Pigment epithelium-derived factor downregulates vascular endothelial growth factor (VEGF) expression and inhibits VEGF–VEGF receptor 2 binding in diabetic retinopathy

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Abstract

It has been shown that the balance between vascular endothelial growth factor (VEGF), a major angiogenic stimulator, and pigment epithelium-derived factor (PEDF), a potent angiogenic inhibitor, is critical for the regulation of vascular permeability and angiogenesis. However, the regulation of the balance is largely unclear. The present study demonstrated that there is a reciprocal interaction between VEGF and PEDF in the retina. PEDF significantly decreased VEGF expression in both retinal capillary endothelial cells (RCEC) and Müller cells. This PEDF effect was confirmed in the retina of rats with oxygen-induced retinopathy. Silencing of the PEDF gene by siRNA in Müller cells resulted in a significant upregulation of VEGF expression at both the RNA and protein levels, suggesting that PEDF is an endogenous negative regulator of VEGF. The further study of the mechanism showed that PEDF inhibited hypoxia-induced increases in VEGF promoter activity, HIF-1 nuclear translocation and mitogen activated protein kinase phosphorylation. These results suggest that PEDF inhibits VEGF expression at the transcriptional level. In addition, PEDF effectively inhibited VEGF binding to RCEC. Moreover, in vitro receptor-binding assay demonstrated that PEDF competed with VEGF for binding to VEGF receptor 2, which may represent a new mechanism for PEDF activity. On the other hand, VEGF significantly downregulated PEDF expression in RCEC, but not in retinal Müller cells, suggesting a VEGF receptormediated process. These results suggest that the reciprocal regulation between VEGF and PEDF may play a role in angiogenic control. The decrease in PEDF levels in the retina is at least partially responsible for the increase in VEGF expression and subsequent vascular leakage and neovascularization in diabetes.

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Introduction

The blood-retinal barrier (BRB) breakdown and retinal neovascularization (NV) are the major retinal vascular dysfunctions leading to vision loss in a variety of retinal diseases such as diabetic retinopathy (DR) (Bresnick 1986, Frank 1991, Aiello et al. 1997, Pelzek & Lim 2002). Although the exact pathogenic mechanisms are uncertain, it is widely accepted that the two counterbalancing systems, angiogenic stimulators such as vascular endothelial growth factor (VEGF) and angiogenic inhibitors such as angiostatin and pigment epithelium-derived factor (PEDF), play important roles in the regulation of vascular functions (Pierce et al. 1995, Dawson et al. 1999, Gao et al. 2001, Gao et al. 2002a). Under normal conditions, there is a balance between these two systems, which is essential for maintaining the quiescence of retinal vasculature and integrity of the BRB (Bussolino et al. 1997, Miller et al. 1997). Under some pathological conditions, such as DR, this balance is disturbed due to the overproduction of angiogenic stimulators and the

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subsequently lead to the BRB breakdown and the overproliferation of capillary endothelial cells, resulting in retinal NV (Forsythe *et al.* 1996, Miller *et al.* 1997, Gao *et al.* 2001, Gao & Ma 2002). VEGF is a major angiogenic factor, playing a crucial

decreased production of angiogenic inhibitors, which

role in normal and pathological angiogenesis (Pe'er et al. 1995, Aiello 1996, Aiello & Wong 2000, Dvorak 2000). VEGF is also referred to as vascular permeability factor (VPF) based on its ability to induce vascular hyper-permeability (Dvorak et al. 1995). In the early stage of streptozotocin (STZ)-induced diabetes, significant increases in retinal VEGF levels have been found to correlate with retinal vascular permeability (Antonetti et al. 1998). This early BRB breakdown can be successfully prevented by VEGF TrapA₄₀, a soluble VEGF receptor fms-like tyrosine kinase $(Flt)/F_c$ chimera (Qaum et al. 2001). In the oxygen-induced retinopathy (OIR), a commonly used model for proliferative diabetic retinopathy (PDR), retinal VEGF levels are significantly elevated, which correlates with the retinal NV progression (Gao et al. 2002b). The VEGF

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levels decline to the normal level when the regression of NV occurs (Gao *et al.* 2002*b*). These findings suggest that VEGF is a key mediator of both vascular hyperpermeability and retinal NV.

PEDF is a neurotrophic factor and a potent angiogenic inhibitor, originally identified from cultured retinal pigment epithelial (RPE) cells (Tombran-Tink et al. 1991, Dawson et al. 1999, Tombran-Tink & Barnstable 2003a). Previously, we have demonstrated that, in OIR rats, PEDF levels in the retina are significantly decreased in contrast to the increased VEGF levels, which leads to an increased retinal VEGF/ PEDF ratio (Gao et al. 2001). This disturbed balance correlates with the formation and progression of retinal NV (Gao et al. 2001, 2002b). The decrease of PEDF is also observed in the vitreous and aqueous humor of the patients with active DR and diabetic macular edema (DME) (Ogata et al. 2001b, 2002, Boehm et al. 2003). After successful laser treatment, the PEDF levels in the vitreous were elevated, correlating with the regression of retinal NV (Ogata et al. 2001a).

