Synergism between vascular endothelial growth factor and placental growth factor contributes to angiogenesis and plasma extravasation in pathological conditions

PETER CARMELIET¹, LIEVE MOONS¹, AERNOUT LUTTUN¹, VALERIA VINCENTI², VEERLE COMPERNOLLE¹, MARIA DE MOL¹, YAN WU³, FRANÇOISE BONO⁴, LAETITIA DEW⁵, HEIKE BECK⁶, DIMITRI SCHOLZ⁷, TILL ACKER⁶, TINA DIPALMA², MIEKE DEWERCHIN¹, AGNES NOEL⁵, INGEBORG STALMANS¹, ADRIANO BARRA², SYLVIA BLACHER⁵, THIERRY VANDENDRIESSCHE¹, ANNICA PONTEN⁹, ULF ERIKSSON⁹, KARL H. PLATE⁶, JEAN-MICHEL FOIDART⁵, WOLFGANG SCHAPER⁷, D. STEPHEN CHARNOCK-JONES⁸, DANIEL J. HICKLIN³, JEAN-MARC HERBERT⁴, DÉSIRÉ COLLEN¹ & M. GRAZIELLA PERSICO²

¹ The Center for Transgene Technology and Gene Therapy, Flanders Interuniversity Institute for Biotechnology, KU Leuven, Leuven, Belgium

²lstituto Internazionale di Genetica e Biofisica, CNR, Naples, Italy

³ImClone Systems, New York, New York, USA

⁴Cardiovascular/Thrombosis Research Department, Sanofi-Synthélabo, Toulouse, France,

⁵Laboratory of Tumor and Developmental Biology, University Liège, Sart-Tilman, Belgium

⁶Department of Neuropathology, FAU Erlangen-Nürnberg, Erlangen, Germany

⁷Department of Experimental Cardiology, Max-Planck-Institute, Nauheim, Germany

⁸Reproductive Molecular Research Group, Department Obstetrics & Gynaecology, University of Cambridge, Cambridge, UK

⁹Ludwig Institute for Cancer Research, Stockholm, Sweden

Abstract: Vascular endothelial growth factor (VEGF) stimulates angiogenesis by activating VEGF receptor-2 (VEGFR-2). The role of its homolog, placental growth factor (PIGF), remains unknown. Both VEGF and PIGF bind to VEGF receptor-1 (VEGFR-1), but it is unknown whether VEGFR-1, which exists as a soluble or a membrane-bound type, is an inert decoy or a signaling receptor for PIGF during angiogenesis. Here, we report that embryonic angiogenesis in mice was not affected by deficiency of PIGF (*Pgf*^{-/-}). VEGF-B, another ligand of VEGFR-1, did not rescue development in *Pgf*^{-/-} mice. However, loss of PIGF impaired angiogenesis, plasma extravasation and collateral growth during ischemia, inflammation, wound healing and cancer. Transplantation of wild-type bone marrow rescued the impaired angiogenesis and collateral growth in *Pgt*^{-/-} mice, indicating that PIGF might have contributed to vessel growth in the adult by mobilizing bone-marrow-derived cells. The synergism between PIGF and VEGF was specific, as PIGF deficiency impaired the response to VEGF, but not to bFGF or histamine. VEGFR-1 was activated by PIGF, given that anti-VEGFR-1 antibodies and a Src-kinase inhibitor blocked the endothelial response to PIGF or VEGF/PIGF. By upregulating PIGF and the signaling subtype of VEGFR-1, endothelial cells amplify their responsiveness to VEGF during the 'angiogenic switch' in many pathological disorders.

Vascular endothelial growth factor (VEGF) stimulates angiogenesis by activating the VEGF tyrosine kinase receptor-2 (VEGFR-2/KDR)^{1,2}. Little is known, however, about other VEGF homologs such as placental growth factor (PIGF)³. Indirect evidence points to a role for PIGF in placental development⁴, whereas recombinant PIGF stimulates angiogenesis in particular conditions and induces vascular permeability when co-injected with VEGF (refs. 5,6), but the role of endogenous PIGF remains unknown^{7,8}. PIGF, VEGF and another homolog, VEGF-B, all bind to VEGF receptor-1 (VEGFR-1/Flt-1)^{9,10}. Loss of VEGFR-1 disrupted normal vascular development¹¹, but deletion of its tyrosine kinase domains allowed normal embryonic angiogenesis¹², indicating that VEGFR-1 might function as an inert 'decoy' by binding VEGF and thereby regulating the availability of VEGFR-1, an alternatively processed type of VEGFR-1 which contains the extracellular ligand-binding domains, but lacks the signaling tyrosine kinase domains^{13,14}.

PIGF has been proposed to stimulate angiogenesis by displacing VEGF from the 'VEGFR-1 sink', thereby increasing the fraction of VEGF available to activate VEGFR-2 (ref. 6). Alternatively, PIGF might stimulate angiogenesis by transmitting intracellular signals through VEGFR-1 (refs. 9,10,16). However, VEGFR-1 has weak tyrosine kinase activity and its intracellular signals are not well understood. PIGF might also affect angiogenesis by forming heterodimers with VEGF, but their role is controversial^{16,17}. Due to lack of available inhibitors of PIGF or VEGFR-1, their role in pathological angiogenesis could not be demonstrated thus far.

To examine the role of PIGF in angiogenesis, we inactivated the gene expressing PIGF (*Pgf*) in mice. Unexpectedly, the absence of PIGF—even in combination with a loss of VEGF-B—had a negligible effect on vascular development. However, PIGF deficiency reduced pathological angiogenesis, permeability and collateral growth in ischemia, inflammation and cancer. These genetic studies unveil a synergism between PIGF and VEGF in pathological angiogenesis, which might have therapeutic implications for either stimulating or inhibiting angiogenesis.

