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Angiogenesis Is Not Impaired in Connective Tissue Growth Factor (CTGF) Knock-out Mice

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SUMMARY Connective tissue growth factor (CTGF) is a member of the CCN family of growth factors. CTGF is important in scarring, wound healing, and fibrosis. It has also been implicated to play a role in angiogenesis, in addition to vascular endothelial growth factor (VEGF). In the eye, angiogenesis and subsequent fibrosis are the main causes of blindness in conditions such as diabetic retinopathy. We have applied three different models of angiogenesis to homozygous CTGF^{-/-} and heterozygous CTGF^{+/-} mice to establish involvement of CTGF in neovascularization. CTGF^{-/-} mice die around birth. Therefore, embryonic CTGF^{-/-}, $CTGF^{+/+}$, and $CTGF^{+/+}$ bone explants were used to study in vitro angiogenesis, and neonatal and mature CTGF^{+/-} and CTGF^{+/+} mice were used in models of oxygen-induced retinopathy and laser-induced choroidal neovascularization. Angiogenesis in vitro was independent of the CTGF genotype in both the presence and the absence of VEGF. Oxygen-induced vascular pathology in the retina, as determined semi-quantitatively, and laser-induced choroidal neovascularization, as determined quantitatively, were also not affected by the CTGF genotype. Our data show that downregulation of CTGF levels does not affect neovascularization, indicating distinct roles of VEGF and CTGF in angiogenesis and fibrosis in eye conditions. (J Histochem Cytochem 55:1139–1147, 2007)

KEY WORDS CTGF

VEGF diabetic retinopathy angiogenesis fibrosis angio-fibrotic switch vitreous

CONNECTIVE TISSUE GROWTH FACTOR (CTGF), a member of the CCN family of growth factors, is known primarily for its pro-fibrotic properties and its potential as a target for therapy in conditions with pathological fibrosis (Franklin 1997; Goldschmeding et al. 2000; Perbal 2004). Anti-CTGF antibodies have been tested in patients with idiopathic pulmonary fibrosis in a Phase 1 clinical trial (Mageto et al. 2004;Van Nieuwenhoven et al. 2005). In addition to its known

pro-fibrotic properties, CTGF has also been reported to induce angiogenesis after exogenous administration in vivo in the cornea and other models (Babic et al. 1999; Shimo et al. 1999). Moreover, CTGF and vascular endothelial growth factor (VEGF) (Witmer et al. 2003) colocalize in vascular cells in human choroidal neovascular membranes (Watanabe et al. 2005), and levels of CTGF are increased in the vitreous of patients with active proliferative diabetic retinopathy (PDR) (Hinton et al. 2004). These findings suggest that CTGF has a causal role in ocular neovascularization as well.

However, combined exogenous administration of CTGF and VEGF in the back of a mouse or in a mouse model of hindlimb ischemia inhibits VEGF-induced angiogenesis because of the binding of VEGF by CTGF

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Received for publication April 12, 2007; accepted June 25, 2007 [DOI: 10.1369/jhc.7A7258.2007].

(Inoki et al. 2002; Jang et al. 2004). When CTGF is upregulated by VEGF (Suzuma et al. 2000; He et al. 2003), it can reduce the bioavailability of VEGF by VEGF–CTGF binding. Furthermore, we recently demonstrated a strong and highly significant correlation of human vitreous CTGF levels with degree of fibrosis, but not with neovascularization, in a large series of patients with various vitreo-retinal disorders, including PDR (Kuiper et al. 2006).

On the basis of the latter findings, we want to challenge the concept that CTGF is an intrinsic ocular angiogenesis factor. We hypothesize that in the eye, CTGF is primarily a pro-fibrotic factor and that CTGF is not essential for angiogenesis.

To test our hypothesis, we investigated the role of CTGF in three angiogenesis models in transgenic mice with absolute or relative CTGF deficiency $(CTGF^{-1}$ or CTGF^{+/-}, respectively). In these mice, we investigated whether CTGF is necessary for spontaneous and VEGFinduced angiogenesis in embryonal bone explants in vitro (Deckers et al. 2001; Van der Pluijm et al. 2003), in oxygen-induced retinopathy in neonatal mice (Agostini et al. 2005), and in laser-induced choroidal neovascularization in adult mice (Lambert et al. 2001,2003).