Although the functions of PEDF and VEGF have been well studied in the eye, the interactions between these two counteracting angiogenic factors are largely unclear. Based on the solid evidence showing the inverse correlation between PEDF and VEGF levels in a variety of vascular diseases, we hypothesize that there exists a reciprocal regulation between these two major angiogenic regulators (Gao *et al.* 2001, Ohno-Matsui *et al.* 2001, Doll *et al.* 2003, Kim *et al.* 2003). In this study, we tested this hypothesis and explored the mechanisms by which PEDF regulates VEGF in retinal cells.

Materials and methods

Experimental animals

Brown Norway (BN) rats were purchased from Charles River Laboratories (Wilmington, MA, USA). Care, use and treatment of all animals in this study were in strict agreement with the guidelines in the Care and Use of Laboratory Animals set forth by the University of Oklahoma.

Experimental reagents

Recombinant human PEDF was purchased from BioProducts MD, LLC (Middletown, MD, USA). Recombinant human VEGF₁₆₅ was purchased from PeproTech Inc. (Rocky Hill, NJ, USA). CoCl₂ and BSA were purchased from Sigma.

Development of OIR rat model

OIR rat model was established as described previously (Gao et al. 2001, Zhang et al. 2004, 2005b). Briefly,

newborn BN rats were exposed to 75% oxygen at postnatal day 7 (P7) for 5 days and returned to room air to develop retinopathy. The intravitreal injection was given at P14 and the eyeball was enucleated at P16.

Cell culture

Primary bovine retinal capillary endothelial cells (RCEC) were isolated and cultured in DMEM supplied with 10% human serum as described previously (Gao *et al.* 2003). A rat Müller cell line (rMC-1) was also used in the experiments. After reaching 80% confluence, cells were exposed to serum-free medium for 12 h before the treatments with desired reagents.

Western blot analysis of VEGF and PEDF

Western blot analysis was performed as described previously (Gao *et al.* 2002*a*). Briefly, 50 µg total protein from each sample were blotted with primary antibodies: anti-VEGF (1:500, Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA) and anti-PEDF (1:1000, Chemicon Inc., Temecula, CA, USA). The same membrane was stripped and reblotted with an anti- β -actin antibody for normalization.

Quantification of VEGF and PEDF by ELISA

The protein concentration was measured with the BioRad DC protein assay. The concentrations of VEGF and PEDF in the cell culture medium were measured using ELISA kits specific for VEGF (R&D Systems, Minneapolis, MN, USA) and PEDF (Chemicon Inc.) according to the manufacturers' instructions.

siRNA transfection in rat retinal Müller cells

PEDF siRNA was synthesized using the silencer siRNA construction kit (Ambion, Austin, TX, USA). The following DNA templates were used to synthesize double-stranded PEDF, siRNA: sense, GGA UUU CUA CUU GGA UGA ATT, and anti-sense, UUC AUC CAA GUA GAA AUC. Transfection was performed with the siPORT lipid transfection reagent (Ambion) following the manufacturer's instructions.

Semi-quantitative RT-PCR analysis

Total RNA was extracted using the RNeasy miniisolation kit (Qiagen). PCR was performed as described previously (Shao *et al.* 2004), using PEDF-specific primers: forward primer, 5'-GTC ACT GCC CGC CAG AAG AAC-3', and reverse primer, 5'-TGT TAC CCA CTG CCC CTT GAA-3'. Specific PCR products were examined by electrophoresis in 1% agarose gel.

Real time RT-PCR analysis

Primers specific for VEGF (VEGF-F, 5'-ggccaactacaacttcttcc-3', VEGF-R, 5'-ccatcatggattcgagaaaa-3') were used for real-time RT-PCR. The PCR was performed using GeneAmp RNA PCR kit and SYBR Green PCR Master Mix (Applied Biosystems, Foster City, CA, USA). The average $C_{\rm T}$ (threshold cycle) of fluorescence units was used to analyze the mRNA levels. The VEGF mRNA levels were normalized by 18 S ribosomal RNA levels. Quantification was calculated as: mRNA levels (percent of control) = $2^{\Delta(C_{\rm T})}$ with $\Delta C_{\rm T} = C_{\rm T, VEGF} - C_{\rm T, 18S RNA}$ and $(\Delta C_{\rm T}) = \Delta C_{\rm T, control} - \Delta C_{\rm T, treatment.}$

Transient transfection and dual luciferase assay

D407 cells, a cell-line derived from human RPE cells, were seeded onto 12-well culture plates at a density of 1.0×10^5 cells/well and grown for 24 h. The VEGF promoter-luciferase reporter in the pGal3 vector was transfected into the cells using Fugene 6 transfection reagent (Roche) according to the manufacturer's instructions. In the case of co-transfection experiments, pRL-TK and pGal3-VEGF promoter constructs were cotransfected at a ratio of 3:2. Constructs of pGal3 (+)and pGal3 (-) were used as positive and negative control respectively. Eighteen hours following the transfection, the cells were exposed to 200 µM CoCl₂ in the presence or absence of 160 nM PEDF for 6 h. Then the cells were washed with PBS, and lysed by passive lysis buffer (Promega). The cell lysate was collected and incubated with the luciferase assay reagent (Promega). Luminescence was measured by a microplate luminometer (Berthold Detection Systems, Pforzheim, Germany). The promoter activity was normalized by the Renilla reniformis luciferase from pRL-TK.