Fig. 1 Normal vascular development but impaired tumor angiogenesis in $Pgf^{\ell_{\infty}}$ mice, **a**, Western blotting for VEGFR-1. Membrane-associated VEGFR-1 (Mr ~190-200 kD) represents only 2% and 30% of total VEGFR-1 in placenta and ischemic myocardium, respectively. Soluble VEGFR-1 (Mr ~100-110kD). Lanes 1-2: placental extract at E9.5 and E.15.5. Lane 3: E15.5 placental extract, mixed with VEGF and PIGF and immuno-precipitated using anti-VEGF and anti PIGF anti-bodies. Lane 4, Myocardial infarct (MI) extract, *b-d*, Labeling of vitelline blood vessels upon binding of [¹²⁵]VEGF₁₆₅ to E9.5 yolk-sac sections (*b*). Excess cold VEGF (*d*) but not PIGF (c) almost completely eliminated labeling, *e* and *f*, Toluidine blue-stained section of a corpus luteum, revealing more numerous and larger vessels in wild-type (e) than in $Pgf^{\ell_{\infty}}$ (f) mice. For clarity, vessels were traced by white lines, *g-i*, Macroscopic pictures of wild-type (*g*), $Pgf^{\ell_{\infty}}$ (*h*) and $Vegf^{\ell_{\infty}}$ (i) ES-cell-derived tumors, *j*, Densities (vessels/optical field) of vessels of different size in wild-type tumors in wild-type mice, and $Pgf^{\ell_{\infty}}$ or $Vegf^{\ell_{\infty}}$ tumors in $Pgf^{\ell_{\infty}}$ mice. \blacksquare , wild-type; \blacksquare , $Pgf^{\ell_{\infty}}$; \blacksquare , $Vegf^{\ell_{\infty}}$, *, P < 0.05 versus wild-type. *k* and *i* Endoglin immunostaining, revealing large vessels in wild-type fibrosarcoma (*k*) but only small capillaries in $Pgf^{\ell_{\infty}}$ fibrosarcoma (*l*). Scale bars: 25 µm (*b-f*), 4 mm (*g-i*) and 50 µm (*k* and *l*).



Normal embryogenesis and subtle vascular defects in $Pgf^{/-}$ mice

We inactivated Pgf by deleting exons 3-6 and confirmed the deletion by Southern- and northern-blot analyses and ELISA (< 0.5 pg PIGF/mg in Pgf' mice; data not shown). Pgf' mice from Pgf' or Pgf' breeding pairs were born at a mendelian frequency and were healthy and fertile. Although PIGF was expressed in wild-type embryos at embryonic day (E)10.5 (33 ± 7 pg/mg protein; n = 6; mean ± s.e.m.), vascular development in Pgf' embryos was normal (data not shown). PIGF levels were 60% lower, however, than VEGF levels (81 ± 4 pg/mg; n = 8) and PIGF was only 10% as potent as VEGF in enlarging embryonic vessels after intracardial injection in cultured E8.5 wild-type embryos (data not shown).

VEGFR-1 is the only known signaling receptor for PIGF. However, western-blot (Fig. 1a), radioligand binding (Fig. 1b-d) and quantitative reverse transcriptase (RT)-PCR (data not shown) analyses revealed that most VEGFR-1 in the embryo from E9.5 onwards was present as a soluble form. Consistent with an inhibitory decoy role of VEGFR-1, intracardial injection of VEGFR-1 antibodies in early-stage embryos stimulated vascular development, whereas VEGFR-2 antibodies blocked it (data not shown). Although more VEGF would be bound by VEGFR-1 in the absence of PIGF, plasma VEGF levels were not reduced in $Pgf^{/-}$ embryos (-65 pg/ml in wild-type embryos versus -84 pg/mg in $Pgf^{/-}$ embryos at E15.5). This might be due to a compensatory increase in production of VEGF in $Pgf^{/-}$ embryos (81 ± 4 pg/mg in wild-type embryos versus 118 ± 11 pg/mg in $Pgf^{/-}$ embryos at E10.5; n = 8; P < 0.01). Other angiogenic molecules (VEGFR-1, VEGF-B, VEGF-C and angiopoietin-1) were comparably expressed in $Pgf^{/-}$ embryos (data not shown).

VEGF-B is another ligand of VEGFR-1 (ref. 18), but loss of VEGF-B (*Vegfb^{-/-}*) does not affect vascular development¹⁹. Although VEGF-B could have rescued vascular development in *Pgf^{-/-}* mice, *Vegfb^{-/-} Pgf^{/-}* mice were born at a normal mendelian inheritance, were apparently healthy and fertile, had a normal life span and were without obvious vascular defects (data not shown). This indicates that the individual VEGFR-1 ligands may have distinct functions in angiogenesis.

We observed no structural vascular defects in $Pgf^{/-}$ mice after birth, except for a subtle remodeling defect of retinal vessels (data not shown). Fewer vessels developed in corpora lutea at 4.5 days of pregnancy in $Pgf^{/-}$ mice (vessels/mm²: 910 ± 62 in 6 wild-type mice versus 430 ± 80 in 6 $Pgf^{/-}$ mice; P< 0.005; Fig. 1e and *f*), but did not impair reproduction. Thus, loss of PIGF impaired VEGF-dependent retinal and luteal angiogenesis.

Fig. 2 Impaired retinal and myocardial angiogenesis in *Pgf* mice, **a** and **b**, Impaired ischemic retinal angiogenesis in *Pgf* mice (fluoro-angiogra-phy). Numerous foci of neovascularization (arrowheads) and large dialated and tortuous retinal vessels in wild-type (a) but not in *Pgf* mice (b). **c** and **d**, Immunostaining of VEGFR-1 in a wild-type mouse, revealing minimal expression during hyperoxia (P12; c) and significant upregulation in retinal vessels (arrowheads) after 1 d of ischemia (P13; d). **e**, Loss of PIGF reduces the number of endothelial cells and neovascular tufts in the vitreous cavity. *, P < 0.05 versus wild-type in the vitreous cavity, **f** and **g**. Thrombomodulin (TM) staining of endothelium, revealing more numerous vessels in wild-type (f) than in *Pgf* (g) myocardial infarcts, **h** and **i**, *In situ* hybridization revealing undetectable PIGF expression in normal myocardium (h) and upregulation of PIGF in ischemic cardiomyocytes and in vessels infiltrating the infarct (h and i; wild-type mice).j and k, Immunostaining revealing that VEGFR-1 expression is undetectable in quiescent myocardial vessels (j), but induced in angiogenic vessels infiltrating the infarct (j and k; wild-type mice). Scale bars: 200 µm (a and b), 25 µm (c and d), 50 µm (f, g and i-k) and 100µm (h).



