Effect of subconjuctival and intraocular bevacizumab injection on angiogenic gene expression levels in a mouse model of corneal neovascularization

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Purpose: This study sought to characterize the expression of angiogenesis-related genes in a mouse model of corneal neovascularization, either untreated or after treatment with a single injection of bevacizumab by three different routes. In addition, the effectiveness of the treatment was compared to a rabbit model.

Methods: A chemical burn was induced in the mid-cornea of the right eye in 119 mice; 56 of them were untreated and 63 were bevacizumab-treated. Neovascularization was evaluated 2, 4, 8, 10, and 14 days later using digital photos, angiography and India ink perfusion. The relative area of new blood vessels was analyzed using slit-lamp examination in vivo and on histological and flat-mount sections. The levels of gene expression involved in the angiogenic process vascular endothelial growth factor [VEGF], insulin-like growth factor-1 [IGF-1], pigment epithelium derived factor [PEDF], and macrophage-inflammatory protein-2 [MIP-2]) were measured by a real-time polymerase chain reaction. Six rabbits underwent the same injury and treatment, and the response was compared to the mouse model.

Results: Neovascularization was first observed two days after injury. The affected section increased from 11.24% (± 7.0) of the corneal area to 47.42% (± 25.4) on day 8 and 50.62% (± 24.7) on day 10. In the mice treated with bevacizumab, the relative area of neovascularization was significantly lower at the peak time points (p<0.005): 24.90% (± 21.8) on day 8 and 28.29% (± 20.9) on day 10. Spontaneous regression was observed on day 14 in both groups, to 26.98% (± 19.9) in the untreated mice and 10.97% (± 10.8) in the bevacizumab-treated mice (p<0.005). Rabbits also showed peak corneal neovascularization on days 8-10, with significant regression of the vessels following intracameral bevacizumab injection. In the mice, intraocular (intravitreal, intracameral) injection was more effective than subconjuctival injection. *VEGF* gene expression was upregulated in both the untreated and treated mice, but was slightly less in the treated mice. *PEDF* gene expression decreased in both the treated and untreated mice. In the untreated group, gene expression peaked (above baseline) at 14 days, and in the untreated mice, it had already peaked by day 8. IGF-1 was upregulated early in the model; at 8 days, there was only a slight change in the untreated group compared to a significant increase in the treated group. MIP-2 was upregulated in both groups in the early stage and returned to baseline on day 14.

Conclusions: Bevacizumab treatment partially inhibits the progressive corneal neovascularization induced by chemical damage in a mouse model. Treatment is more effective when administered via the intraocular than the subconjunctival route. The clinical findings are compatible with the angiographic and histologic data and are supported by molecular analysis showing a partial change in expression of proangiogenic genes. The molecular mechanisms involved in corneal neovascularization and inflammation warrant further exploration. These findings may have important therapeutic implications in the clinical setting.

Angiogenesis is one of the key factors in tumor progression and metastasis. Anti-angiogenic therapy inhibits tumor angiogenesis and promotes apoptosis of existing blood vessels, thereby intercepting the tumor's supply of oxygen and nutrients. Furthermore, by normalizing vascular permeability,

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anti-angiogenic therapy can improve the delivery of therapeutic agents to tumor cells.

A normal, healthy cornea is devoid of both blood and lymphatic vessels [1-3], but inflammatory conditions such as chemical burns, trauma or infection can cause angiogenesis [4-6]. The development of new blood vessels in the cornea, called neovascularization, is a final pathway common to all insults that are resistant to treatment and may lead to severe impairment of vision. Corneal neovascularization has been reported in 4.14% of patients presenting for general

ophthalmologic care in the USA, representing an estimated 1.4 million individuals [7].

Until recently, the mainstay of anti-angiogenic therapy in the cornea was nonspecific anti-inflammatory drugs and sometimes anti-angiogenic steroids, but they were often unable to prevent or stop neovascularization [8]. Even surgical treatment with corneal transplantation frequently failed, as the new vessels induced an inflammatory rejection of the graft [7,9]. However, the rapid progress in angiogenesis research in the last few years has led to the development of several novel, specific anti-angiogenic drugs for use in both oncology and ophthalmology [10]. As the vast majority of these agents are low-molecular-weight compounds with pharmacokinetic profile and rapid clearance rate, researchers have found that by conjugating them to polymeric carriers, their solubility and specificity may be increased and their pharmacokinetics is improved [11,12]. Recent reports describe the development of polymeric nanospheres that selectively target the activated vascular endothelium and can deliver encapsulated anti-angiogenic agents in cases of inflammatory disease with an angiogenic component [13].