Analysis of hypoxia-induced factor- α (HIF-1 α) nuclear translocation

Cultured RCEC were trypsinized, counted and washed three times in PBS. Nuclear and cytoplasmic proteins were extracted using NE-PER Nuclear and Cytoplasmic extraction kit (Pierce, Rockford, IL, USA) according to the manufacturer's protocol. The same amount of nuclear and cytoplasmic proteins was applied to Western blot analysis using an anti-HIF-1 antibody (Santa Cruz Biotechnology, Inc.) at a 1:500 dilution.

Determination of MAP kinase activation

The total proteins from the cultured RCEC or dissected retinas from OIR rats were isolated on 12% SDS-PAGE. The phosphorylation of p42/p44 was measured using a phosphorylation-specific antibody (1:1000, Santa Cruz Biotechnology, Inc.) and normalized by the total p42/p44 levels.

VEGF-binding assay in retinal endothelial cells

VEGF (PeproTech) was labeled with ¹²⁵I using the Chloromine T ¹²⁵I labeling kit (ICN Pharmaceuticals, Inc., Costa Mesa, CA, USA) following the manufacturer's protocol as described previously (Gotte *et al.* 2002, Gao *et al.* 2003). The same amount of ¹²⁵I-VEGF (4 nM) was incubated with primary RCEC for 1 h. Unbound ¹²⁵I-VEGF was removed by four washes. The cells were then lysed, and the ¹²⁵I-VEGF bound to the cells was quantified by a γ -counter, calculated to molar concentration based on VEGF standards.

In vitro VEGF receptor-binding assay

Microtiter plates (Corning Inc., Corning, NY, USA) with 96 wells were coated with 5 nM carrier-free human recombinant VEGF receptor (KDR)/IgG Fc (R&D Systems) in PBS overnight at 4 °C. The wells were then washed three times with PBS containing 0.1% (v/v) Tween 20 (PBS-T) and blocked with 1% BSA in PBS for 2 h. The plates were then incubated for 2 h with 1 nM recombinant human VEGF₁₆₅ in the presence of various concentrations of PEDF or BSA, ranging from 0.1 to 1 000 nM as control. After three washes, a biotinylated anti-VEGF antibody (R&D Systems) was added to the wells and incubated for another 2 h, followed by incubation with horseradish peroxidase-conjugated streptavidin for 20 min. Then the color reagents were added, and the results were measured by a microplate reader (Perkin-Elmer Wallac, Inc., Boston, MA, USA). All the experiments were carried out in triplicate and repeated three times.

Statistical analysis

Statistical analysis employed the Student's *t*-test. Statistical difference was considered significant at P < 0.05.

Results

PEDF decreased VEGF levels in cultured RCEC and Müller cells under normal and hypoxic conditions. RCEC is the major component of the BRB and serves as the primary target of VEGF (Hatcher *et al.* 1998). Cultured primary RCEC have been shown to produce high levels of VEGF (Simorre-Pinatel *et al.* 1994). The retinal Müller cell has been recognized as a major source of VEGF in the retina (Hata *et al.* 1995). We first determined the effect of PEDF on the expression of VEGF in cultured RCEC and Müller cells. Under normal culture conditions, VEGF is

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detected in both RCEC and Müller cells. After treatment of the cells with 160 nM PEDF for 24 h, VEGF levels were dramatically decreased in both the RCEC and Müller cells (Fig. 1A). In RCEC exposed to hypoxia, PEDF has also shown a potent inhibitory effect on VEGF expression in a PEDF concentrationdependent manner (Fig. 1B).

Positive feedback regulation of PEDF in retinal Müller cells

Recombinant PEDF was added to and incubated with retinal Müller cells for 24 h in the presence of 200 μ M cobalt chloride (CoCl₂) or hypoxia (<5% O₂ concentration). The cells were harvested by trypsinization and thoroughly washed by PBS. The intracellular PEDF levels were determined by western blot analysis. The results showed that, after the CoCl₂ (Fig. 2A) or hypoxia (Fig. 2B) treatment, the PEDF expression was drastically decreased. Treatment with exogenous PEDF significantly upregulated the expression of endogenous PEDF in a concentration-dependent manner (Fig. 2A, B). Western blot analysis showed that the PEDF treatment downregulated VEGF expression in the same samples (Fig. 2B).

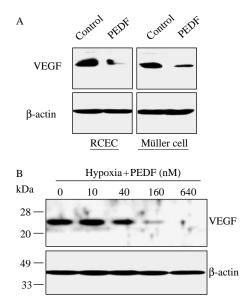


Figure 1 Inhibition of VEGF expression by PEDF in RCEC and Müller cells under normal and hypoxic condition. (A) RCEC and Müller cells were exposed to 160 nM PEDF for 24 h. The intracellular VEGF levels were determined by Western blot analysis. The same membrane was stripped and reblotted with an anti- β -actin antibody. (B) RCEC were exposed to 1% oxygen with different concentrations of PEDF for 24 h. The VEGF levels in the cell lysate were determined by Western blot analysis. The same membrane was stripped and reblotted with the anti- β actin antibody.