Impaired tumor angiogenesis in Pgf^{-} mice

PIGF is variably upregulated in tumors^{8.20-22} and its role in tumor angiogenesis remains ill defined. Therefore we grew embryonic stem (ES)-cell-derived tumors of either genotype in athymic *nu/nu* wild-type or Pgf' mice for four weeks. When both tumor and host expressed PIGF, tumors were large and highly vascularized (4 ± 1 g; *n* = 8; Fig. 1g and j). In contrast, when they both lacked PIGF, tumors remained small and poorly vascularized (1.0 ± 0.3 g; *n* = 8; Fig. 1h and j), comparable to the reduced growth and angiogenesis of VEGF-deficient tumors (0.3 ± 0.1 g; *n* = 10; Fig. 1 i and j). When either tumor or host produced PIGF, tumor growth and angiogenesis were intermediate, indicating that production of PIGF by tumor- and host-derived tissue contributed to tumor angiogenesis. Coverage of tumor vessels by smooth muscle cells was not affected by *Pgf* genotype. We confirmed the role of PIGF in tumor angiogenesis using ras-transformed fibroblasts. *Pgf'*⁻ fibrosarcomas were

smaller than wild-type fibrosarcomas when implanted in *nu/nu* wild-type mice $(2.8 \pm 0.2 \text{ g versus } 5.5 \pm 0.6 \text{ g}; \prod = 7; P < 0.05)$, and, although vascular densities were comparable in both tumors (~550 vessels/mm²), *Pgf^{/-}* fibrosarcomas contained vessels with a smaller lumen than wild-type fibrosarcomas $(110 \pm 18 \ \mu\text{m}^2 \text{ versus } 210 \pm 12 \ \mu\text{m}^2; n = 7; P < 0.05; Fig. 1k \text{ and } I)$. As revealed by *in situ* hybridization analysis of wild-type fibrosarcomas, PIGF was expressed in endothelial and tumor cells, whereas VEGFR-1 was highly expressed in tumor vessels (data not shown). Wild-type tumors expressed large amounts of PIGF ($85 \pm 12 \ \text{pg/mg}$ protein; n = 5) and VEGF ($430 \pm 40 \ \text{pg/mg}$ protein; n = 5). However, reduced tumor angiogenesis in *Pgf^{/-}* tumors did not result from reduced expression of VEGF, VEGF-B or VEGF-C (data not shown). Thus, loss of PIGF impaired growth of VEGF-dependent tumors.

Fig. 3 Impaired pathological arteriogenesis in Pgf' mice, a, Time course of skin wound healing. \blacksquare , wild-type; \bigcirc , Pgt'~ mice; n = 15 mice; P < 0.05 versus wild-type, b and c, Macroscopic view of impaired skin wound healing in Pgf' (c) as compared with wild-type (b) mice, d and e, Smooth muscle α -actin (SMA) staining, revealing impaired coverage of newly formed vessels by smooth muscle cells (arrowheads) in Pgf' (e) as compared with wild-type (d) mice, f and g, PIGF immunostaining of human skin, revealing minimal expression in normal skin (f) and upregulation during chronic eczema (g). Scale bars: 4 mm (b and c), 50 µm (d-g).



Reduced angiogenesis in ischemia in Pgf^{-} mice

During ischemic retinopathy, increased VEGF levels stimulate vitreous neovascularization. Previous studies reported minimal changes in PIGF transcripts²³ or increased PIGF protein levels²⁴, but the role of PIGF in retinal neovascularization remains unknown. We induced capillary dropout in the retina by exposing seven-day-old (P7) mice to 80% oxygen for five days. Upon return to normoxia at P12, the hypovascular retina became ischemic, upregulated VEGF and induced venous dilation, arterial tortuosity and capillary growth in the vitreous chamber (Fig. 2a and e). Loss of PIGF significantly protected mice against intravit-reous neovascularization (n = 19; P < 0.005; Fig. 2b and e), venous dilatation and arterial tortuosity (Fig. 2 b). During retinal ischemia, VEGFR-1 was induced in vascular sprouts (Fig. 2 c and *d*), whereas VEGF and PIGF were significantly upregulated (pg VEGF/mg protein: 98 ± 12 at P12 versus 850 ± 18 at P13; n = 7; P < 0.005 and pg PIGF/mg protein: 10 ± 2 at P12 versus 850 ± 18 at P13; n = 7; P < 0.005 and pg PIGF/mg protein: 10 ± 2 at P12 versus 850 ± 18 at P13; n = 7; P < 0.005 and pg PIGF/mg protein: 10 ± 2 at P12 versus 38 ± 3 at P13; n = 3; P < 0.05). VEGF levels were comparable, however, in both genotypes (data not shown).