A major focus of the research into anti-angiogenic therapy is vascular endothelial growth factor (VEGF), which is known to promote several steps in angiogenesis, including proteolytic activities, endothelial cell proliferation, endothelial cell migration and capillary tube formation [6]. Its essential role in normal embryonic vasculogenesis and angiogenesis was supported by findings that inactivation of a single VEGF allele in mice resulted in death of the embryo [14]. VEGF exerts its activity by binding to several highaffinity transmembrane endothelial cell receptors, especially VEGFR-1 (Flt-1) and VEGFR-2 (KDR/Flk-1). This leads to intracellular receptor phosphorylation, which in turn triggers the relevant intracellular downstream receptor pathways [15]. VEGF has been found to play a role in corneal neovascularization in both experimental models and humans [16-18]. Measurements of VEGF molecules and receptors in diseased corneas yielded higher concentrations than in normal or avascular abnormal corneas. Treatment strategies targeting the VEGF signaling pathway include anti-VEGF antibodies, soluble receptors binding directly to the VEGF ligand, anti-VEGF receptor (VEGFR) antibodies and VEGFR tyrosine kinase inhibitors. Anti-VEGF therapy has proven highly effective in treating advanced colon cancer and is currently being tested in gastric cancer [19]. In several studies, the manipulation of various anti-VEGF agents led to effective inhibition of corneal neovascularization [3,20].

Bevacizumab (Avastin; Genentech, San Francisco, CA) is a full-length recombinant humanized murine monoclonal antibody that binds to and inhibits the biological activity of all five human VEGF-A isoforms: VEGF115, VEGF121, VEGF165, VEGF189, and VEGF206 [21]. It prevents VEGF-A from ligating to its endothelial receptors, VEGFR-1 and

VEGFR-2 [22,23], but does not neutralize other members of the VEGF gene family, such as VEGF-B or VEGF-C [24-26]. The antibody was engineered by assembling VEGF-A binding residues from the murine-neutralizing antibody into the framework of a human immunoglobulin [27]. Bevacizumab has been approved by the US Food and Drug Administration for use in the treatment of metastatic colorectal cancer. Since mid-2005, it has been applied offlabel in the treatment of ocular disease and has shown promising short-term results in alleviating intraocular neovascular conditions [28-30].

Reports that the systemic application of bevacizumab in models inhibited inflammatory animal neovascularization [20,23] led to the assumption that bevacizumab might also be clinically beneficial in patients who do not respond to conventional steroid therapy. This was followed by successful experiences with bevacizumab applied as eye drops [31-33] or subconjunctivally [34-36] in patients with progressive corneal neovascularization. Topical treatment was found to be safe and efficient, without local or systemic adverse effects [37,38]; some authors suggested that it could serve as a pre-transplantation treatment option [39]. Further studies reported a reduction in corneal inflammation and choroidal neovascularization after subconjunctival injection of bevacizumab [35,40-43], and a reduction in neovascular glaucoma after intracameral injection [44,45]. Intravitreal injection is currently the most common route of treatment for age-related macular degeneration with choroidal neovascularization [46].

Studies have shown that there is a balance in the cornea between angiogenic molecules such as VEGF and erythropoietin (EPO) and anti-angiogenic molecules such as pigment epithelium derived factor (PEDF). Angiogenesis and erythropoiesis represent adaptive responses to hypoxia and are upregulated in ischemic conditions. PEDF, a 50-kDa protein secreted by the retinal pigment epithelium, inhibits the growth of new blood vessels, most likely via the PI3K/Akt pathway, and shows an inverse pattern of expression to VEGF and EPO [47]. Intravitreal bevacizumab injections have been found to reduce aqueous VEGF and increase PEDF in patients with choroidal neovascularization secondary to age-related macular degeneration or pathologic myopia [48].

Insulin-like growth factor (IGF)-1 and its receptor, IGF-1R, have also been implicated in choroidal neovascularization. IGF-1 participates in angiogenesis of the developing retina in newborns [49]. Although it failed to enhance endothelial tube formation in vitro [50], it increased VEGF secretion in cultured ARPE-19 cells [51]. Accordingly, IGF-1R inhibitors may be useful tools in the treatment of conditions associated with neovascularization.

The proangiogenic CXC chemokine macrophage inflammatory protein (MIP)-2 induces endothelial cell chemotaxis [52] and promotes angiogenesis in cancer

TABLE	1	MICE	USED	IN THE	STUDY

	Model-untreated	Bevacizumab-treated (n=63)			
Procedure	(n=56)	SC (n=22)	IV (n=20)	IC (n=21)	Total (n=63)
In vivo digital photographs	32	20	18	19	57
India ink (flat cornea)	2	2	2	2	6
Fluorescein angiography	4	-	-	-	-
Histology (H&E)	8	8	8	8	24
Corneal flat mounts	4	=	=	-	-
Antibody staining	-	-	4	-	4
Real-time PCR	10	12	12	19	57

Several mice underwent more than one procedure. Abbreviations are: SC-subconjunctival, IV-intravitreal, IC-intracameral.

conditions [53]. MIP-2 regulates the production of VEGF, and like VEGF, it mediates the regulation of corneal neovascularization caused by bacterial infection [54]. One study showed that following infection, both the corneal resident cells and infiltrating neutrophils produced MIP-2 and VEGF, and the significantly high levels of both genes paralleled the extensive corneal neovascularization seen at later stages of the disease. Anti-MIP-2 antibody treatment significantly reduced both VEGF expression and corneal neovascularization [54]. However, these findings are controversial. Others have reported that following alkali injury in mice, the injection of neutralizing anti-mouse MIP-2 antibodies had no effect on corneal neovascularization [55].