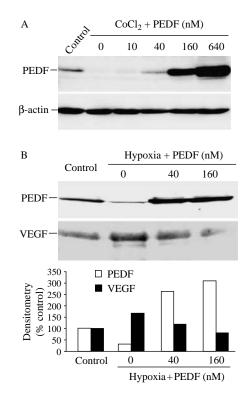


Figure 2 Autoregulation of PEDF and downregulation of VEGF by PEDF in retinal Müller cells. (A) Müller cells were exposed to $200 \ \mu M \ CoCl_2$ with different concentrations of PEDF for 24 h. The intracellular PEDF levels were determined by Western blot analysis. The same membrane was stripped and reblotted with the anti- β -actin antibody. (B) Cultured Müller cells were exposed to 1% oxygen with different concentrations of PEDF for 24 h. The intracellular PEDF level was determined by Western blot analysis. The same membrane was stripped and reblotted with the anti- β -actin antibody. (B) Cultured Müller cells were exposed to 1% oxygen with different concentrations of PEDF for 24 h. The intracellular PEDF level was determined by Western blot analysis. The same membrane was stripped and reblotted with the anti-VEGF antibody. The relative changes of PEDF and VEGF levels were semi-quantified by densitometry as shown in lower panel.

PEDF upregulated endogenous PEDF and downregulated VEGF expression in the retina of rats with OIR

The effect of PEDF on the expression of endogenous PEDF and VEGF was determined in the retina of rats with OIR. The PEDF used for intravitreal injection was expressed and purified from Escherichia coli, which has a molecular weight of 45 kDa, lower than native PEDF (50 kDa) due to the lack of glycosylation. At P14, PEDF was injected into the vitreous of one eye at $1 \,\mu g/eye$ and PBS was injected into the contralateral eye as control. At 48 h after the injection, PEDF and VEGF levels in the retina were determined by western blot analysis. The results showed that endogenous PEDF levels in the retina were significantly higher in the PEDF-injected eyes than in the control eyes (Fig. 3A). Endogenous PEDF had a different molecular weight compared with the recombinant PEDF for the injection, indicating that the elevated retinal PEDF levels are indeed from endogenous expression. The VEGF levels, in contrast,

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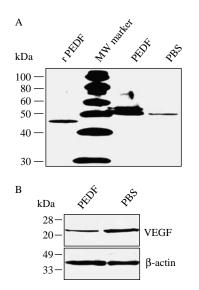


Figure 3 Autoregulation of PEDF and downregulation of VEGF by PEDF in the retina of rats with OIR. Newborn BN rats were exposed to 75% oxygen from age P7 to P12 and returned to room air to induce retinopathy. At P14, PEDF (1 μ g/eye) was injected intravitreally into one eye and PBS into the contralateral eye as control. At 48 h after the injection, the eyeballs were enucleated and the retinas were dissected. The PEDF levels (A) and VEGF levels (B) in the retina were determined by Western blot analysis. rPEDF, recombinant PEDF expressed and purified from *E. coli*.

were drastically decreased in the PEDF-injected eyes compared with the contralateral control eyes (Fig. 3B).

Silencing of PEDF increased VEGF secretion in cultured retinal Müller cells. At 24 h after PEDF siRNA transfection, RT-PCR showed that the PEDF mRNA was largely eliminated (Fig. 4A). The PEDF protein levels in the culture medium were also decreased as shown by ELISA (Fig. 4B). In the same cells, the mRNA and protein levels of VEGF were determined by quantitative real-time RT-PCR and ELISA respectively. The results showed that, after silencing of PEDF, the expression of VEGF was significantly increased at both the mRNA (Fig. 4C) and protein levels (Fig. 4D), suggesting that PEDF is an endogenous, negative regulator of VEGF expression in retinal Müller cells.

PEDF decreased VEGF promoter activity under hypoxic conditions but not under normoxic conditions. To further explore the mechanism responsible for the downregulation of VEGF expression by PEDF in the retina, we determined the effect of PEDF on the VEGF promoter activity in D407 cells. After exposing the transfected cells to CoCl₂ for 6 h, the VEGF promoter activity was significantly increased, indicating a hypoxiainduced activation in VEGF promoter (Fig. 5A, P < 0.0001). PEDF at 160 nM completely blocked the increase in VEGF promoter activity induced by CoCl₂, suggesting that the downregulation of VEGF by PEDF in hypoxia is at least partially through the inhibition of VEGF gene transcription (Fig. 5A, P < 0.0001). Moreover, under normoxic conditions, PEDF had no effect on the VEGF promoter activity (Fig. 5A, P > 0.15), suggesting that PEDF plays an important role in maintaining the normal level of VEGF in the retina.

PEDF inhibits HIF-1 α activation and protein kinase phosphorylation. HIF-1 α is the major transcriptional factor mediating the regulation of VEGF expression (Hata *et al.* 1995). In cultured RCEC exposed to hypoxia with or without PEDF treatment for 24 h, the nuclear and cytoplasmic proteins were isolated from the treated cells. Western blot analysis showed that, under normal culture conditions, HIF-1 α is almost undetectable in the nucleus (Fig. 5B). Exposure to hypoxia for 24 h resulted in a significant increase in the nuclear HIF-1 α level, indicating activation of HIF-1 induced by hypoxia (Fig. 5B). PEDF at 40 and 160 nM dramatically inhibited the hypoxiainduced HIF-1 α nuclear translocation (Fig. 5B).