We analyzed revascularization of myocardial infarcts after ligating the left anterior descending coronary artery²⁵. Compared with wild-type mice, $Pgf^{/-}$ infarcts contained approximately 20% fewer vessels (vessels/mm²: 134 ± 9 in wild-type mice versus 111 ± 5 in $Pgf^{/-}$ mice; n = 12 to 24; P < 0.05; Fig. 2f and g). VEGF and PlGF levels (expressed per mg protein) were increased in infarct borders where angiogenesis occurs: normal myocardium produced 22 ± 2 pg VEGF and 8 ± 1 pg PlGF, whereas infarcts produced 52 ± 6 pg VEGF and 102 ± 8 pg PlGF

(n = 5; P < 0.05). Expression of VEGF, VEGFR-2 and endoglin in infarcts was comparable in both genotypes (data not shown). PIGF and VEGFR-1 were undetectable in quiescent vessels in normal myocardium, but were upregulated in angiogenic vessels and macrophages infiltrating ischemic myocardium (Fig. 2*h*-*k*). Administration of recombinant hPIGF-1 to $Pgf^{/-}$ mice rescued infarct revascularization (data not shown), which was associated with increased accumulation of macrophages (Mac-3-positive area/infarct area: $1.8 \pm 0.4\%$ after vehicle versus $5.0 \pm 0.7\%$ after PIGF; n = 4; P < 0.05). Taken together, loss of PIGF impaired VEGF-dependent retinal and myocardial angiogenesis.

Role of PIGF during skin wound healing

Healing of skin incisions progressed more slowly in Pgf'' mice versus wild-type (Fig. 3a-c). Both genotypes contained comparable densities of vessels in unwounded skin (~220 vessels/mm²) and of new capillaries infiltrating into the wound (vessels/mm²: 240 ± 50 in wild-type mice versus 180 ± 50 in Pgf' mice; n = 5; P = NS). However, wild-type mice contained more smooth muscle-coated vessels than naked vessels (vessels/mm²: 75 ± 18 smooth muscle α -actin-positive vessels versus 40 ± 7 naked; n = 5), whereas the opposite was true for Pgf' mice (vessels/mm²: 84 ± 13 naked versus 30 ± 10 mature vessels; n = 5; P < 0.05 versus wild-type; Fig. 3d and e). VEGFR-1 and VEGFR-2 were induced in wounded skin vessels (data not shown), whereas PIGF expression was upregulated in endothelial cells and in the hyperplastic epidermis at the wound edge, where new vessels formed (Fig. 3f and g). VEGF expression was comparably increased in the hyperplastic epidermis in both genotypes (see below). Coverage of pulmonary vessels by smooth muscle cells was also impaired in Pgf' mice after exposure to hypoxia (data not shown).

| Effect | Treatment | Wild-type | Pgf'- |
|--------------------------------|---|-------------|--------------|
| Apoptosis | Vehicle | 100 | 100 |
| (% of vehicle control) | $\mathbf{DICE}(100 \text{ ng/ml})$ | 07 ± 0 | 06 ± 10 |
| | PIGF (100 ng/ml) | $9/\pm 9$ | 96 ± 10 |
| | VEGF (100 ng/ml) | 22 ±8* | 93 ± 10 |
| | VEGF (100 ng/ml) + PP2 (100 μ M) | ND | 102 ± 5 |
| | VEGF + PIGF (50 ng/ml each) | ND | $50 \pm 13*$ |
| | VEGF + PlGF (100 ng/ml each) | ND | $25 \pm 5*$ |
| | VEGF + PIGF (100 ng/ml each) + anti-VEGFR-1 (50 μ g/ml) | ND | 100 ± 11 |
| | VEGF + PlGF (100 ng/ml each) + anti-mPlGF (50µg/ml) | ND | 75 ± 9 |
| | VEGF + PIGF (100 ng/ml each) +PP2 (100 μ M) | ND | 107 ± 5 |
| | VEGF/PIGF heterodimer (100 ng/ml) | ND | 17±7* |
| Migration (migrating cells) | Vehicle | 5±3 | 6±2 |
| | PlGF (100 ng/ml) | 7 ±2 | 8±2 |
| | VEGF (100 ng/ml) | $88 \pm 9*$ | 24 ±2* |
| | VEGF + PlGF (100 ng/ml each) | 92 ±5* | 97 ±3* |
| | bFGF (50 ng/ml) | 96 ±6* | 102 ±13* |
| | bFGF (50 ng/ml) + PlGF (100 ng/ml) | $94\pm5*$ | $100 \pm 6*$ |
| Proliferation (cells/well) | Vehicle | 10±1 | 10±1 |
| | PlGF (100 ng/ml) | 11 ± 2 | 8±2 |
| | VEGF (100 ng/ml) | $34 \pm 2*$ | $15 \pm 1*$ |
| | VEGF + PlGF (100 ng/ml each) | 35 ±2* | 38 ±2* |
| | bFGF (50 ng/ml) | $39 \pm 3*$ | 43 ±2* |
| | bFGF (50 ng/ml) + PIGF (100 ng/ml) | $34 \pm 2*$ | $39\pm2*$ |

Table 1; Role of PIGF and VEGF in endothelial survival, migration and proliferation

Data represent the mean \pm s.d. of 9 to 12 experiments. Migration and proliferation are expressed in absolute number of migrating and proliferating cells; apoptosis is expressed as a percent of the control (vehicle-treatment). Anti-mPIGF, anti-murine PIGF antibodies; ND, not determined. *, P < 0.05 versus control (vehicle).

Reduced collateral growth in *Pgf*^{-/-} mice

We studied growth of collateral arteries after ligation of the femoral artery. Collaterals in non-occluded limbs were small in both genotypes, but became tortuous and enlarged in wild-type but not Pgf^{-} mice after ligation (Fig. 4a-c). PIGF levels were 45% higher in ligated vessels than in control vessels (4.5 ± 0.9 pg PIGF/mg protein versus 3.1 ± 0.9 pg PlGF/mg protein; n = 4; P < 0.05 by paired t-test). Macrophages, known to play an essential role in collateral growth²⁶, infiltrated more collaterals in wild-type than in Pgf' mice (68% of 66 wild-type collaterals versus 43% of 67 $Pgf^{/2}$ collaterals; P < 0.05 by χ -square analysis; n = 5 mice; Fig. 4d and e). Macrophages may have been recruited by PIGF, produced by activated endothelial cells, as PIGF is known to be chemoattractive for monocytes/macrophages²⁷. Activated macrophages themselves may be another source of PIGF in growing collaterals, as they produce PIGF in culture ($8 \pm 2 Pgf$ per 1 × 10³ hprt transcripts; n = 5) and stained strongly for PIGF in vivo (Fig. 4h). During collateral growth, extravasated fibronectin provides a scaffold for migrating smooth muscle cells. Fibronectin leaked out in 70% of 80 wild-type collaterals, but in only 25% of 83 Pgf' collaterals after ligation (n = 5 mice; P < 0.05 by γ -square; Fig. 4f and g). A role for PIGF in vascular permeability was further demonstrated in other assays (see below). Thus, PIGF may stimulate collateral growth by affecting monocyte recruitment, plasma extravasation and growth of endothelial and smooth muscle cells (see below).