The present study was conducted in a mouse model of chemically-induced corneal neovascularization. The aim of the study was to evaluate the effect of treatment with bevacizumab by three different routes of injection (subconjuctival, intracameral, and intravitreal). We characterized the damage clinically and angiographically and measured the areas of neovascularization in vivo and on histological and flat-mount sections. Molecular analysis of changes in the levels of expression of angiogenesis-related genes was performed with real-time polymerase chain reaction (PCR). In addition, given the low affinity of bevacizumab to murine VEGF [56], we also qualitatively investigated the ocular effects of bevacizumab treatment in a rabbit model.

METHODS

Animals: The effect of injection route of bevacizumab on corneal neovascularization following a chemical burn was investigated in 119 C57BL57 male mice aged 6–8 weeks and weighing 20–25 g (Harlan Laboratories, Jerusalem). In addition, the qualitative effect of treatment was studied in six female outbreed commercial rabbits (Kfar HaNagid, Israel) weighing 2–2.5 kg and aged six months.

All protocols were conducted in accordance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research and were approved and monitored by the Animal Care Committee of Rabin Medical Center. The animals were housed under a 14 h:10 h light-dark cycle with standard chow and water ad libitum.

Induction of corneal neovascularization:

Mouse model—The mice were placed under general anesthesia by intramuscular injection of combined ketamine/xylazine (80 mg/kg and 8 mg/kg, respectively) supplemented by topical anesthesia (proparacaine hydrochloride 0.5%). The silver nitrate cauterization technique described by Mahoney and Waterbury [57] was used to induce corneal neovascularization. Briefly, under the operating microscope, an applicator stick coated with 75% silver nitrate and 25% potassium nitrate with a diameter of 1.8 mm was pressed on the central right cornea of each animal for 10 seconds. Excess chemical reagent was removed by rinsing the eyes with 5 ml of balanced salt solution (NaCl 0.9%). The left eyes were not injured and served as controls. To increase the reproducibility of the injuries, a single investigator (N.G.-C.) cauterized all animals.

Rabbit model—A similar technique was used to induce central chemical burn in the right eye of the rabbits. *Bevacizumab injections:*

Mouse model—Immediately after cauterization, the treated mice (n=63, Table 1) received a single injection of bevacizumab (25 mg/ml) into the right eye via a Hamilton syringe with a 30-gauge tip (Hamilton, Reno, NV). Injections were delivered by one of three routes: subconjuctival: 2.5 mg/0.1 ml (n= 22), intravitreal: 0.75 mg/0.03 ml (n= 20), or intracameral: 0.5 mg/0.02 ml (n= 21). The left eyes were untreated.

Rabbit model—Bevacizumab was injected into the anterior chamber immediately after induction of chemical burn to the rabbit cornea (n=6).

Digital photographs: To quantify the extent of neovascularization, photographs were taken at 2, 4, 8, 10 and 14 days after cauterization in the mice and rabbits with a Canon A610 digital camera attached to a slit-lamp microscope. The corneal surface area containing neovascular vessels was measured on the photographs as the percentage

TARI	F 2	LIST OF	PRIMERS.

	E 1 (7) 2))	D (21.51)
Gene	Forward (5'-3')	Reverse (3'-5')
VEGF	CACGACAGAAGGAGCAGAA	CGCTGGTAGACGTCCATGA
MIP-2	CTGTATTCCCCTCCATCGTG	CTCGTCACCCACATAGGAGTG
IGF-1	AGAGACCCTTTGCGGGGC	CGGATAGAGCGGGCTGCTT
PEDF	GGACTCTGATCTCAACTGCAAGA	GGAGGAAGAAGATGATGCTCATAC
ACTB	TAGGCACCAGGGTGTGATGGT	CATGTCGTCCCAGTTGGTAACA

of the total area of the cornea. Image analysis was performed on each cornea using a public domain Java image processing program developed at the National Institutes of Health, Washington, D.C.

India ink perfusion: Mice (n=8; six treated, two untreated) were anesthetized with ketamine/xylazine on day 10 after corneal cauterization and perfused with 1 ml of waterproof Sanford-Higgins Black India Ink for drawing (Sanford, Oakbrook, IL). The eyes were removed and the corneas (with limbal regions) were isolated, radial-cut and flat-mounted. The samples were reviewed under a light microscope for new vessels and photographed.

Fluorescein angiography: To confirm the growth of new pathological vessels, untreated mice (n=4) were anesthetized for fluorescein angiography on day 10 after corneal cauterization. Animals were injected with 0.04 ml of 25% sodium fluorescein (AK-Fluor 25% AMP; Akorn, Decatur, IL), and the procedure was performed using a digital fundus camera (Topcon TRC 50x, Farmingdale, NY). Thereafter, the mice were euthanized, and flat corneas filled with the fluorescein dye were analyzed under the microscope.

