

## Differential Expression of Placenta Growth Factors and Their Receptors In the Normal and Pregnancy-Induced Hypertensive Human Placentas

Placental development requires extensive angiogenesis and the invasion of the maternal decidua by the trophoblasts. Adequate and organized interaction of vascular endothelial growth factors (VEGF), placenta growth factors (PlGF), and their receptors are essential for a normal development and function of the placenta. In this study, we evaluated the expressions of PlGFs and their receptors, mRNAs by Northern blotting, in situ hybridization and RT-PCR in the normal and pregnancy-induced hypertensive (PIH) placentas. The expression level of PlGF-2 mRNA was lower in the PIH placentas compared to control as assessed by Northern blotting and in situ hybridization. PlGF mRNA was mainly localized to the vasculosyncytial membrane of placental villi and villous stroma. The expression of PlGF receptor-1 (PlGFR-1) was significantly increased in the PIH placentas compared to the normal ones. These results suggest that the alteration of PlGF-2 and PlGFR-1 mRNA expressions in the placenta are related to the pathogenesis of PIH.

Key Words : Growth Factors; Receptors, Growth Factor; Placenta; Human; Pre-Eclampsia

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## INTRODUCTION

Pregnancy-induced hypertension (PIH) usually develops during the second half of the pregnancy and accompanies proteinuria and edema. PIH is estimated to affect 7 to 10% of all pregnancies in the United States (1), making common complications associated with fetomaternal death and morbidity.

While the etiology of PIH remains unclear, it is known to be associated with endothelial damage and dysfunction (2). There are also an increased concentration of vasoconstrictive agents such as endothelin (3), increased capillary permeability (4), and the endothelial damage is also associated with neutrophil (5).

An impaired trophoblastic invasion of the maternal placental bed is considered to be the initiator of endothelial damage in PIH (6). This defect might result in poor fetoplacental perfusion along with a release of factors into the maternal circulation, leading to a widespread endothelial damage and the resulting clinical symptoms of PIH (7). This placental dysfunction could be mediated by the factors present in the maternal circulation as a result of impaired trophoblastic invasion

(6). Angiogenic growth factors are also thought to play a substantial role in the placentation, and can be strong endothelial cell activators (8).

Recently, placenta growth factor (PlGF), a dimeric secreted factor that shares close amino acid homology (53% identity) to vascular endothelial growth factor (VEGF), was known as a potent angiogenic growth factor capable of inducing the proliferation, migration, and activation of endothelial cells (9). Unlike VEGF, abundant expression of PlGF is restricted to the placenta (10). Alternative splicing of *PlGF* mRNA produces at least three polypeptides, PlGF-1, PlGF-2 and PlGF-3, which have different secretory patterns, heparin-binding affinities and dimerization properties (10, 11).

Ideal placental development requires an adequate and organized interaction of vascular growth factors and their receptors, including VEGF and PlGF; both VEGF and PlGF act through their common tyrosine kinase receptors (12). The expression of VEGF and KDR (PlGFR-2/VEGFR-2) is strongest during early gestation and decrease as pregnancy advances (13-15) while PlGF and Flt-1 (PlGFR-1/VEGFR-1) increase towards term (16, 17). In the present study, we examined the expression patterns and localization of PlGFs and their recep-

tors in the normal and PIH human placentas at term to get insight into their relation to PIH.

## MATERIALS AND METHODS

### Tissue collection and preparation for in situ hybridization

Placentas used for analyses were collected from 8 normal [37 (4 cases), 38 (2 cases), 39 wk (2 cases)] and 5 PIH [36 (2 cases), 37, 38, 39 wk] mothers at caesarean section delivery in Gyeongsang National University Hospital (GNUH). The Human Subject Research Committee of GNUH approved these human tissues for the present study. The parameters to define PIH were systolic, and diastolic blood pressures above 140 and 90 mmHg, respectively, in at least two consecutive measurement, and proteinuria of  $>0.3$  g/24 hr, in women after 20-week of gestation. The placentas (approximate weight: 100 mg) were snap frozen in liquid nitrogen immediately after collection for total RNA extraction. For in situ hybridization, the placentas were dissected on ice, cut into cubes fit into slide glass, and washed with phosphate buffered saline (PBS, pH 7.0). And then those tissues were post-fixed by a 48 hr-immersion in 4% neutrally buffered paraformaldehyde and cryoprotected by immersion in 20% sucrose in PBS for 24 hr. The tissues were frozen using OCT compound (Sakura Finetek, CA, U.S.A.), and 12  $\mu$ m sections were cut with a cryostat (Leica, Heidelberg, Germany) and thaw-mounted on gelatin-coated slides, dried and stored at  $-70^{\circ}\text{C}$  until use.

