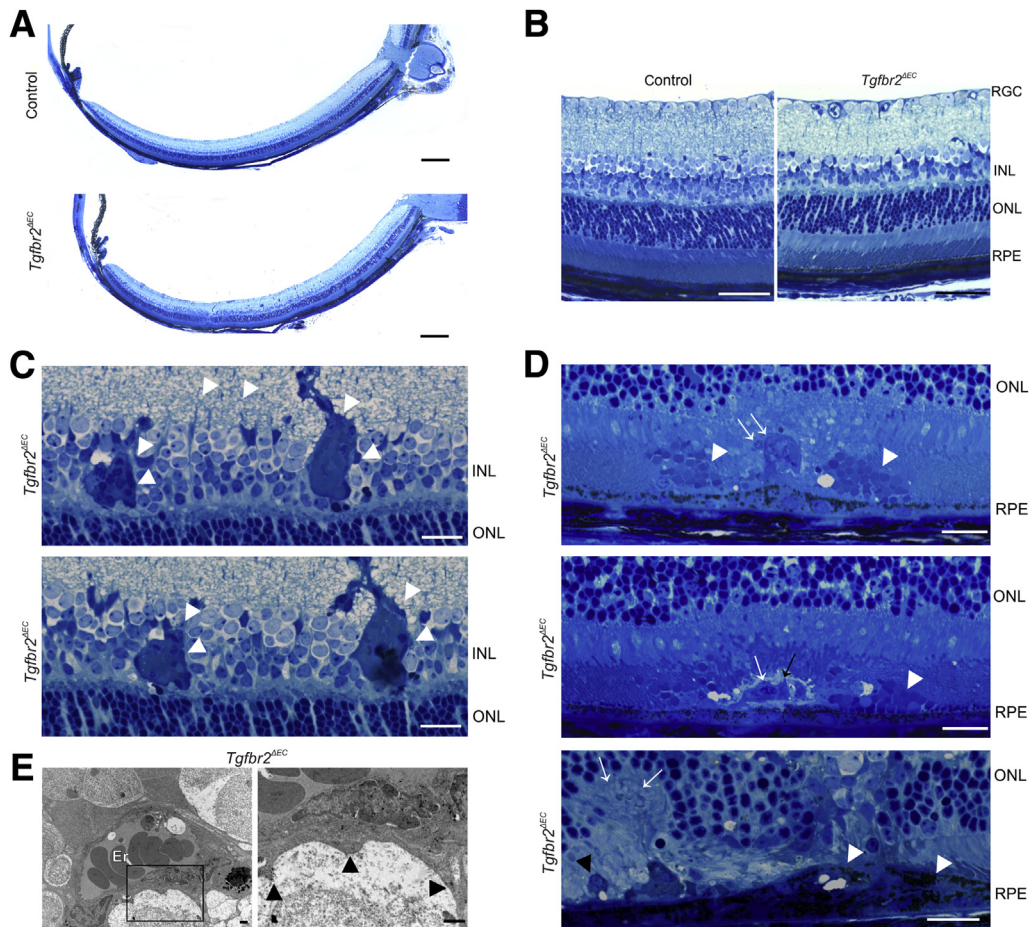


Figure 8 Structural changes of the retina in *Tgfb2 Δ EC* mice. Stained semithin sections of a 6-week-old *Tgfb2 Δ EC* mouse and its control littermate, according to the method of Richardson et al.³⁷ **A:** Retinal hemispheres of the control mouse and the *Tgfb2 Δ EC* littermate. **B:** Detailed magnification of the retina and choroid. **C:** Dilated retinal vessels (**white arrowheads**) are seen in the inner nuclear layer (INL) of *Tgfb2 Δ EC* mice. **D:** In the retinal/choroidal interface, endothelial cells (**white arrows**), erythrocytes (Ers; **white arrowheads**), mononuclear cells (**black arrowhead**), and ragged edged extravascular material (**black arrow**) are seen between degenerating photoreceptor outer segments. **E:** Transmission electron microscopy of an intraretinal vessel in a 6-week-old *Tgfb2 Δ EC* animal. Electron-dense material is visible around the intraretinal vessel (**left panel**); the **boxed area** is shown in higher magnification (**right panel**). The **arrowheads** point to the electron-dense extravascular material with ragged edged appearance. Scale bars: 200 μ m (**A**); 50 μ m (**B**); 20 μ m (**C** and **D**); 1000 nm (**E**). ONL, outer nuclear layer; RGC, retinal ganglion cell; RPE, retinal pigment epithelium.