Materials and Methods

Genetically Modified Mice

Male BalbC/129sv $CTGF^{+/-}$ mice (Ivkovic et al. 2003) were crossbred with $CTGF^{+/-}$ female C57Bl/6J mice (Harlan; Horst, The Netherlands). Seventeen-day-old embryos $(CTGF^{+/-})$, $CTGF^{+/+}$, and $CTGF^{-/-}$) were removed from pregnant $CTGF^{+/-}$ mice, and metatarsals were dissected as described previously (Deckers et al. 2001). In addition, male BalbC/129sv $CTGF^{+/-}$ mice were crossbred with $CTGF^{+/+}$ female C57Bl/6J mice (Harlan). The offspring ($CTGF^{+/-}$ and $CTGF^{+/+}$ mice) were used for the ocular angiogenesis studies. Homozygous knock-outs ($CTGF^{-/-}$) were not available for these experiments, because these mice are not viable after birth (Ivkovic et al. 2003).

All mice were genotyped using PCR. Animal experiments were performed in compliance with the Association for Research in Vision and Ophthalmology Statement for the Use of Animals in Ophthalmic and Vision Research. The animals were maintained under a 12:12-hr light/dark cycle and had free access to food and water.

In Vitro Angiogenesis Assay of Embryonal Mouse Metatarsals

Metatarsals were dissected, cultured, and stained as previously described (Deckers et al. 2001; Van der Pluijm et al. 2003). Briefly, four left metatarsals and four right metatarsals of each of 17 $CTGF^{+/-}$, 4 $CTGF^{+/+}$, and 2 $CTGF^{-/-}$ embryos were isolated and separately cultured in 24-well plates for 14 days. The experiment was conducted in triplicate (metatarsals of three breeds). After 72 hr of culture, the 10% fetal calf serum–containing medium was replaced by serum-free

medium with (right metatarsals) or without (left metatarsals) 50 ng/ml recombinant human VEGF-A (Oncogene; Sanbio, Uden, The Netherlands). At day 14, the cultures were fixed in zinc macrodex formalin (ZnMF) fixative containing 0.1 mmol/liter Tris-acetate, pH 4.5, 0.5% $ZnCl₂$, 0.5% Zn acetate, 5% dextran, and 3.8% paraformaldehyde for 15 min and subsequently stained using a rat anti-mouse monoclonal antibody against an endothelial cell marker [CD31, PECAM-1, or ER-MP12; kindly provided by Pieter J.M. Leenen (Slieker et al. 1993)]. Following this, metatarsals were washed three times in PBS and then incubated with 40% methanol/1% $H₂O₂$ in PBS for 30 min at room temperature to abolish endogenous peroxidase activity. After three washes with PBS, nonspecific staining was blocked by incubation in a solution of 0.3% skimmed milk powder (Boehringer; Mannheim, Germany) in TTBS (50 mmol/liter Tris-HCl, pH 7.4, 150 mmol/liter NaCl, and 0.05% Tween 20) for 1 hr at 37C, followed by incubation with the ER-MP12 antibody (dilution, 1:33) overnight at 4C.

The following day, after three washes in 100 mmol/liter Tris-HCl, pH 7.4, 150 mmol/liter NaCl, and 0.05% Tween 20, cultures were incubated with biotinylated sheep anti-rat secondary antibody (dilution, 1:300; Amersham, Buckinghamshire, UK) for 45 min and streptavidin-horseradish peroxidase complex for 30 min at 37C. Peroxidase activity was visualized using its chromagenic substrate 3-amino-9-ethyl carbazole (Sigma; St. Louis, MO).

Quantification of CD31-positive Tubular Structures. Digitized images were acquired using a digital camera (Nikon; Melville, NY). Captured images were exported as JPEG files and further processed using Powerpoint software. Total area of CD31-positive tubular structures per metatarsal was determined by computerized image analysis using Image-Proplus software (Media Cybernetics; Silver Spring, CA).