As the MAP kinase pathway is known to play a role in the VEGF regulation, we have determined the effect of PEDF on the activation of MAP kinase. Serum-deprived RCEC were stimulated with bFGF, and phosphorylation of p42/p44 was determined by western blot analysis. A 5-min challenge with basic fibroblast growth factor (bFGF) resulted in a two-fold increase in p42/p44 phosphorylation when compared with those in the unchallenged cells (data not shown). PEDF decreased the bFGF-induced p42/p44 phosphorylation by fourfold in RCEC (Fig. 5C). Similarly, we have also determined the effect of PEDF on MAP kinase activation in the retina. PEDF was injected into the vitreous of OIR rats at P14, and the retinas were dissected and pooled for MAP kinase phosphorylation assay at 48 h after the injection of PEDF. Consistent with the results in cultured RCEC, PEDF decreased p42/p44 phosphorylation two-fold in the retina of the rats with OIR (Fig. 5D).

PEDF competes with VEGF for binding to RCEC. Incubation of ¹²⁵I-VEGF (4 nM) with RCEC for 1 h resulted in specific binding of VEGF to VEGF receptor (VEGFR) on RCEC. To determine the competition between PEDF and VEGF for VEGFR binding, the same amount of ¹²⁵I-VEGF was added to RCEC together with increasing concentrations (40, 400, and 4000 nM) of unlabeled PEDF or plasminogen kringle 5 (K5), another potent angiogenic inhibitor. Unlabeled VEGF (400 nM) was used as a positive control. In the presence of excess amounts of unlabeled PEDF, VEGF bound to RCEC was decreased in a PEDF concentration-dependent manner (Fig. 6A). In contrast, K5 in the same concentration range did not inhibit VEGF binding with VEGFR, suggesting different mechanisms of action between PEDF and K5 (Fig. 6A), although they both reduce vascular permeability and inhibit retinal NV.

PEDF competed with VEGF for binding to VEGFR2 *in vitro*. To further explore whether the effect of PEDF on the blockade of VEGF binding to endothelial cells

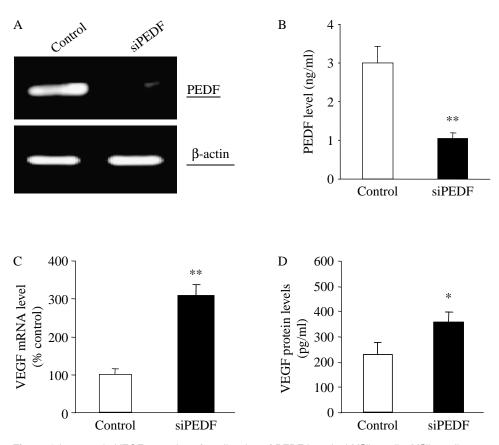


Figure 4 Increase in VEGF secretion after silencing of PEDF in retinal Müller cells. Müller cells were transfected with a PEDF siRNA or control, and cultured for 24 h. (A). The mRNA level of PEDF in the cells was analyzed by RT-PCR and normalized by β -actin levels. (B). The PEDF secreted into the culture medium was measured by ELISA (mean ±s.D., n=3). (C). The mRNA level of VEGF in the same cells was determined by real time RT-PCR and normalized by 18S RNA. The results were expressed as percentage of control (mean ±s.D., n=3). (D). The protein levels of VEGF in the culture medium were measured by ELISA (mean ±s.D., n=3). (D). The protein levels of VEGF in the culture medium were measured by ELISA (mean ±s.D., n=3). (D). The values statistically different from the control are indicated by *P < 0.05, **P < 0.01.

is through the competition with VEGF for binding to the VEGF receptor, we performed an *in vitro* receptorbinding assay. VEGFR2, which is recognized as the major form of VEGF receptor responsible for the activity of VEGF in inducing vascular permeability and angiogenesis, was immobilized on a 96-well plate. The binding of VEGF to VEGFR2 was measured by an ELISA-based method (R&D Systems). The results showed that the binding of VEGF to VEGFR2 was significantly inhibited by excess PEDF in a PEDF concentration-dependent manner (Fig. 6B). Moreover, the binding was not affected by the same concentrations of albumin (negative control) and heat-inactivated PEDF protein, suggesting that the competition of PEDF with VEGF on VEGFR2 binding is a specific effect (Fig. 6B).