our data from *Tgfb2 Δ eye* mice, we did not observe an accumulation of F4/80-positive cells in the choroid of *Tgfb2 Δ EC* mice (**Figure 9B**) (control: 95.0 ± 8.40 , $n = 4$; *Tgfb2 Δ EC*: 111.67 ± 10.41 , $n = 3$; $P \geq 0.05$). In contrast, F4/80-positive cells were increased in the retina (control: 22.5 ± 1.71 , $n = 4$; *Tgfb2 Δ EC*: 108.0 ± 3.51 , $n = 3$; $P \leq 0.001$). In addition, immunoreactivity for collagen IV in the RPE/BM region was seen in basal laminae of choriocapillaris and RPE and did not markedly differ between *Tgfb2 Δ EC* mice and controls (**Figure 9C**). Transmission electron microscopy confirmed the findings seen by light microscopy and the presence of degenerated photoreceptor outer segments, multilayered RPE, and extravasated erythrocytes (**Figure 9D**). Moreover, plasma-derived electron-dense material did not pass the RPE tight junctions in controls but accumulated between RPE and photoreceptor outer segments in *Tgfb2 Δ EC* mice, indicating breakdown of the RPE barrier (**Figure 9E**). Overall, our data obtained in

Tgfb2 Δ EC mice strongly support the conclusion that deletion of T β RII in vascular endothelial cells alone is sufficient to promote the formation of CNV.

Discussion

We conclude that the deletion of TGF- β signaling in the ocular microenvironment is sufficient to induce CNV and other phenotypic characteristics of AMD in humans. Lack of endothelial TGF- β signaling is sufficient to trigger the onset of CNV, whereas its lack in the RPE is not relevant in this context. This conclusion is based on the following: i) the generation of mice with T β RII deficiency in the entire microenvironment of the eye, in vascular endothelial cells, or in the RPE; ii) the frequent detection of capillaries with fenestrated, PLVAP-positive endothelium that originate from the choriocapillaris and traverse the RPE to