Mouse Model of Oxygen-induced Retinopathy

Retinopathy was induced as previously described (Agostini et al. 2005). Briefly, 7-day-old litters and the mother were kept for 5 days in a sealed container ventilated by a mixture of oxygen and compressed air with a final oxygen concentration of 76 \pm 2% (Oxygen Monitor 180, Kontron Instruments, Bletchley, UK; sensor, Linde Heimox; Unterschleissheim, Germany). Afterwards, animals were kept under normal conditions for 6 days. On postnatal day 18, the litters were deeply anesthetized by inhalation of 2.5–3.5% isoflurane mixed with oxygen, with a flow of 2 liters/min in a vaporizer (Vapor 19.3; Dräger, Lubeck, Germany). Retinal vessels of the litters were visualized by perfusion with fluorescein isothiocyanate (FITC) coupled to high-molecular-weight dextran (FD 2000 S; Sigma-Aldrich, Steinheim, Germany) dissolved in 0.9% NaCl at a concentration of 50 mg/ml. Perfusion was performed by injecting 1 ml of the fluorescein solution into the right atrium with a 30-gauge cannula, after carefully opening the thorax. Three to 4 min after perfusion, the flow of isoflurane was increased and the mice were sacrificed. Eyes were enucleated, and pieces of the tail were collected for PCR genotyping.

After enucleation, the eyeballs were fixed for 4 hr in 4% buffered formaldehyde at room temperature. After removal of the anterior segment, retinas were mounted on siliFigure 1 Retinal flat mount of mouse eye with oxygen-induced retinopathy showing blood vessels after perfusion with fluorescein isothiocyanate (FITC) coupled to high-molecular-weight dextran. The retina was divided into three concentric zones: the inner zone around the optic disc (A), the middle zone (B), and the outer zone (C) for scoring of avascular areas on a central–peripheral axis. Vascular proliferations were quantified by counting blood vessel tufts and cluster formation (presumed extra-retinal neovascularization) in each of 12 equally sized sections of the retina (numbers). $Bar = 1$ mm.

conized microscope slides (SuperFrost Plus; Menzler-Gläzer, Braunschweig, Germany) using Vectashield mounting medium (Vector; Burlingame, CA), an anti-fading agent, and 15 mmol/liter NaN₃ (Dako; Glostrup, Denmark). Retinal whole mounts were covered by a coverslip and kept in the dark at 4C until analysis.

Quantification of Retinal Vascular Changes. Retinal whole mounts were analyzed using a standard fluorescence microscope. Slides were masked to avoid bias. Whole mounts that had been incompletely perfused were excluded from analysis. Based on a scoring system described by Higgins and colleagues (Higgins et al. 1999; Yossuck et al. 2000), retinas were divided into three concentric zones, with the papilla in

the center (Figure 1). This system takes different aspects of vascular morphology into account; these are based on funduscopic features of proliferative retinopathy. Phenomena such as central and peripheral avascularity, blood vessel tufts, extra-retinal neovascularization, and blood vessel tortuosity, were scored in the three retinal zones (Table 1). The higher the score (range 0–13), the more severe the retinopathy.

Mouse Model of Laser-induced Choroidal Neovascularization

Choroidal neovascularization was induced in mice by laser burns as previously described (Lambert et al. 2001,2003). Briefly, mice were anesthetized with an intraperitoneal in-

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jection of Avertin (Aldrich; Milwaukee WI). Both pupils were dilated with topical administration of 0.5% tropicamide (0.5% Tropicol; Bournonville Pharma, The Hague, The Netherlands) and four burns were delivered at the 3, 6, 9, and 12 o'clock positions around the optic disc, using a green argon laser $(532 \text{ nm}; 50\text{-}\mu\text{m})$ diameter of the spot size; 0.05-sec duration; 400 mW; Figure 2) and a coverslip as contact lens. Animals were sacrificed after 14 days.

To quantify choroidal neovascularization, mice were injected intravenously with $200 \mu l$ of 50 mg/ml FITCconjugated dextran (Sigma) in PBS (pH 7.4). Immediately after, one eye per mouse was fixed in 1% paraformaldehyde, pH 7.4, for 60 min at room temperature. Retinas were removed, and the choroid was mounted flatly for confocal microscopy using Vectashield mounting medium. The other eye was fixed in buffered 3.5% paraformaldehyde and frozen in liquid nitrogen for cryostat sectioning.

Quantification of Choroidal Neovascularization. Choroidal neovascularization was quantified as previously described (Lambert et al. 2001,2003). Briefly, serial sections were cut throughout the entire area of each burn using a cryostat (Leica; Meyer Instruments, Houston, TX), and the thickest lesions (at least five sections per lesion) were selected for quantification. With the use of image analysis (Micro Image version 3.0 for Windows 95/NT; Olympus, Hamburg, Germany), neovascularization was estimated by the ratio (B/C) of the thickness from the bottom of the pigmented choroidal layer to the top of the neovascular membrane (B) to the thickness of the intact pigmented choroid adjacent to the lesion (C) (Figure 3). A mean value of the B/C ratio was attributed to each laser impact.