VEGF decreased PEDF expression and secretion in RCEC but not in Müller cells. The effect of VEGF on the expression of PEDF was investigated in cultured RCEC 24-h incubation of RCEC with 40 ng/ml VEGF, the PEDF expression was drastically decreased (Fig. 7A). In retinal Müller cells, however, the same concentration of VEGF had no effect on the expression of PEDF (Fig. 7A). The same membrane was stripped and reblotted with an anti-VEGF antibody, which showed that VEGF induced a significant increase of endogenous VEGF expression in RCEC, but not in Müller cells (Fig. 7A and B), consistent with the cell type-specificity in its effect on PEDF expression. The effect of VEGF on the regulation of PEDF was further confirmed by measuring the PEDF secretion into the culture medium of RCEC. The ELISA result showed that VEGF decreased PEDF secreted from RCEC in a VEGF concentration-dependent manner (Fig. 7B). These results suggest that the effect of VEGF on the regulation of PEDF production and its autoregulation is endothelial cell-specific.

and Müller cells. Western blot analysis showed that after

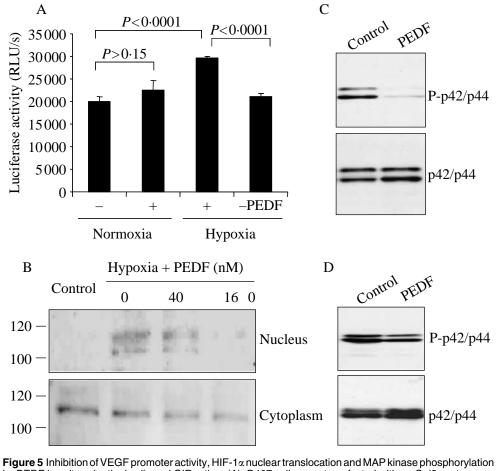


Figure 5 Inhibition of VEGF promoter activity, HIF-1α nuclear translocation and MAP kinase phosphorylation by PEDF in cultured retinal cells and OIR retina. (A). D407 cells were transfected with a pGal3 vector containing the VEGF promoter and a luciferase reporter gene, in triplicate. Eighteen hours following transfection, the cells were exposed to 200 μM CoCl₂ in the presence or absence of 160 nM PEDF for 6 h. The luciferase activity was measured and normalized as described in Materials and methods. (B). Primary RCECs were exposed to 1% oxygen in the presence or absence of various concentrations of PEDF for 24 h. The nuclear protein and cytoplasmic proteins were isolated from the cells. The HIF-1α levels in the nucleus and in the cytoplasm were determined by Western blot analysis using the same amounts of proteins. (C). Primary RCECs were stimulated with bFGF in the presence or absence of 160 nM PEDF for 5 min. The phosphorylated p42/p44 was semi-quantified by Western blot analysis using a phosphorylation-specific antibody and normalized by total p42/p44 levels. (D). At 48 h after the PEDF intravitreal injection in OIR rats, the phosphorylated and total p42/p44 levels in the retina were determined by Western blot analysis.

Discussion

In recent years, accumulating evidence suggests that the decrease of endogenous angiogenic inhibitors plays an important role in pathological vascular leakage and angiogenesis (Cao 2001, Gao *et al.* 2001, Boehm *et al.* 2003). PEDF, a serine proteinase inhibitor, has been found as a major anti-angiogenic factor in the eye (Dawson *et al.* 1999). High levels of PEDF in the transparent avascular ocular compartments, including the cornea and the vitreous, suggest its critical role in maintaining the local avascularity via its anti-angiogenic activity (Karakousis *et al.* 2001). In diabetic patients with aqueous humor are significantly decreased, which predicts the progression of retinopathy (Ogata *et al.* 2001*b*, 2002, Boehm *et al.* 2003). Our previous study demonstrated that in OIR rats, the decrease in PEDF and the increase in VEGF in the retina correlate with the elevation of vascular permeability and retinal NV, suggesting that a local balance between PEDF and VEGF is crucial for the regulation of retinal vascular function (Gao *et al.* 2001, Zhang *et al.* 2005*b*). The present study confirmed this finding by showing decreased PEDF expression and increased VEGF expression in hypoxia-exposed retinal Müller cells

active retinopathy, PEDF levels in the vitreous and

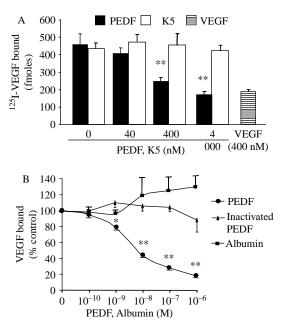


Figure 6 Blockade of the binding of VEGF to VEGFR2 by PEDF. (A) ¹²⁵I-VEGF was incubated with primary RCEC in the absence and presence of excess amounts of PEDF or K5 as indicated. Unlabeled VEGF was used as a positive control. The binding of VEGF on RCEC was measured. Bars represent the bound VEGF (fmoles/well, means \pm s.b., n=3), and ** indicates the values statistically different from the control (VEGF alone) (P<0·01). (B) Immobilized VEGFR2 was incubated with VEGF in the absence and presence of various concentrations of PEDF, heat-inactivated PEDF (100 °C for 5 min) or albumin. The binding of VEGF to VEGFR2 was detected as described in Materials and methods section. Values statistically different from the albumin control are indicated as *P<0·05, **P<0·01.