Fig. 4 Impaired collateral growth in Pgf^{/-} mice, a and b, Macroscopic view of collateral arterioles (visualized by bismuth gelatinography; arrowheads) in the hindlimb after ligation of the deep femoral artery (FA), revealing enlarged, tortuous collaterals in wildtype (a) but not in Pgf⁻⁽⁻(b) mice, c, Quantification of the lumen area of the collateral vessels in the non-occluded and occluded limb, and the rescue of adaptive arteriogenesis by transplantation of wild-type bone marrow in Pgf^{-/-} mice (+/+ --> -/-). s.e.m. of at least 10 mice. *, P< 0.05 versus non-ligated. d and e, Mac3 immunos-taining, revealing more numerous macrophages accumulating and infiltrating in wild-type (d) than in $Pgf^{/-}(e)$ collaterals 3 d after occlusion, f and g, Fibronectin immunostaining, revealing larger extravasated fibronectin deposits around wild-type (f) than around $Pgf^{/-}(g)$ collaterals 3 d after occlusion. h, PIGF immunostaining of macrophages in a human brain infarct. Scale bars: and 50 µm (d-g), 10 µm (h).





Loss or inhibition of PIGF reduced plasma extravastion

Recombinant PIGF stimulates vascular leakage⁵, but no role for endogenous PIGF in plasma extravasation has been identified. Topical administration of VEGF on microvessels in cremaster muscle stimulated extravasation of fluorescent dextran in wild-type mice, but only minimally in Pgf^{-} mice (Fig. 5a-e). Notably, histamine induced a comparable response in both genotypes, indicating that PIGF specifically amplified the effect of VEGF (Fig. 5e). We observed a similar reduced plasma extravasation in Pgf^{-} mice in the Miles assay during skin wound healing, after subcutaneous implantation of a perforated polyethylene ball or in a model of delayed skin hypersensitivity (data not shown). PIGF levels were upregulated in endothelial cells and in hyperplastic keratinocytes in wounded or inflamed skin and increased from lower than 1 pg/mg protein in normal to 27 ± 6

pg/mg protein in wounded skin (n = 5; P < 0.005). VEGF was comparably upregulated in both genotypes (2 ± 1 pg/mg versus 97 \pm 15 pg/mg in wild-type mice; n = 5; P < 0.05; and 4 ± 1 pg/mg versus 108 \pm 15 pg/mg in $Pgf^{/-}$ mice; n = 5; P < 0.05), whereas endothelial VEGFR-1 expression was induced in wounded skin (data not shown). Few specific therapeutic strategies are currently available to block plasma extravasation. Neutralizing anti-PIGF antibodies reduced vascular leakage in wild-type mice after application of mustard oil on their ears (relative absorbance units × 1000/ear: 13 \pm 3 after mineral oil; 53 \pm 7 after mustard oil plus control IgGs; 26 \pm 5 after mustard oil plus anti-PIGF; n = 5; P < 0.05 versus control antibodies; Fig. 5f and g). We obtained similar results after intradermal injection of VEGF (data not shown). Recombinant PIGF restored the reduced plasma extravasation in $Pgf^{/-}$ mice (data not shown).

Role of PIGF in mobilization of bone-marrow-derived cells

VEGF recruits bone-marrow-derived cells to sites of pathological angiogenesis, but the role of PIGF remains undefined^{28,29}. Because bone-marrow-derived cells contribute to capillary ingrowth in matrigel implants (data not shown), mice were transplanted with congenic bone marrow and capillary ingrowth in matrigel, supplemented with the VEGF₁₆₅ isoform, and quantified by measuring the hemoglobin (Hb) content per implant. Matrigel angiogenesis was abundant when both the donor bone marrow and the host vessels produced PIGF (0.29 ± 0.05 g Hb/dl; n = 7), but minimal when the bone marrow and host lacked PIGF (0.05 ± 0.01 g Hb/dl; n = 7; P < 0.05). Capillaries still infiltrated the ma-trigel in wild-type mice transplanted with Pgf' bone marrow (0.26 ± 0.07 g Hb/dl; n = 9), indicating that PIGF production by vessel-wall-associated endothelial cells was sufficient for angiogenesis still occurred (0.17 ± 0.07 g Hb/dl; n = 7), indicating that production of PIGF by bone-marrow cells can stimulate angiogenesis at distant sites. Transplantation of wild-type bone marrow in Pgf'^{-} mice also rescued the impaired collateral enlargement after ligation of the femoral artery, possibly by mobilizing PIGF-producing monocytes/macrophages to the collaterals (Fig. 4c). PIGF specifically modulated the VEGF response, as angiogenesis in matrigel supplemented with bFGF was comparable in both genotypes (0.28 ± 0.02 g Hb/dl in Pgf'^{-} mice; n = 15; P = NS).