In addition, choroidal neovascularization was also quantified by confocal visualization of flat-mount choroids. The retinas were removed, and the choroid was mounted flatly for confocal microscopy using anti-fading Vectashield mounting medium. The spatial distribution of FITC fluorescence was examined using a Leica TCS SP2 inverted confocal laser microscope (Leica Microsystems; Mannheim, Germany) equipped with an argon laser and an acousto-optical tunable filter for excitation intensity. Digitized images were acquired using a $10\times$ [numerical aperture (NA) 0.4] objective or a 63 \times (NA 1.2) Plan-Apo water-immersion objective at $1024 \times$ 1024 pixel resolution. FITC was visualized by using an excitation wavelength of 488 nm. Emission light was recorded at 500–555 nm. For each lesion, serial optical sections were

ser photocoagulation.

Figure 3 Estimation of choroidal neovascularization after induction by laser photocoagulation in hematoxylin-stained cryostat sections of the area containing the entire extent of the burn in mouse eyes. Neovascularization is determined by the ratio of the thickness of the lesion (B) to the adjacent intact pigmented choroidal layer (C).

recorded with a z-step of $1.67 \mu m$. After successive scanning for each interval, three-dimensional images of fluorescence were reconstructed by using Leica confocal software. Images were acquired under identical conditions, and we ensured that the maximal fluorescence signal was not saturating the photomultiplier tubes. Captured images were exported as TIFF files and processed further using Photoshop software (Adobe; San Jose, CA). Quantification of the confocal images was realized by measurement of the surface areas of fluorescence (Scion Image for Windows beta 4.0.2; Scion, Frederick, MD).

Statistics

Differences between metatarsals incubated in the presence or absence of VEGF were calculated using the nonparametric paired Wilcoxon signed rank test. Data were corrected for age differences between breeds as determined by differences in degree of calcification of the bones and for the fact that part of the data were obtained from metatarsals of one mouse (repeated measurements). Thus, results for $CTGF^{+/+}$, $CTGF^{+/-}$ and $CTGF^{-/-}$ mice (genotype effects and effects of VEGF) were calculated using a repeated measurement model with metatarsal as repeated subject with correction for degree of calcification. Results were considered significantly different at $p<0.05$.

Scores of retinal vascular changes in retinal whole mounts were not normally distributed and therefore were expressed as median and range. Differences between scores of retinal vascular changes and choroidal neovascularization in $CTGF^{+/+}$ and $CTGF^{+/-}$ mice were statistically analyzed using the nonparametric Mann-Whitney test with $p=0.05$ as level of significance.

Results

In Vitro Angiogenesis Assay

Figure 2 Induction of choroidal neovascularization in mice by la- In total, 92 metatarsals of 4 $CTGF^{+/+}$, 17 $CTGF^{+/-}$, ser photocoagulation.

presence or absence of VEGF, and each of these metatarsals demonstrated outgrowth of vessels (Figure 4). Owing to failure of attachment of some of the metatarsals, 15 out of 16 CTGF^{+/+}, 60 out of 68 CTGF^{+/-}, and 7 out of 8 $CTGF^{-/-}$ metatarsals incubated in the absence of VEGF, and 15 out of 16 $CTGF^{+/+}$, 50 out of 68 CTGF^{+/-}, and 5 out of 8 CTGF^{-/-} metatarsals incubated in the presence of VEGF were analyzed. Metatarsals of CTGF^{+/+}, CTGF^{+/-}, and CTGF^{-/-} embryos showed similar outgrowth of vessels. A genotype effect could not be detected, whether treated with VEGF or not (treated group, $p=0.50$; untreated group, $p=0.76$; Figure 5). The median area of vascular outgrowth of metatarsals of $CTGF^{+/+}$ and $CTGF^{+/-}$ embryos cultured in the presence of VEGF was significantly larger than that of those cultured in the absence of VEGF ($p \le 0.001$). The CTGF^{-/-} metatarsals also showed a larger median area of vascular outgrowth under VEGF stimulation. However, this difference was not statistically significant ($p=0.225$; Figure 5).