Histological study: After euthanization of both the treated and untreated groups (n=24, Table 1), both eyes from each subject were enucleated and embedded in paraffin, and 5-μm thick sagittal sections were stained with hematoxylin and eosin and examined under a light microscope. The disruption of the corneal structure and the presence of neovascularization were documented. Measurements were performed in triplicate at 20 μm intervals at all corneal thicknesses.

Corneal flat-mount immunostaining: Mice (n=4) were euthanized by CO₂ inhalation on day 8 after cauterization to study the endothelium. The eyes were enucleated and the corneas isolated and fixed in 4% formaldehyde. Immunochemistry staining for the endothelial marker was performed to detect pathological blood vessels in the cornea. Briefly, corneal flat-mounts were rinsed several times in phosphate buffered saline, blocked with 1% bovine serum albumin, stained with Rat Anti-Mouse Pecam-1 [CD31] monoclonal antibody overnight (1:100; CBL1337, Chemicon, Bellirica, MA), washed three times for 5 min with PBS and stained with goat anti-rat IgG-1 for 1 h at room temperature (1:200; A11077, Molecular Probes, Eugene, OR).

Antibody staining for intraocular bevacizumab: Eyes (n=4) were enucleated 1 day after intravitreal injection, fixed in

formalin, embedded in paraffin wax, sectioned and deparaffinized. Donkey anti-human Cy3-labeled IgG (dilution 1:100; 69901, Jackson ImmunoResearch Laboratories, West Grove, PA) was used for bevacizumab detection. This polyclonal antibody binds to epitopes of both Fc and Fab portions of human IgG. Untreated control eyes were stained in the same way. The sections were then examined with a confocal microscope (Zeiss 510).

Molecular analysis: Corneal tissue was dissected and snapfrozen in liquid nitrogen. Total RNA was isolated using a TRIzol™ reagent (Invitrogen, Life Technologies, Carlsbad, CA), according to the manufacturer's protocol, and then reverse-transcribed into cDNA using random hexamers (Bioline, London, UK) and Moloney murine leukemia virus (M-MLV)-reverse transcriptase (Promega, Madison, WI).

Two-stage real-time quantitative PCR (Sequence Detection System, Prism 7900; Applied Biosystems, Inc., Foster City, CA) was applied to evaluate the genes encoding *VEGF*, *IGF-1*, *MIP-2*, *PEDF*, and beta actin (*ACTB*) after cauterization (for primer list, see Table 2). Mouse β -actin was used to normalize the cDNA input levels. Reactions were performed in a 20 μ l volume containing 4 μ l cDNA, 0.5 μ M each of forward and reverse primers and buffer included in the master mix (SYBR® Green I; Applied Biosystems, Inc.). Duplicate reactions were performed for each gene to minimize individual tube variability, and an average was taken for each time point. Threshold cycle efficiency corrections were calculated, and melting curves were obtained using cDNA for each individual gene PCR assay.

PCR cycling conditions consisted of an initial denaturation step of 95 °C for 10 min followed by 40 cycles of 15 s denaturation at 95 °C and 1 min of annealing and extension at 60 °C. Standard curves were obtained using untreated mouse cDNA for each gene PCR assay. The results were quantified using a comparative threshold cycle (Ct) method, also known as the $2^{-\Delta\Delta Ct}$ method [58], where: $\Delta\Delta Ct = \Delta Ct$ (sample) – ΔCt (reference gene).

For each treatment, the levels of expression in the cauterized right eye were compared to the untreated eye, which served as an internal control.

Statistical analysis: Differences between groups were analyzed using an unpaired Student t-test. Significance was set at p<0.05.

TABLE 3. RELATIVE AREA OF NEOVASCULARIZATION (IN VIVO MEASUREMENT BY IMAGEJ).

	Model of corneal neovascularization		Bevacizumab treate		
	(n=32)		Intraocular in	jection (n=37)	Sub-conjuctival
Time (days)	No treatment	Total (n=57)	Intra-cameral (n=19)	Intra-vitreal (n=18)	injection (n=20)
2	11.24±7.02	10.98±8.79 p=0.88	14.41±2.98 p=0.06	11.05±2.49 p=0.91	7.05±2.88 *p=0.012
4	19.70±8.96	14.76±8.79 *p=0.01	16.51±6.27 p=0.18	14.68±5.05 p=0.03	15.69±6.00 p=0.09
8	47.42±25.45	27.10±21.84 *p=0.0001	26.28±6.53 *p=0.005	23.51±6.51 *p=0.001	32.16±15.21 p=0.04
10	50.62±24.74	28.29±20.92 *p=0.001	19.86±1.23 *p=0.0004	24.20 ± 14.87 p=0.02	39.73±14.51 P=0.27
14	26.98±19.90	10.97±10.84 *p=0.002	5.28±5.90 *p=0.01	4.16±6.43 *p=0.009	23.48±17.32 p=0.68

The asterisk indicates a p<0.05.