PIGF modulates the endothelial response to VEGF

To determine whether the amplification of the VEGF response by PIGF resulted from direct stimulation of endothelial cells, we studied capillary outgrowth in intact aortic rings. We used $Pgf^{/-}$ aortic rings and serum to avoid effects of endogenously produced PIGF. At 20 ng/ml, PIGF or VEGF alone minimally stimulated capillary outgrowth (< 1 capillary/aorta). However, a combination of VEGF and PIGF strongly stimulated capillary outgrowth (8 ± 1 capillaries/aortic ring; 4 ± 2 branches/capillary; 0.37 ± 0.05 µm capillary length; n = 9). A higher concentration of VEGF (60 ng/ml) stimulated capillary outgrowth from $Pgf^{/-}$ aortic rings, indicating that PIGF amplified, but did not determine the VEGF response (13 ± 2 capillaries/aortic ring; 8 ± 2 branches/capillary; 0.35 ± 0.02 µm capillary length). A high concentration of PIGF (50 ng/ml) stimulated capillary outgrowth (13 ± 3 capillaries/aortic ring; n = 5). Because PIGF alone (in the absence of VEGF) did not stimulate isolated endothelial cells (see below), PIGF probably stimulated capillary outgrowth by amplifying endogenous VEGF in aortic rings. We observed a similar amplification of the VEGF-response by PIGF in other endothelial cells (data not shown).

We also observed the interaction between VEGF and PIGF using capillary endothelial cells from both genotypes (Table 1). VEGF stimulated the migration, growth and survival of wild-type cells. Compared with the wild type response, VEGF stimulated migration by 25%, proliferation by 40% and survival by 7% in Pgf' cells. However, both genotypes responded comparably to bFGF. A high concentration of PIGF alone did not affect the response of wild-type endothelial cells to VEGF, presumably because these cells produced sufficient PIGF (data not shown). However, even at low doses, PIGF dose-dependently restored the impaired VEGF-response of Pgf' cells to a combination of VEGF plus PIGF, indicating that PIGF activated VEGFR-1. VEGF/PIGF het-erodimers also stimulated the survival of Pgf' endothelial cells (Table 1). Thus, PIGF, though ineffective by itself, amplifies the VEGF response by activating VEGFR-1.

Fig. 5 Modulation of permeability and of VEGF-response by PIGF. *a-d* Intravital microscopy, revealing extravasation of fluorescent dextran from microvessels in the cremaster muscle at 0 (a and c) and 60 (*b* and *d*) min after topical application of VEGF in wild-type (a and *b*) but not in Pgf^{-c} (*c* and *d*) mice. Scale bars: 50 µm (*a-d*). *e*, Quantification of extravasated dextran, revealing extravasation in response to VEGF and histamine in wild-type mice, but only to histamine in Pgf^{-c} mice. Inwild-type, *Pgf^{-c}* mice. *Pgf^{-c}* mice. *Pgf^{-c}* mice. *Pgf^{-c}* mice is reduced by antibodies against PIGF (*g*) but not by control IgGs (f). *h*, Schematic model of the role of VEGF, PIGF, VEGFR-1 and VEGFR-2. In the embryo, VEGF induces angiogenesis by activating VEGFR-2. VEGFR-1 is primarily expressed as an inhibitory soluble receptor (sVEGF-R1) which functions as a 'sink' for VEGF, with only minimal levels of membrane-associated VEGFR-1. *i*, In pathological conditions, PIGF is upregulated and stimulates angiogenesis via displacement of VEGF from VEGF-R1 to VEGF-R2 and via activation of VEGFR-1, of which a larger fraction is membrane-associated. PIGF-mediated activation of VEGFR-1 synergizes with VEGF-mediated activation of VEGFR-2 to induce a stronger angiogenic response. Dotted lines represent low expression; solid lines denote high levels of expression. The inhibitory pathway is indicated in red.



Upregulation and signaling of membrane-associated VEGFR-1

The inhibition of the endothelial response to PIGF by VEGFR-1 antibodies indicated that VEGFR-1 transmitted intracellular signals. For PIGF to cause signaling, VEGFR-1 must be membrane-associated. In mid-term placentas, only $2 \pm 0.5\%$ of total VEGFR-1 was membrane-associated, whereas $30 \pm 2\%$ of VEGFR-1 was membrane-bound in ischemic myocardium (n = 3; P < 0.005; Fig. 1a). Quantitative RT-PCR analysis confirmed the increased membrane-localization of VEGFR-1 during pathological angiogenesis (data not shown). To examine whether PIGF induced intracellular signaling, we studied the role of the Src-family kinases in the response of Pgf' endothelial cells to VEGF and PIGF (Table 1). Src-kinases have recently been implicated in responses to VEGF (refs. 30,31), but their role in the response of Pgf' endothelial cells to VEGF, but completely blocked the PIGF-dependent amplification of the VEGF response in Pgf' endothelial cells.

DISCUSSION

PIGF and VEGFR-1 have received little attention in angiogenesis research³. The present analysis indicates that PIGF—via activation of VEGFR-1—specifically potentiates the angiogenic response to VEGF, but not to bFGF. In contrast to the essential role of VEGF in physiological and pathological angiogenesis^{1,2,32}, the role of PIGF is restricted to pathological conditions and is therefore a possible target for therapy.

Several mechanisms might explain why PIGF had a negligible role in vascular development (Fig. 5h and *i*). First, PIGF may be unable to displace VEGF from VEGFR-1, because of its lower expression levels in the embryo and its lower affinity for VEGFR-1 compared with VEGF (ref. 3). Second, VEGF was upregulated in compensation for the absence of PIGF. Third, the minimal levels of membrane-associated VEGFR-1 in the embryo might be insufficient to transmit PIGF-dependent angiogenic signals. VEGFR-1 is predominantly soluble and antibodies against VEGFR-1 stimulate angiogenesis in embryos, confirming that VEGFR-1 is primarily an inert 'sink' for VEGF during development^{11,12}. Fourth, the need for amplification of the VEGF response by PIGF in the adult might be explained by a requirement for greater responses to VEGF to mediate pathological rather than embryonic angiogenesis. In at least several pathological conditions, VEGF and PIGF levels exceeded those found in the embryo. VEGF-B did not rescue vascular development of *Pgf*^{-/-} embryos, even though it also binds to VEGFR-1 (refs. 9,10), raising the question of whether the individual VEGFR-1 ligands have distinct functions *in vivo*.