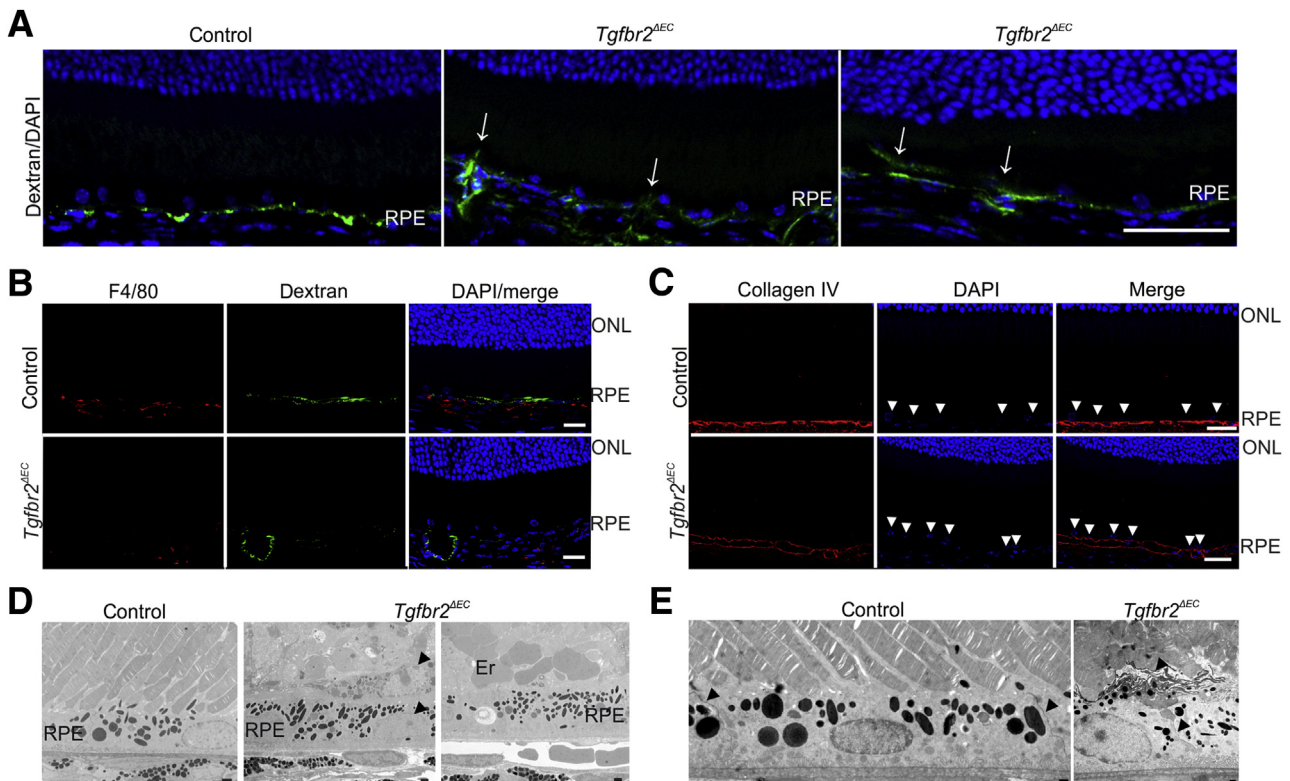


Figure 9 Structural changes of the retinal/choroidal interface in *Tgfb2^{ΔEC}* mice. **A:** Fluorescein isothiocyanate (FITC)—dextran—perfused retinal meridional sections of a 6-week-old *Tgfb2^{ΔEC}* mouse and its control littermate. **White arrows** point toward tracer leakage in the retinal pigment epithelium (RPE; **middle panel**) and choroidal vessels (**right panel**) invading the RPE. Nuclei are DAPI stained (blue). **B:** FITC-dextran (green)—perfused and F4/80 (red)—immunostained sections. The control and the *Tgfb2^{ΔEC}* littermate show immunoreactivity for F4/80 (red)—positive cells in the choroid, which are not higher in number in the mutant. Nuclei are DAPI stained (blue). **C:** Immunoreactivity for collagen IV (red) in the RPE/Bruch membrane (BM) region of 6-week-old mice. Basal laminae of choriocapillaris and RPE are regularly labeled in control and mutant. Nuclei are DAPI stained (blue), and RPE nuclei are marked by **white arrowheads**. **D** and **E:** Transmission electron microscopy of RPE/BM in 4- to 6-week-old mice. **D:** In the control, photoreceptor outer segments, RPE, and BM are of normal structure. In the *Tgfb2^{ΔEC}* mouse, areas of degenerated photoreceptor outer segments, multilayered RPE (**black arrowheads** point toward nuclei of RPE cells), and extravasated erythrocyte (Er) are present. **E:** Plasma-derived electron-dense material does not pass the RPE tight junctions in controls (**black arrowheads**). In contrast, in the mutant, it accumulates between RPE and photoreceptor outer segments (**black arrowheads**). Scale bars: 50 μ m (**A**); 20 μ m (**B** and **C**); 1000 nm (**D**); 500 nm (**E**). ONL, outer nuclear layer.

anastomose with retinal capillaries; iii) the presence of basal lamina—like deposits around the RPE basal infoldings and of areas with multilayered RPE; iv) the degeneration of photoreceptor outer segments; and v) the finding of distinct accumulations of F4/80-positive macrophages in the choroid.