RESULTS

Area of neovascularization:

Mouse model—The mean burn area in the center of the cornea measured 10% of the total corneal area in the untreated mice. In all mice subjected to chemical burn, corneal neovascularization started on day 2 and reached its maximum on days 8–10. We calculated the relative area of corneal neovascularization as 11.24% (± 7.0) on day 2, 19.70% (± 8.9) on day 4, 47.42% (± 25.4) on day 8 and 50.62% (± 24.7) on day 10. Spontaneous regression of the neovascularization was detected on day 14, when the calculated area of neovascularization was 26.98% (± 19.9) of the corneal area (Table 3).

In the bevacizumab-treated mice, the relative area of neovascularization in the corneas was lower at all time points than in the untreated, cauterized mice (Table 3). Differences between the groups were significant on days 8, 10 and 14 (p<0.005).

Qualitative rabbit model—In the rabbits (n=6), the mean burn area measured 10% of the corneal area of the untreated eyes. Neovascularization was induced by the chemical burn in all rabbits starting on day 2 and achieved its maximum on day 8. Neovascularization covered less than 20% of the corneas at sacrifice (Figure 1).

The intracameral (anterior chamber) injection of bevacizumab reduced the neovascularization in the rabbits. Moreover, the limbal congestion noted in the untreated model did not develop.

Route of injection: All three injection routes of bevacizumab — intravitreal, intracameral and subconjuctival — reduced growth of new abnormal vessels in the mice (Figure 2H-J). However, intraocular injections were the most effective (p<0.005, 8 and 10 days after injury).

Fluorescein angiography: In the neovascular model, following an intraperitoneal injection of fluorescein, angiography demonstrated new vessels in the cornea on day 8 (maximum growth). The vessels were located at 360 degrees along the periphery cornea, arising from the limbus and extending towards the central scar. The new vessels showed early leakage that increased with time (Figure 2C-E).

Immediately following fluorescein angiography, the eyes were enucleated for flat-mount studies to validate the area of neovascularization (Figure 2F-G). Intraluminal fluorescein dye analysis demonstrated peripheral neovascularization of about 50% of the corneal area at 8 days after cauterization (Figure 2G).

India ink perfusion: Flat-mount corneal studies following perfusion with India ink were conducted 10 days after cauterization. Patent vessels were noted in the corneal periphery covering 50% of the corneal area (Figure 2L). After bevacizumab treatment, the area of neovascularization was significantly reduced (Figure 2M-O).

Histology: Histological sections stained with hematoxylin and eosin showed scarring in the epithelial and anterior stroma of the central corena after cauterization. New pathological blood vessels were located in the superficial stroma, filled with erythrocytes (Figure 3). Neovascularization was detected in both untreated and treated corneas.

Staining for endothelial marker: Blood vessels in the flatmount corneas were stained with CD31, as shown in Figure 4. Staining was negative for endothelial marker in the control avascular cornea.

Staining for intraocular bevacizumab: One day after intravitreal injection of 0.75 mg in 0.03 ml of bevacizumab, staining with anti-human Cy3TM conjugated affine pure

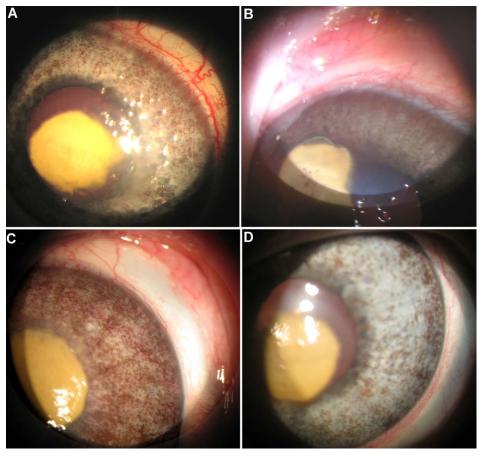


Figure 1. Corneal neovascularization in rabbits. **A-B**: Clinical appearance of neovascularization in a rabbit cornea 10 days (peak) after chemical burn induction. Note the central scar (yellow) covering approximately 10% of the corneal area. **C-D**: Intracameral bevacizumab-treated cornea at 10 days. Note the reduced neovascularization.

fragment donkey anti-human IgG revealed bevacizumab in the anterior chamber, filtrating into the cornea (Figure 5B). No such findings were detected in the negative controls (cauterization without bevacizumab treatment; Figure 5A).

Molecular analysis:

Vascular Endothelial Growth Factor—In the untreated eyes, VEGF expression peaked on day 10 and decreased slightly on day 14, in line with the clinical observations. The bevacizumab-treated eyes showed a slight increase in VEGF expression from day 2 (2.49 fold) to day 14 (3.43 fold) (Table 4, Figure 6A).

Pigment Epithelium Derived Factor—PEDF showed an inverse pattern of expression to VEGF. After untreated cauterization, PEDF gene expression was 0.35 on day 2, with a mild increase toward baseline until day 10; by day 14, expression measured 3.54 fold. With bevacizumab treatment, PEDF levels decreased on day 2 (0.44 fold), followed by a 2-fold increase on days 8 and 10, and a 5.48-fold increase on day 14 (Table 4, Figure 6B).