Previous studies reported that exogenously administered PIGF stimulates angiogenesis or permeability in particular conditions^{5,6}, but they did not evaluate the endogenous role of PIGF in the adult. The present findings indicate that by affecting vascular growth and remodeling, PIGF contributes to the pathogenesis of several disorders with high morbidity. Although our findings do not exclude the possibility that PIGF activates endothelial cells independently of VEGF (refs. 15,33), PIGF was found to stimulate angiogenesis by synergizing with VEGF. When new blood vessels form in the adult endothelial cells become more responsive to VEGF by upregulating PIGF and VEGFR-1, (Fig. 5h and *i*). Apart from this autocrine pathway, PIGF from paracrine sources (malignant cells, macrophages, ischemic cardiomyocytes and so forth) might also enhance the angiogenic activity of VEGF. PIGF might increase tumor angiogenesis depending on whether tumors up-regulate PIGF (refs. 8, 20-22). The effect of PIGF on the vasculature was quantitatively and qualitatively dependent on the tissue microenvironment. For instance, PIGF stimulated endothelial growth in ischemic retina, but promoted arterial remodeling in ischemic limbs. In addition, depending on the tumor, PIGF increased the number or the size of new vessels. A similar tissue-specific role for VEGF is now being recognized⁵, even though the underlying mechanisms remain largely undefined.

Several mechanisms might explain the role of PIGF in pathological, but not physiological angiogenesis. First, PIGF and VEGFR-1 are minimally expressed in adult quiescent vasculature, but are markedly upregulated during pathological conditions. Second, PIGF might enhance the angiogenic response to VEGF by forming VEGF/PIGF heterodimers, which have been detected in tumors and are upregulated by hypoxia^{16,17}. Third, PIGF activates VEGFR-1 to transmit angiogenic signals-as supported by our findings that antibodies against VEGFR-1 blocked the angiogenic activity of PIGF in vitro (this study) and in vivo (manuscript in preparation). If the only function of PIGF was to 'fill up an inert VEGFR-1 sink', antibodies against VEGFR-1 should stimulate, not inhibit, angiogenesis. The increased membrane-localization of VEGFR-1 in pathological angiogenesis, as compared with embryonic angiogenesis, and the inhibition of PIGF by the Src-kinase inhibitor are also consistent with a role for PIGF and VEGFR-1 in angiogenic signaling. Rather than contradicting the previous suggestion that PIGF stimulates angiogenesis by displacing VEGF from the 'VEGFR-1 sink^{5,6}, our data provide evidence for an additional role of PIGF in transmitting angiogenic signals. Because both mechanisms could be operational at the same time (for example, PIGF displaces VEGF from VEGFR-1 and activates VEGFR-1), it remains to be determined to what extent and under which conditions VEGFR-1 is a decoy and/or a signaling receptor for PIGF. Fourth, PIGF is a chemoattractant for inflammatory cells, a hallmark of pathological angiogenesis and collateral growth^{25,32}. The reduced recruitment of macrophages likely contributed to the reduced collateral growth in Pgf^{-/-} mice, whereas transplantation of wild-type bone marrow rescued their collateral expansion. Finally, PIGF might also stimulate pathological angiogenesis by mobilizing bone-marrowderived cells. Whether PIGF affects mobilization of bone-marrow-derived an-gioblasts or hematopoietic stem cells remains unknown.

Conditional VEGF gene-inactivation and inhibitor studies in adult mice have revealed that the adult quiescent vasculature becomes less dependent on VEGF for its maintenance³⁴. However, during pathological conditions— such as ischemia, inflammation or malignancy—angiogenic endothelial cells are stimulated by increased VEGF levels. The molecular mechanisms by which adult endothelial cells are induced to rapidly grow ('angiogenic

switch') are still incompletely understood. Our findings indicate that PIGF might contribute to this switch by amplifying VEGF. An outstanding question is therefore whether PIGF could be used for therapeutic angiogenesis of ischemic cardiovascular disease, or whether inhibition of PIGF might suppress pathological angiogenesis in tumors, inflammatory disorders and diabetic retinas (manuscript in preparation). The present findings that anti-PIGF antibodies blocked permeability indicate a novel treatment strategy for vascular leakage syndromes as well.

METHODS

Generation of *Pgf*^{/-} mice.

To inactivate the Pgf gene in embryonic stem (ES) cells, pPNT. *Pgf* was constructed containing a 8-kb Ncol fragment (exons 1 and 2 and part of exon 3 until nucleotide 265 of the murine *Pgf* cDNA), a neomycin phosphotransferase (neo) cassette, and a 5-kb BamHI fragment (exon 7). Probes for Southern-blot analysis included a 380-bp fragment obtained by PCR using 5'-GAATTCAATGAGTTAAGGGTG -3' and 5'-AATACTACAGTTATAGACTA -3' primers and a neo probe. *Pgf^{-/-}* ES clones were obtained by culturing *Pgf^{-/-}* ES clones in 1800 μ g/ml G418. Targeted ES clones were used to generate chimeric mice, which were test-bred with Swiss females for germline transmission as described³⁵.

Gene expression, morphology, antibodies and embryo culture.