Oxygen-induced Retinopathy

In total, 13 CTGF^{+/+} mice (26 eyes) and 8 CTGF^{+/-} mice (16 eyes) were analyzed by FITC–dextran perfusion to examine the retinal response after oxygen exposure (Figure 1). Seven eyes of $CTGF^{+/+}$ mice and one eye of a $CTGF^{+/-}$ mouse were excluded from analysis because of incomplete perfusion. In total, 19 $CTGF^{+/+}$ and 15 $CTGF^{+/-}$ retinal whole mounts were analyzed using the scoring system shown in Table 1. There was no difference in median score of the retinal vascular changes between the $CTGF^{+/+}$ retinal whole mounts (5.0; range, 1–9) and the CTGF^{+/-} whole mounts (6.0; range, 4–10; Mann Whitney, $p=0.107$; Figure 6).

Laser-induced Choroidal Neovascularization

In total, four CTGF^{+/+} mice and six CTGF^{+/-} mice received four laser burns per eye. Twelve out of the 16 spots in the eyes of the $CTGF^{+/+}$ mice and 20 of the 24 spots in the eyes of the $CTGF^{+/-}$ mice induced neo-

Figure 4 Vascular outgrowth of metatarsals as stained immunohistochemically with anti-CD31 antibodies after being cultured in the absence (A,C,E) or presence (B,D,F) of vascular endothelial growth factor (VEGF). (**A,B**) connective tissue
growth factor (CTGF)^{+/+} metatarsals; (**C,D**) $CTGF^{+/-}$ metatarsals; (E,F) CTGF^{$-/-$} metatarsals. Vascular outgrowth is enhanced by VEGF independently of the CTGF genotype. Bar $= 1$ mm.

Figure 5 Graphic representation of the differences in median area of vascular outgrowth (expressed as numbers of pixels \times 1000) between the metatarsals cultured with and without VEGF for CTGF $^{+/+}$, CTGF^{+/-}, and CTGF^{-/-} mice. Data are expressed as median with range. The box indicates values from the 25th to the 75th quartile.

vascularization and were included in the analysis of the B/C ratios in cryostat sections.

Figure 7 shows confocal images of FITC–dextranperfused mouse retinas. Only 8 of 16 choroidal impact areas of $CTGF^{+/+}$ mice and 11 of 24 choroidal impact

Figure 6 Median scores with range of vascular pathology in retinal
whole mounts of CTGF^{+/+} and CTGF^{+/-} mice in the oxygen-induced retinopathy model. There is no significant difference in neovascular response between $CTGF^{+/+}$ and $CTGF^{+/-}$ mice.

areas of $CTGF^{+/-}$ mice could be included for quantitative analysis. Background fluorescence was too high in the areas of the other impacts. Neither quantification method showed statistical differences between experimental choroidal neovascularization in $CTGF^{+/+}$ and CTGF^{+/-} mice (Mann Whitney, $p=0.39$ in Figure 8A and $p=0.85$ in Figure 8B, respectively).

Discussion

The observations in the mouse models in the present study are in line with our previous findings in patients that intravitreal CTGF levels do not correlate with degree of neovascularization (Kuiper et al. 2006). Vascular outgrowth from embryonic metatarsal explants in vitro of homozygous $CTGF^{-/-}$ knock-out mice did not differ significantly from vascular outgrowth from metatarsals of wild-type $CTGF^{+/+}$ mice or heterozygous $CTGF^{+/-}$ mice, in both the presence and absence of VEGF. However, the large variation in all experimental groups that were incubated in the presence of VEGF may hide a different response in the three genotypes. Nevertheless, our data indicate that CTGF is not required for (VEGF-induced) angiogenesis in this model. Our findings do not exclude that CTGF on its own and/ or excessive levels of (exogenous) CTGF may induce angiogenesis, but redundancy of CTGF in ocular angiogenesis in vivo is also suggested by the lack of significant differences between heterozygous mice and wild-type mice in the two ocular angiogenesis models. Heterozygous $CTGF^{+/-}$ mice have been identified as a valid model of decreased CTGF availability, because they express \sim 50% lower protein levels of CTGF (Kuiper 2006; Roestenberg 2007). Furthermore, these mice fail to develop a thickened glomerular basement membrane during experimental diabetes mellitus (Roestenberg 2007).