Insulin-Like Growth Factor-1—In the untreated eyes, IGF-I expression increased on day 2, returned to baseline on day 8 and increased again on day 14 (2.88 fold). With bevacizumab treatment, levels decreased on day 2, significantly increased on day 8 (3.6 fold), peaked on day 10

(5.4-fold) and decreased on day 14 (1.94 fold; Table 4, Figure 6C).

Macrophage-Inflammatory Protein-2—MIP-2 expression significantly increased in both the untreated and treated mice on day 2 and then dropped to near-normal levels by day 14. However, in the treated group, the levels increased to over 300 fold, followed by a reduction to 200 fold. In the untreated group, the maximal level achieved was only 166 fold (Table 4, Figure 6D).

DISCUSSION

The avascular, transparent nature of the cornea makes it possible for researchers to detect and measure new blood vessel development [59]. The present study molecularly and histologically characterized a mice model of corneal neovascularization and measured the effect of bevacizumab treatment on the appearance of new vessels and the changes in angiogenesis-related genes using three different routes of injection: intravitreal, intracameral or subconjuctival.

We found that neovascularization appeared two days after the chemical burn was induced, which then progressed from the limbal vascular plexus toward the cauterization site in the central cornea. Maximal vessel growth was observed 8–10 days after injury, with spontaneous regression thereafter.

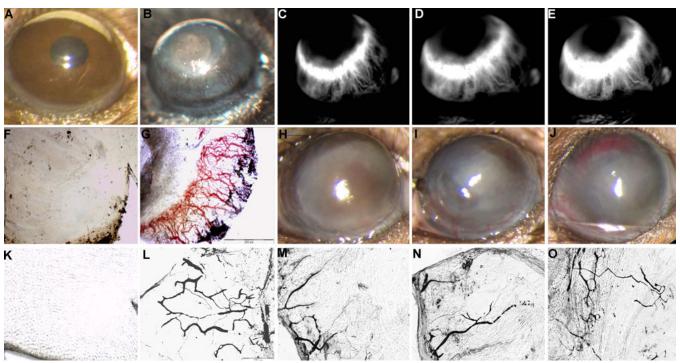


Figure 2. Clinical appearance of neovascularization in a mouse cornea. Comparison between normal eye, without blood vessels on the cornea (A) and 8 days following cauterization without treatment, with high level of neovascularization (B). Less blood vessels observed after bevacizumab treatment by (H) intravitreal injection, (I) intracameral injection or (J) subconjunctival injection at the same time point. Fluorescein angiography 8 days after chemical burn induction showing (C) early leakage and (D-E) increased late leakage. Flat mount cornea of the control mouse showing transparent cornea without blood vessels (F). In the neovascular corneal model, studies with fluorescein dye (G) and India ink (K, black) reveal new vessels 8 days after induction of chemical burn, compared to the control avascular cornea (K), and a reduced level of NV detected following bevacizumab treatment (M-O).

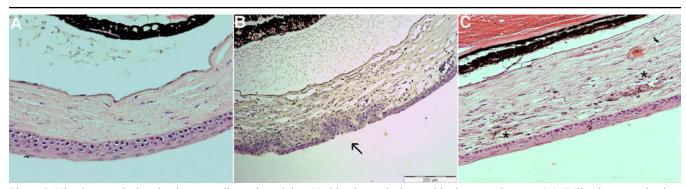


Figure 3. Histology analysis using hematoxylin-eosin staining. No blood vessels detected in the normal cornea (**A**). Following cauterization, development of the scar (arrow) was observed in the epithelial and anterior stroma of the center cornea (**B**). New pathological blood vessels (**C**) were located in the superficial stroma, filled with erythrocytes (*).

These findings are in agreement with other reports of corneal neovascularization in mouse [3], rat [60], rabbit [17] and guinea pig [61] models. On day 10, the maximal area of neovascularization in our study measured 51% of the corneal area, compared to 23.5% in the guinea pig after 10 days [61], and 72% [34], 68% [60] and 63% [20] in rats after one week. The reason for the lower rate in the guinea pig is unclear, though it is probably unrelated to the different corneal size.

In vivo analysis of our model with fluorescein angiography demonstrated early leakage. Analysis with India

ink showed the complex of the vessel tree in the flat-mount cornea. Histologically, the vessels were located in the superficial stroma, resembling the pathology described in human corneas following keratoplasty [62].

A single injection of bevacizumab reduced the corneal neovascularization relative to the untreated, injured mouse model. The effect was greater with the intraocular route (anterior chamber or vitreous) than with the subconjuctival route. The effect of a subconjuctival bevacizumab injection has been previously investigated in rat [63], guinea pig [61]

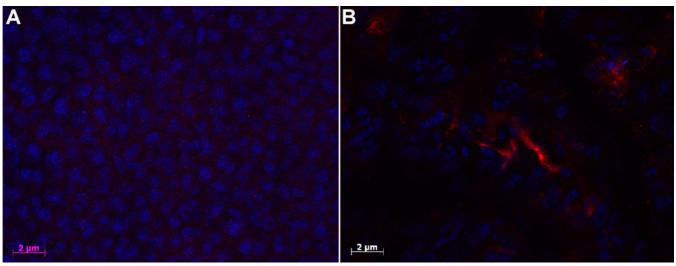


Figure 4. Corneal flat-mount immunostaining for endothelial marker. Flat-mount corneas were stained with endothelial marker using anti-CD31 antibody. No positive staining detected in a control mouse (avascular cornea; **A**) compared to blood vessels detected in the flat mount cornea 8 days after cauterization (red staining; **B**).