TGF- β Functions at the Retinal/Choroidal Interface to Prevent CNV

The results of our study clearly support the concept that a major function of TGF- β signaling in choroid and retina is the stabilization of the choroidal and retinal vascular beds that are each essential for neuronal integrity in the sensory retina. This function includes the prevention of neo-angiogenic processes in the two capillary beds that would otherwise cause neuronal dysfunction and death in the retina. It is of interest that the sensitivity to the induced lack of TGF- β signaling differs between the two vascular beds, as the formation of microaneurysms, leaky capillaries, and

hemorrhages of the retinal vasculature is only seen when TGF- β signaling is deleted in the ocular microenvironment shortly after birth.²¹ In contrast, the formation of CNV is seen regardless if TGF- β signaling is interfered with in newborn or 3- to 4-week-old animals. A likely explanation is the fact that the retinal vasculature of the mouse eye forms and differentiates in the first 2 weeks after birth⁵² and may be more vulnerable to lack of TGF- β signaling during that period. A critical contributing factor may be the failure of pericyte differentiation around retinal capillaries that results from deficiency of TGF- β signaling during that period.²¹ Formation of the choroidal vasculature is completed in late embryonic life, and the choriocapillaris is mature at birth.⁵³

The coating of the choroidal vessels with neural/glial antigen 2—positive pericytes in early- and late-induced *Tgfb2^{Δeye}* and *Tgfb2^{ΔEC}* animals was comparable to controls, which appears to indicate that lack of pericyte presence or differentiation is not a requirement for CNV. Data from genetically modified mice with a specific

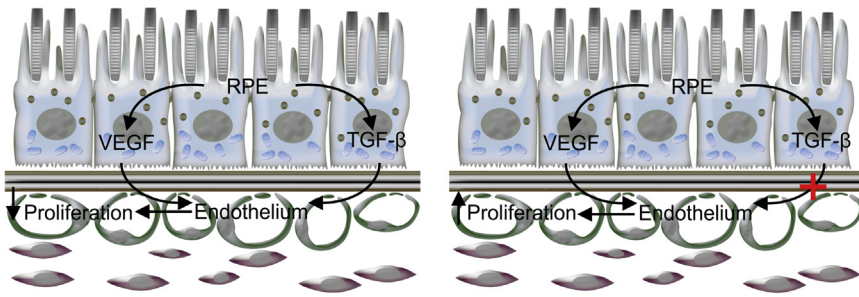


Figure 10 Schematic of signaling events at the retinal-choroidal interface. **Left panel:** Normally, the retinal pigment epithelium (RPE) secretes high amounts of both transforming growth factor- β (TGF- β) and vascular endothelial growth factor (VEGF) that both ensure an appropriate physiological microenvironment for the choriocapillaris, including its VEGF-mediated maintenance, and TGF- β -mediated inhibition of proliferation. **Right panel:** The deletion of *Tgfr2* in endothelial cells results in the specific disability of TGF- β to act on endothelial cells (red cross). This results in an imbalance of the effects of VEGF/TGF- β , a scenario that promotes the proliferation of the vascular endothelium of the choriocapillaris in the direction of the RPE and, finally, results in choroidal neovascularization.

deficiency of platelet-derived growth factor-B in endothelial cells further challenge the role of pericytes in the formation of CNV as these mice show a varying degree of pericyte loss without developing CNV.⁵⁴ The higher sensitivity of the choriocapillaris to lack of TGF- β signaling, as opposed to the mature retinal vasculature, may also be caused by the presence of the high amounts of VEGF at the retinal/choroidal interface that are continuously secreted by the RPE, the only source of VEGF in the back of the adult eye.⁵⁵ Secretion of VEGF occurs in a polarized manner to the RPE basolateral side facing the choriocapillaris^{56,57} and is expected to convey a proliferative signal to endothelial cells of the choriocapillaris that are no more under the influence of constitutive TGF- β signaling in animals with a deletion of T β RII.