and RCEC. Although a number of studies have shown that oxygen concentration regulates the retinal levels of VEGF and PEDF, interactions between these two counterbalancing angiogenic factors are largely unclear (Gao et al. 2002a, Eichler et al. 2004, Werdich et al. 2004). The present study demonstrated for the first time that there is a reciprocal regulation between VEGF and PEDF in the retina. In cultured RCEC, PEDF downregulated hypoxia-induced VEGF expression at least partially via the inhibition of MAP kinase phosphorylation and HIF-1 nuclear translocation. In retinal Müller cells and in the retina of the OIR rats, PEDF upregulated the expression of endogenous PEDF. Moreover, silencing of PEDF alone significantly upregulated VEGF expression and secretion, suggesting that PEDF is an endogenous inhibitor of VEGF expression. In addition to the regulation of VEGF expression, PEDF effectively inhibited VEGF binding to retinal endothelial cells. In vitro studies demonstrated that PEDF competed with VEGF binding to VEGFR2. This competitive blockade of the VEGF-VEGFR2 binding may be responsible for the PEDF effects on VEGF-induced permeability and angiogenesis. On the other hand, VEGF also significantly reduced PEDF expression and upregulated the expression of endogenous VEGF in endothelial cells, but not in Müller cells, suggesting a VEGF receptor-mediated regulation. These findings revealed a complex reciprocal regulation between PEDF and VEGF, which may be an important mechanism responsible for maintaining the balance between the angiogenic stimulator system and angiogenic inhibitor system in the retina.

VEGF is a potent VPF and a major angiogenic stimulator with endothelial cell-specific mitogenic activity and plays a crucial role in both normal and pathological angiogenesis (Dvorak et al. 1995, Pierce et al. 1995, Aiello 1996, Aiello et al. 1997). In diabetic patients with proliferative retinopathy, VEGF levels are significantly increased in the vitreous and the retina, and successful laser treatment decreased vitreous VEGF levels by 75% (Adamis et al. 1994, Aiello et al. 1994, Pe'er et al. 1995). In patients with DME, significantly elevated VEGF levels in the aqueous humor are correlated with the severity of macular edema (Funatsu et al. 2003). These findings suggest that over-production of VEGF in the retinal diabetics is a major cause of DME as well as retinal NV. In the retina, VEGF is produced by multiple cell types, including the RPE, pericytes, endothelial, glial, Müller, and ganglion cells (Pe'er et al. 1995, Dorey et al. 1996, Lu & Adamis 2002). Among them, Müller cells and RPE are believed to be the major source of VEGF in the retina, and endothelial cells to be the primary target of VEGF (Pierce et al. 1995, Dorey et al. 1996). Cultured vascular endothelial cells also express high levels of VEGF in an autocrine manner (Simorre-Pinatel et al. 1994). In the present study, we have investigated the effect of PEDF on the regulation of VEGF expression in cultured retinal Müller cells and RCEC. Our results demonstrated that, under normoxic and hypoxic conditions, PEDF significantly downregulated VEGF expression in both RCEC and Müller cells as well as in the retina of the OIR rat model. Moreover, in cultured RPE cells, PEDF inhibited the hypoxiainduced increase in VEGF promoter activity, but had no effect on VEGF promoter activity under normoxic conditions. These results suggested that PEDF is a negative regulator of VEGF expression in the retina. In the normal eye, the high levels of PEDF in the vitreous and the retina inhibits the over-expression of VEGF, which, in turn, maintains the quiescence of retinal vascular endothelial cells and the integrity of BRB. Under hypoxia and other pathological conditions, in addition to the upregulation of VEGF induced directly by the primary insults, the decreased PEDF levels also contribute to the over-production of VEGF. The fact that silencing of PEDF gene expression alone is sufficient to increase VEGF expression further supports the role of PEDF as an endogenous inhibitor of VEGF

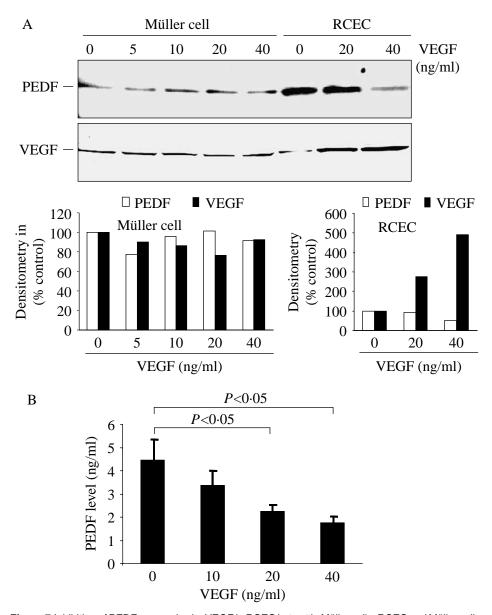


Figure 7 Inhibition of PEDF expression by VEGF in RCEC but not in Müller cells. RCEC and Müller cells were exposed to different concentrations of VEGF for 24 h. (A). The intracellular PEDF levels were determined by Western blot analysis. The same membrane was stripped and reblotted with an anti-VEGF antibody. The relative levels of PEDF and VEGF were semi-quantified by densitometry analysis. (B). The PEDF levels in the medium of cultured RCEC were measured by ELISA (mean \pm s.D., n=3).

expression and suggests that the reduced PEDF levels contribute to over-production and subsequent activation of vascular endothelial cells.