Western and northern blotting, real-time RT-PCR, immunostaining and in situ hybridization were performed as described³⁵. VEGFR-1 was immunoblotted after incubation with heparin-sepharose using VEGFR-1-specific antibodies (Sigma; clone Flt11). To demonstrate binding of VEGF and PlGF to soluble VEGF-R1, 500 µg placental extract was incubated with 500 ng VEGF and 1 µg PlGF-2 for 1 h at room temperature, immunoprecipitated with anti-VEGF and anti-PIGF antibodies and immunoblotted for VEGF-R1. Radioligand binding studies with $[^{125}I]$ -labeled VEGF₁₆₅ were performed as described", except that a 60-fold molar excess of cold PIGF or VEGF was used as competitor. The following antibodies were used for immunostaining: rat antibody against mouse VEGFR-1 (MF1; Imclone, New York, New York; characterization will be reported elsewhere: goat antibody against mouse VEGFR-2 (R&D Systems, Abingdon, UK), rat antibody against mouse VEGFR-2 (DC101; Imclone)³⁶; and rabbit antibody against human PIGF (#PA211, ReliaTech; Braunschweig, Germany). A murine PIGF-specific ELISA was set up, using goat antibody against murine PIGF antibodies (R&D), but murine VEGF was quantified with a commercially available ELISA (R&D). The following primers were used for RT-PCR of Pgf. 5'-TTCAGTCCGTCCT-GTGTCCTT-3'; 5'-GCACACAGTGCAGACCTTCA-3'; and 5'-AC-CACAGCAGC CACTACAGCGACTCA-3'. Expression levels of Pgf were normalized for hprt. The following immunoneutralizing antibodies were used: VEGFR-2 (DC101, Imclone)³⁶; VEGFR-1 (MF1, Imclone); PIGF-2 (R&D); and isotype-matched control IgGs (Imclone and ICN, Costa Mesa, California)³⁷. The recombinant proteins used were PIGF-2 (ReliaTech), VEGF₁₆₅ and VEGF/PIGF heterodimer (both from R&D).

Models of angiogenesis.

Morphometry of angiogenesis in neonatal mice³⁸, transplantation of congenic bone marrow (C57BI6 background)²⁵, ingrowth of capillaries in matrigel (containing VEGF or bFGF)³⁹ and analysis of ES-cell-derived⁴⁰ and fibrosarcoma⁴¹ tumors were described. Luteal angiogenesis was analyzed on toluidine blue-stained semi-thin sections of ovaries 4.5 d after mating. Ischemic retinal neovascularization was induced by exposing P7 neonatal mice to hyperbaric (80%) oxygen for 5 d and subsequently to room air for another 5 d, and then analyzed by fluorescent retinal angiography and counting endothelial cell on retinal cross-sections. Venous dilatation and arterial tortuosity were semi-quantitatively scored (scale: 0-3). Vascular remodeling during skin wound healing was analyzed within 4 d after applying a 15-mm, full-thickness skin wound⁴². To induce limb ischemia, the right femoral artery was occluded distal to the branch site of the deep femoral and the popliteal artery. After 7 d, mice received 0.1% adenosine, were perfused with fixative and bismuth-gelatin contrast medium for angiography. Collaterals in the adductor muscle were used for morphometry. Myocardial infarction and infarct revascularization were performed as described²⁵. Human PIGF-1 (10 µg active dimer; Geymonat SpA, Anagni, Italy) was delivered using an osmotic minipump (Alzet, type 2001, Someren, the Netherlands).

Vascular permeability.

Mustard oil was painted on the ears of Swiss mice, and extravasation of Evans blue determined⁴³. Antibodies were injected i.v. (300 μ g/kg) 30 min before injection of Evans blue and application of mustard oil. For intravital analysis, the mouse cremaster muscle was exposed, and a flexible ovoid ring introduced into the cremaster pouch

to allow transillumination of the muscle, covered with carbogen-bubbled prewarmed mineral oil^{44} . Within 10 min after i.v. injection of FITC-labeled dextran, 150,000 (100 mg/kg), 10 µl of VEGF₁₆₅ (260 nM) or histamine (500 nM) was topically administered. Video images of fluorescent microcirculation were recorded throughout the experiment for off-line analogous video image processing of the light intensity in a selected area of the video picture, expressed as gray values (arbitrary units).

In vitro angiogenesis assays.

Capillary outgrowth from cultured rings of mouse aorta in gels of rat-tail interstitial collagen (1.5 mg/ml) was performed as described for the rat⁴⁵, except that 2.5% mouse serum was used. To quantify the number, length and branching of vessels, a scanned image of the aortic ring was converted to a binary image (aortic ring and capillaries in blue with black background; fibroblasts were subtracted). To obtain mouse capillary endothelial cells, mice were injected s.c. with 500 µl of matrigel containing bFGF (100 ng/ml) and heparin (100 µg/ml). After 7 d, the matrigel pellet was enzymatically dispersed and endothelial cells were cultured in M131 medium supplemented with 5% MVGS (Gibco-BRL). Cells were starved in medium with 0.5% serum for 24 h, after which they were stimulated with human VEGF₁₆₅ and/or murine PIGF-2, or bFGF (all from R&D) for 24 h. Cultures were then analyzed for the total cell number (proliferation) or the number of cells migrated after scrape-wounding (migration). For apoptosis studies, cells were cultured in RPMI 1640 medium containing 10% FCS (Life Technologies), 100 µg/ml heparin and 30 µg/ml endothelial cell growth supplement. Apoptosis was induced by withdrawal of growth factors (0.1% FCS) and quantified by measuring cytoplasmic his-tone-associated DNA fragments (mono- and oligonucleosomes) as described³⁷. Pyrazolopyrimidine (PP2) was from Calbiochem (Beeston Nottingham Forest, UK).

Acknowledgments

We thank P. Schaeffer, K. Bijnens, A. Bouché, S. De Cat, M. De Moi, K. De Roover, E. Gils, B. Hermans, S. Jansen, L. Kieckens, A. Manderveid, K. Maris, A. Sahii, T. Vancoetsem, A. Vandenhoeck, P. Van Wesemael and S. Wyns for assistance. This work was supported in part by the European Community (Biomed BMH4-CT98-3380), Actie Levenslijn (#7.0019.98), F.W.O. (GO12500), AIRC and ISS (Programma Nazionaie AIDS 1998), the Deutsche Krebshilfe 10-1302-Ri 3 and BMBF 01 KV9922/6.A. Noel and L Devyare supported by the F.N.R.S.

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