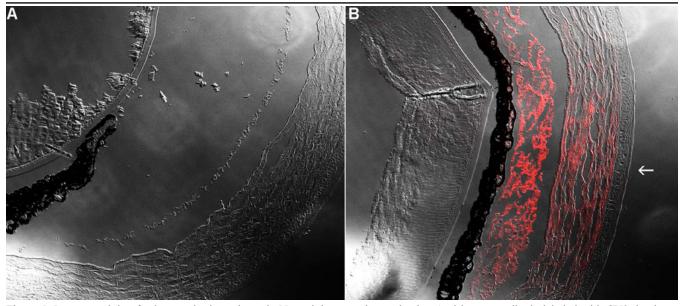


Figure 5. Immunostaining for intraocular bevacizumab. No staining was detected using Anti-human antibody labeled with CY3 dye in an untreated mouse after chemical burn injury (A) without bevacizumab injection. Bevacizumab staining identified in the anterior chamber, filtrating into the cornea, following intraocular injection one day after injury (B).

and rabbit models [21,43,64] and in humans [41]. Hurmeric et al. [61] noted a significant reduction in corneal neovascularization when bevacizumab was injected subconjuctivally immediately after cauterization and again on day 3 after injury. Delayed injections (days 3 and 5) had a significantly weaker effect. The adjuvant injection as well as species differences may account for the superior results of this study compared to ours. In an earlier rabbit model [43], subconjunctival injection of bevacizumab caused significant inhibition of corneal neovascularization.

The intravitreal route is commonly used nowadays, with only minor complications, in patients who require repeated injections for age-related macular degeneration [46], Best disease [65] or macular edema of central retinal vein occlusion [66]. In the present study, intravitreal injection of bevacizumab was associated with a 50% reduction in corneal neovascularization compared to the untreated model (p<0.001; Figure 2). There were no injection-related complications. Surprisingly, intravitreal injection had a better effect on regression of the corneal neovascularization than injections into the anterior chamber, which come into contact with the cornea. However, the difference was not statistically significant. Immunostaining was performed to better understand the location of the intravitreally-injected

Time po	oint 2 (days	8	days	10	days	14 d	lays
Gene	Model	Treated	Model	Treated	Model	Treated	Model	Treated
VEGE	2.98	2.49	3.80	2.96	4.10	3.27	3.64	3.43
	(± 0.04)	(± 1.95)	(± 5.50)	(± 2.78)	(± 3.30)	(± 2.32)	(± 2.41)	(± 1.71)
PEDF	0.35	0.44	0.65	2.0	0.84	1.78	3.54	5.48
	(± 0.48)	(± 0.09)	(± 0.19)	(± 0.7)	(± 0.08)	(± 0.78)	(± 0.37)	(± 3.97)
IGF-1	1.65	0.66	0.90	3.60	1.37	5.41	2.88	1.94
	(± 1.65)	(± 0.45)	(± 0.31)	(± 1.76)	(± 0.85)	(± 4.96)	(± 0.68)	(± 0.89)
MIP-2	166.43	332.53	75.25	197.11	76.25	170.83	11.45	4.63

TABLE 4. GENE EXPRESSION IN THE UNTREATED AND BEVACIZUMAB-TREATED CORNEAS.

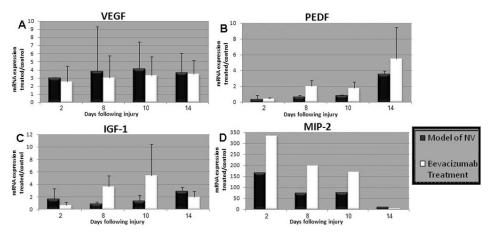


Figure 6. Molecular analysis of gene expression levels after chemical cauterization in the untreated and bevacizumab treated mice. VEGF expression up-regulated in all time points, with a slight decrease following bevacizumab treatment (A). PEDF showed an inverse pattern of expression VEGF (B). IGF-1 expression increased between days 8 and 10 following injury, with a higher level in the bevacizumab treated group (C). MIP-2 expression significantly increased in both the untreated and treated mice on day 2 and then dropped to near-normal levels by day 14, with higher level in the bevacizumab treated eyes (D).

bevacizumab, and it revealed that the agent was in the anterior chamber and penetrating the cornea as early as day 2 (Figure 5). Accordingly, others reported that ranibizumab (Lucentis) could be detected in the aqueous humor three days after intravitreal administration [67]. These authors attributed the drug's effectiveness to the shift from the vitreous to the aqueous humor, and possibly to the presence of the drug in the anterior chamber one day after induction of injury [67]. We assumed that in our study, bevacizumab reached the corneal stroma by passive diffusion. Chen et al. [43] reported positive staining for bevacizumab in the corneal stroma between 3-14 days following injury induction in several different rabbit models of corneal neovascularization. They suggested that bevacizumab may be useful in preventing corneal neovascularization in the acute phase of various kinds of corneal inflammation. In a study of the effectiveness of bevacizumab in a rabbit model of retinal detachment, Dib et al. [68] detected subretinal drug molecules 2 h after intravitreal injection. Given the absence of any retinal holes or tears, they too attributed their findings to a mechanism of diffusion throughout the retinal layers.