HIF-1 is a major transcriptional factor regulating VEGF expression under hypoxia (Forsythe *et al.* 1996, Ozaki *et al.* 1999). The present study demonstrated that the nuclear HIF-1 α level was elevated significantly in the retinal endothelial cells insulted by hypoxia, correlating with increased VEGF expression. Moreover, the total

amount of HIF-1 α was drastically increased under hypoxic conditions (data not shown). Incubation with PEDF significantly reduced the nuclear HIF-1 α levels, indicating an inhibited HIF-1 nuclear translocation. These results suggest that inhibiting HIF-1 activation is responsible, at least partially, for the downregulated VEGF expression by PEDF. Furthermore, we determined if p42/p44 MAP kinase activation, an important signaling pathway in VEGF expression, is involved in the regulation of VEGF by PEDF (Milanini *et al.* 1998). Our results demonstrated that PEDF significantly inhibited the activation of p42/p44 in the retinal endothelial cells as well as in OIR retina, suggesting that the downregulation of VEGF by PEDF may be, at least in part, via inhibition of the MAP kinase activation. It is noteworthy that we have used a 24-h time scale in the experiments, as we found that the nuclear HIF-1 level was significantly higher at this time point. Although our results showed that PEDF inhibited MAPK activation and, thus, may directly block the downstream activation of HIF-1, we cannot exclude the possibility that this inhibitory effect of PEDF may be secondary, i.e. via blocking the positive feedback regulation of VEGF.

In addition to the regulation of VEGF expression, the present study, for the first time, demonstrated that PEDF effectively inhibited VEGF binding to VEGFR2 on RCEC and in isolated VEGFR2 in vitro. VEGFR2 is the major type of receptor responsible for the activity of VEGF on vascular permeability and angiogenesis (Aiello et al. 1995, Ozaki et al. 2000, Witmer et al. 2002). Several studies have shown that blockade of the activation of VEGFR2 induced by VEGF binding successfully abrogated retinal NV (Aiello et al. 1995, Ozaki et al. 2000). In the past decade, although PEDF has been well studied as a potent angiogenic inhibitor, the molecular mechanisms for its anti-angiogenic and anti-permeability activities are still uncertain. Our results demonstrated that PEDF significantly competed with the VEGF binding to VEGFR2, while the competitive capacity was lost in heat-inactivated PEDF. In endothelial cells, we have used an excess amount of PEDF to ensure the competitive effect of PEDF (Gotte et al. 2002). The structural basis responsible for the competition between PEDF and VEGF for binding to VEGFR2 remains to be elucidated.

PEDF is a multifaceted factor with neurotropic and anti-angiogenic functions (Tombran-Tink & Barnstable 2003b, Tombran-Tink et al. 1991). Although PEDF as a major endogenous angiogenic inhibitor has been extensively studied, the regulation of PEDF is not yet well understood (Tombran-Tink & Barnstable 2003b). Our study demonstrated that PEDF is secreted from RCEC and Müller cells. Under hypoxic conditions, PEDF secretion was significantly decreased along with the increase in VEGF expression. This result is consistent with the recent study reported by Eichler et al. (2004), which shows that hypoxia downregulates PEDF expression at the RNA level by 1.7-fold and at the protein level by two-fold in retinal Müller cells. Although hypoxia did induce a significant increase in VEGF expression and secretion, the addition of exogenous VEGF did not cause any change in the expression of PEDF in Müller cells, suggesting that hypoxia-induced downregulation of PEDF expression

in Müller cells cannot be ascribed to the elevated VEGF levels. On the other hand, VEGF significantly downregulates PEDF expression in retinal endothelial cells, but not in Müller cells, suggesting that the regulation of PEDF by VEGF is dependent on VEGF receptor, which is expressed in endothelial cells but not in Müller cells.

Another important finding of the current study is that PEDF positively autoregulated endogenous PEDF expression in cultured Müller cells and in the retina of diabetic rats. PEDF is a major endogenous angiogenic inhibitor in the retina. Moreover, PEDF has recently been identified as a potent anti-inflammatory factor and anti-permeability factor (Liu *et al.* 2004, Zhang *et al.* 2005*a*). Furthermore, a decrease in ocular PEDF levels is believed to contribute to the pathogenesis of DR. Therefore, upregulation of endogenous PEDF levels in the retina will be beneficial in diabetic patients. The mechanisms by which PEDF is regulated by VEGF and by PEDF itself via positive feedback regulation remain to be determined in the future.

In summary, our study demonstrated that there exists a reciprocal regulation between PEDF and VEGF in the retina, which may represent an important mechanism for maintaining the balance between the two angiogenic regulatory systems. PEDF downregulates VEGF expression via inhibition of MAPK-mediated HIF-1 activation and competes with VEGF binding to VEGFR2. The decrease in PEDF levels is, at least partially, responsible for the increase in VEGF expression and the subsequent BRB breakdown and retinal NV in diabetic retinopathy. This study suggests that the angiogenic stimulator system and angiogenic inhibitor system have complex interactions, which may represent an important mechanism for maintaining the homeostasis of angiogenesis. The detailed mechanism responsible for these interactions remains to be elucidated.

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