The fact that angiogenesis is not impaired in the CTGF knock-out mouse is novel and surprising. Previous studies have shown that CTGF is sufficient to induce angiogenesis, directly or indirectly, under experimental conditions in vitro and in vivo (Aiello et al. 1998; Babic et al. 1999; Ferrara 1999a,b; Shimo et al. 1999; Campochiaro and Hackett 2003). In these studies, exogenous CTGF induced angiogenesis in rat cornea (Babic et al. 1999), in chicken chorio-allantoic membrane, and in rCTGF-containing implants in the backs of mice (Shimo et al. 1999). Our findings in human PDR (Kuiper et al. 2006) and in three distinct angiogenic mouse models as described here question the applicability of this concept to ocular disease. The ocular models used in our study mimic pathological conditions occurring in the human eye. In the oxygeninduced retinopathy model, preretinal angiogenesis is caused by retinal hypoxia and VEGF overexpression, a situation similar to retinopathy of prematurity and

Figure 7 Confocal images of FITC–dextran-perfused retinas showing the vasculature in a CTGF^{+/+} (A) and a CTGF^{+/-} mouse (B). Bar = 80 μ m.

PDR (Witmer et al. 2003). In the laser-induced model of choroidal neovascularization, the angiogenic response is part of a wound-healing response after disruption of Bruch's membrane, a situation similar to myopic and agerelated macular degeneration. Our data suggest that as far as the final angiogenic response is concerned, CTGF is a dispensable factor in the complex interplay of hypoxic signaling and VEGF- and wound-healing–driven angiogenesis. Our previous studies indicated that CTGF plays a role in the later stages of PDR, after neovascularization, and induces subsequent fibrosis (Kuiper et al. 2006). CTGF can be induced by increased VEGF (Suzuma et al. 2000; He et al. 2003) and/or advanced glycation end products (Hughes et al. 2007) that are the result of diabetic conditions.

The literature supports this concept. In other organs, the role of CTGF as a pro-fibrotic factor is wellestablished (Franklin 1997; Grotendorst 1997; Duncan et al. 1999; Mori et al. 1999; Goldschmeding et al. 2000; Yokoi et al. 2001,2004; Perbal 2004; Van Nieuwenhoven et al. 2005). Furthermore, it has been shown recently that hypoxic induction of CTGF is directly mediated by the transcription factor hypoxiainducible-factor-1 (HIF-1; Higgins et al. 2004), whereas CTGF inhibits VEGF-dependent angiogenesis by promoting proteasomal HIF-1 α degradation in tumors (Chang et al. 2006; Smith et al. 2006; Tosetti et al. 2006). This shows that the interplay between VEGF and CTGF in angiogenesis and fibrosis is complex, but evidence is accumulating that CTGF does not induce angiogenesis.

Figure 8 Induction of choroidal neovascularization by laser in retina of CTGF^{+/+} and CTGF^{+/-} mice. There is no difference between the two groups as determined by B/C ratio (A) and surface area of choroidal neovascularization, expressed as μ m² \times 1000 (B).

In the eye, CTGF (mRNA and protein) is upregulated by VEGF in retinal endothelial cells (Suzuma et al. 2000; He et al. 2003) and colocalizes with VEGF in neovascular subretinal membranes of age-related macular degeneration patients (Watanabe et al. 2005). In these membranes, CTGF is also found in retinal pigment epithelial cells (Watanabe et al. 2005), which can transdifferentiate into myofibroblasts, the main cell type driving fibrosis. CTGF expression has also been found in myofibroblasts in fibrovascular membranes of patients with PDR (Watanabe et al. 2005) and in pericytes in the retina of PDR patients (Kuiper et al. 2004). Pericytes and myofibroblasts are closely related cell types.

In summary, our data, obtained under various experimental conditions in vivo and in vitro, show that CTGF gene expression, or the level of expression, is not a critical determinant for angiogenesis induced by VEGF, hypoxia, or tissue damage. However, our results do not exclude that CTGF by itself can induce angiogenesis under certain conditions. These data and those from our previous studies suggest that CTGF drives fibrosis rather than angiogenesis in pathological eye conditions.

Acknowledgments

This study was supported by a grant from the Diabetes Fonds Nederland (Grant 2001.042), the Edmond and Marianne Blaauwfonds, and by Les Amis des Aveugles (Ghlin).

The authors thank Lotte Wieten (Department of Pathology, University Medical Center, Utrecht, The Netherlands) for technical assistance, Patrick Motte (Unité de Biologie Cellulaire Végétale, Department of Life Sciences, University of Liege, Belgium) for the confocal imaging, and Dr. Pieter J.M. Leenen (Department of Immunology, Erasmus University, Rotterdam, The Netherlands) for the anti-mouse endothelium antibody.

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