Bevacizumab only partially inhibited neovascularization in our mouse model compared to its almost 90% inhibition of

choroidal neovascularization in human eyes. The difference might be species-related, as rat, rabbit and guinea pig models all showed a better response to treatment. Yu et al. [41] found bevacizumab to be species-specific, with weak interaction with murine VEGF relative to human VEGF due to differences in the spatial molecular structure.

Although our qualitative analysis yielded a better response to bevacizumab in rabbits than in mice, we nevertheless found a consistent trend of a decrease in neovascularization following bevacizumab injection by different routes. In the wake of advances in the treatment of eye disease involving pathologic angiogenesis, interest has been directed to developing reliable and reproducible animal models [69]. Models of anterior segment neovascularization include the corneal micropocket assay, which is used to study the influence of specific molecules/proteins in angiogenesis, and suture-induced or corneal chemical injury, as in our study, which more closely mimics the complex nature of human disease [69]. The mouse model of corneal neovascularization is well-recognized as a reliable and valid model of all diseases that involve neovascularization [69]. The cornea is also wellvisualized and easily accessible for qualitative and quantitative measurement. Accordingly, we were able to characterize the relative changes in the angiogenesis-related genes and to identify new aspects of the mechanisms underlying corneal neovascularization in the mice.

VEGF is required for angiogenesis [70]. In our model, there was an upregulation of VEGF mRNA expression in both the untreated and bevacizumab-treated mice, although there was slightly less in the treated group (Table 4). Yan et al. [71] conducted an in vivo study of VEGF and TSP2 expression after corneal alkali burn. They found that the level of VEGF increased at 6 h after injury and reached its maximum at 12 h. The level then increased again to additional peaks at 96 h and 192 h. The VEGF-positive reaction was concentrated mainly in the corneal stroma. This is in accordance with our detection of bevacizumab in the corneal stroma using immunohistochemical staining (Figure 5). In contrast, the expression of PEDF in the untreated mice decreased until the late stages of angiogenesis, and peaked only at 14 days following cauterization during the process of wound healing (Table 4). In the treated mice, the expression of PEDF decreased early and then increased progressively until day 14. These findings reflect the complicated interaction, with many factors involved in the promotion and inhibition of vascularization in vivo [72-74].

IGF-1 is known to induce angiogenesis [75] and may explain the upregulation at the early state of neovascularization process. In our model, IGF-1 was upregulated early in the untreated mice, with higher levels in the bevacizumab-treated group than at all the other later points. A previous study that used an ischemic mouse model also reported an increase in IGF-1 three days following injury [76]. Eight days after cauterization, there was a slight change in IGF-1 expression in the untreated mice, compared to a 3.6-fold increase in the bevacizumab-treated group. We assume that when VEGF was neutralized by bevacizumab, other proangiogenic factors, such as IGF-1, came into play.

MIP-2 is the murine analog of IL-8, a member of the proinflammatory C-X-C family of cytokines that activates the release of neutrophils during inflammation. IL-8 also serves as a proangiogenic factor during corneal neovascularization. Similarly to IGF-1, we found an immediate, early increase in MIP-2 mRNA expression after chemical injury, which was greater in the bevacizumab-treated than in the untreated group. This finding may also be a result of the inhibition of VEGF. This finding is consistent with other studies of molecular expression during angiogenesis [77-79], but is not consistent with the conclusion of Lu et al. [55]. In this study, the authors used a similar murine model of corneal neovascularization and concluded that although MIP-2 mediated neutrophil infiltration to the corneal neovascularization, it made only a minor contribution to its development. A reduction in inflammatory cell infiltration of chemically burned rat corneas following bevacizumab injection was also reported by Oh et al. [40]. However, they

noted a reduction in the expression of various cytokines, namely interleukin-2 and -6 and interferon-gamma. It is noteworthy that in this study, bevacizumab treatment led to only a borderline reduction in corneal neovascularization.

In summary, bevacizumab treatment partially inhibits the progressive corneal neovascularization induced by chemical injury in a mouse model. Treatment via the intraocular route is more effective than the subconjuctival (periocular) route. The clinical findings are compatible with the angiographic and histologic findings, and are supported by molecular analysis showing changes in the expression of proangiogenesis genes after treatment. Further studies of the molecular mechanisms involved in neovascularization are needed in the animal models used in this study and in other animal models as well. The ability to directly block target proteins or peptides or signaling pathways holds significant promise for the development of new, targeted anti-angiogenic therapies as well as for the optimization of existing anti-angiogenic drugs polypeptides [11].

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