TGF- β and VEGF as Part of a Homeostatic System to Maintain Integrity of the Choriocapillaris

Not only VEGF but also TGF- β 1 and TGF- β 2 are present at high amounts at the retinal/choroidal interface.²⁰ It is tempting to speculate that both factors are critical parts of a homeostatic system designed to maintain structure and function of the choriocapillaris. In this system, VEGF would be required to maintain the extreme high density of the choriocapillaris and its fenestrations,^{57,58} whereas TGF- β s antagonize any proliferative properties of VEGF signaling on the vascular endothelium. Failure in the balance of this homeostatic system would cause ablation of the choriocapillaris or induce its proliferation, finally leading to CNV (Figure 10). There is direct evidence for such a homeostatic system from studies in genetically engineered mice in which lack of VEGF and/or increase in TGF- β activity leads to ablation of the choriocapillaris,^{14–18} whereas an increase in VEGF^{59,60} and/or a lack of TGF- β activity (results of our present study) have direct opposite and proliferative effects. For the initiation of CNV, lack of TGF- β activity appears to be more relevant than sole increase of VEGF activity, as mice with overexpression of VEGF in the RPE develop an intrachoroidal neovascularization, but no CNV.^{59,60} VEGF

and TGF- β s induce the transcription of one another in multiple cell types^{61–66} and may well do so in the RPE, establishing an autoregulatory feedback system designed to maintain structure and function of the choriocapillaris. We realize that this concept needs to be validated in further studies that should also aim at a complete identification of the signaling network that drives endothelial proliferation and CNV formation downstream of TGF- β signaling deficiency.

Potential Neuroprotective Effects of TGF- β

Late-induced *Tgfr2*^{4eye} mice showed a dramatic deterioration of the retina. The prolonged presence and formation of CNV, and its detrimental effects on structure and function of the retina, may explain this finding. It may also indicate a participation of TGF- β signaling in a neuroprotective pathway independent of the formation of CNV. We recently showed that TGF- β signaling protects retinal neurons from developmental programmed cell death.³³ It is well possible that it also contributes to maintenance of adult neurons by protecting them from apoptosis. In support of such a scenario are findings reported by Walshe et al,⁶⁷ who neutralized TGF- β in the adult mouse eye via expression of soluble endoglin, a TGF- β inhibitor. Apoptosis of retinal ganglion cells was observed, as were functional deficits detected by ERG. Neuroprotective activities of TGF- β s were reported for multiple types of neurons throughout the central nervous system, such as in the striatum,⁶⁸ spinal cord,⁶⁹ substantia nigra,⁷⁰ or hippocampus.⁷¹ Similarly, there is increasing evidence that VEGF signaling is important for the trophic maintenance of neurons and their survival after injury, effects that appear to be mediated via the VEGF receptor-2.^{72–76} Mice with a constitutive high expression of VEGF in retinal ganglion cells are protected from retinal ganglion cell degeneration after axotomy via VEGF receptor-2 and downstream activation of extracellular signal-regulated kinase 1/2 and Akt pathways.⁷⁷ Although TGF- β /VEGF are antagonists in their actions at the retinal/choroidal interface and the choriocapillaris, they

may cooperate in their neuroprotective activities. Both functions would serve the ultimate goal to maintain structure and function of the retina.

The Formation of BlamD-Like Material Is Not Required for CNV

Although CNV was consistently observed in the eyes of both early- and late-induced *Tgfb2^{Δeye}* mice, and in those of *Tgfb2^{ΔEC}* mice, other findings were predominant only in the eyes of early-induced *Tgfb2^{Δeye}* mice. This includes the formation of homogeneous extracellular material between the basal infoldings of the RPE and internal to the RPE basal lamina. The material was similar in electron density and structure to that of basal lamina deposits (BlamDs), a characteristic finding in patients with early age-related macular degeneration.^{78–80} BlamD-like changes were commonly observed in early-induced *Tgfb2^{Δeye}* mice, only rarely observed in late-induced *Tgfb2^{Δeye}* mice, and not observed in *Tgfb2^{ΔEC}* mice, suggesting that they are not a requirement for CNV and breakdown of the RPE barrier. Our immunohistochemical data indicate that the homogeneous extracellular BlamD-like material contains collagen type IV, an observation that correlates with the observation that basal lamina proteins (eg, collagen IV and laminin) are present in BlamDs in human patients with AMD.^{78,81} BlamD formation in AMD is likely caused by RPE dysfunction.⁷⁹ A comparable dysfunction might more easily happen in early-induced *Tgfb2^{Δeye}* mice, in which the RPE is not yet fully differentiated. Still, a common structural change of the RPE in our models with CNV was the loss of an epithelial monolayer and the formation of multilayered RPE cells. Beneath and above those multilayered RPE cells, newly formed vessels were observed, similar to type 1 (sub-RPE, occult) and type 2 (subretinal, classic) CNV in human patients. Direct deletion of TGF-β signaling in RPE cells appears not to be relevant for formation of BlamD-like material, as it was not observed in *Tgfb2^{ΔRPE}* mice.

Neovascularization Is Associated with Macrophage Accumulation

A common finding in *Tgfb2^{Δeye}* mice with CNV was the marked accumulation of F4/80-positive macrophages in the region of the retinal/choroidal interface. Comparable findings have been described for patients experiencing AMD.⁸² It is reasonable to assume that signaling molecules released from proliferative capillary endothelial cells attracted the cells. Reactive macrophages might contribute to the loss of the RPE barrier in our mice and its transition from an epithelial layer to a multilayer in areas of CNV. Our findings in *Tgfb2^{ΔRPE}* mice indicate that lack of TGF-β signaling in RPE cells is not required for this process. Currently, we do not know how proliferating choroidal endothelial cells manage to break through the RPE barrier, but we trust that our mouse models are appropriate tools to identify the causative molecular

events. The expression of signaling molecules characteristic for reactive macrophages/microglia, such as inducible nitric oxide synthase, Il-6, tumor necrosis factor-α, and monocyte chemoattractant protein-1/chemokine (C-C motif) ligand, was elevated in the retinal mRNA of early-induced *Tgfb2^{Δeye}* mice, but not in late-induced mice. We attribute this observation to the marked changes in the inner retina of early-induced mice that are absent in late-induced mice. Still, those molecules might well be elevated in the microenvironment of the choroidal/retinal interface, where macrophages accumulate in mice with CNV.

TGF-β Signaling and Blood-Retinal Barrier

Leakage of high-mol. wt. FITC-dextran and erythrocytes into the retina and subretinal space, after induced TGF-β signaling deficiency, implicates disruption of the inner and outer blood retinal barriers. Deficiency of the TGF-β signaling pathway in endothelial cells is likely the major contributor in this scenario. Comparable, mice with an endothelial-specific deficiency of SMAD4, an intracellular downstream mediator of TGF-β signaling, or mice with a deletion of TβRII specifically in endothelial cells in the brain^{83,84} show perinatal intracerebral hemorrhages and a breakdown of the blood-brain barrier. Furthermore, the virus-driven expression of soluble endoglin results in an inhibition of TGF-β1 signaling in the murine retina, a situation that also causes breakdown of the blood-retinal barrier, most likely mediated through a decreased association of the tight junction proteins occludin and zona occludens-1.⁶⁷

TGF-β Signaling in Human Patients with AMD

Several independent case-control genome-wide association studies detected an association of two synonymous polymorphisms in exon 1 of the high-temperature requirement A1 (*HTRA1*) gene, with a high risk to develop AMD.^{85–87} The gene product HTRA1 appears to play a causative role in CNV.^{88–90} Intriguingly, a recent study provided evidence that the two synonymous HTRA1 variants influence their protein interaction with TGF-β1, leading to an impaired regulation of TGF-β signaling.⁹¹ Moreover, a collaborative genome-wide association study identified *TGFBR1*, the gene encoding for the TGF-β1 type I receptor, as a new susceptibility gene for AMD, further highlighting the importance of the TGF-β signaling pathway in the context of AMD.⁸⁵

Conclusion

Our findings emphasize the importance of TGF-β signaling as a key player in the development of ocular neovascularization and implicate a fundamental role of TGF-β signaling in the pathogenesis of AMD. A more thorough understanding of this role at the retinal/choroidal interface has the distinct potential to lead to the development of novel

treatments strategies preventing CNV in patients experiencing AMD.

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Supplemental Data

Supplemental material for this article can be found at <http://dx.doi.org/10.1016/j.ajpath.2017.06.018>.

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