### Hematoxylin and eosin staining

The frozen sections were washed with tap water for 5 min, immersed in hematoxylin for 2 min and checked for complete staining in tap water. Eosin staining was carried out for 3 min. Sections were dehydrated through a graded series of alcohol (70 to 100% ethanol, 3 min each), cleared in xylene, cover-slipped and observed with a light microscope.

### Northern blot analysis

Total RNA was extracted from placenta, using acid guanidium thiocyanate-phenol chloroform method, and quantified by UV spectrophotometer. Twenty micrograms of total RNA was subjected to electrophoresis in a 1.2% agarose-formaldehyde gel and then capillary-transferred to a nylon membranes (Nytran, pore size: 0.45  $\mu$ m, Schleicher & Schuell, Keene, NK). The partial cDNAs of human PIGF-1/-2 were obtained from human placenta using RT-PCR and cloning. Then we confirmed PIGF cDNAs after sequence analysis and used as template for probe synthesis. Probes were labeled by random priming with  $\alpha$  [ $^{32}\text{P}$ ]-dCTP, and purified with a Sephadex G-50 nick column (Pharmacia Biotech, Uppsala,

Sweden). Hybridization was carried out for 16-18 hr at  $42^{\circ}\text{C}$  in 20 mL of hybridization buffer [50% Deionized Formamide, 5  $\times$ SSPE (20  $\times$ SSPE: 3 M NaCl, 0.2 M  $\text{NaH}_2\text{PO}_4$ , and 20 mM EDTA, pH 7.4), 5  $\times$ Denhardt's solution (1  $\times$ Denhardt's solution, 0.02% polyvinyl-pyrrolidone, 0.02% Ficoll, 0.02% BSA), 0.1% SDS, 1 mg/mL heated-denatured salmon sperm DNA]. Membranes were washed in 2  $\times$ SSPE with 0.1% SDS at room temperature for 10 min twice and in 0.1  $\times$ SSPE with 0.1% SDS at  $42^{\circ}\text{C}$  for 10 min. The membranes were exposed to radiography film (Fuji RX & Fuji Co, Tokyo, Japan) for 3 days at  $-70^{\circ}\text{C}$ .

### In situ hybridization

The  $^{35}\text{S}$ -UTP-labeled probes were generated by *in vitro* transcription from partial cDNA clone amplified from PIH placentas. The probes were purified with Sephadex G-50 nick column and radioactivity was measured on a  $\beta$ -counter. Pre-hybridization was carried out for 1 hr at  $37^{\circ}\text{C}$  in pre-hybridization buffer [50% Deionized Formamide, 0.25 M NaCl, 1  $\times$  Denhardt's solution, 10 mM Tris (pH 8.0), 1 mM EDTA (pH 8.0), 10% Dextran Sulfate, yeast tRNA (250  $\mu$ g/mL), 50 mM DTT]. Hybridization of the slides with anti-sense  $^{35}\text{S}$ -UTP-labeled RNA transcripts ( $7 \times 10^5$  cpm per section) in prehybridization buffer was performed at  $60^{\circ}\text{C}$  for 20 hr. Sections were washed in 4  $\times$ SSC, treated with 20  $\mu$ g/mL of RNase A, washed in 2  $\times$ SSC, and followed with a further wash in 0.1  $\times$ SSC containing 1.0 M DTT at room temperature. Sections were dehydrated through a graded series of alcohol (50% to 100% ethanol) and air dried at room temperature, then autoradiographed for 7 days at  $-70^{\circ}\text{C}$  using Amersham  $\beta$ -max hyperfilm (Amersham, NJ, U.S.A.). Slides were then dipped in NTB2 autoradiographic emulsion (Kodak, NY, U.S.A.) and exposed in the dark at  $4^{\circ}\text{C}$  for 14 days, and developed with D-19 developer, fixed with rapid fixer (both from Kodak, NY, U.S.A.), counter-stained with cresyl violet, and mounted.

### RT-PCR analysis of PIGFR-1 and PIGFR-2 mRNAs

Complementary DNA was synthesized in a mixture consisting of 1  $\mu$ g total RNA from each placenta, 200 unit of RNaseH Muloney murine leukemia virus (MMLV) reverse transcriptase (Promega, Madison, WI, U.S.A.) in 20  $\mu$ L of the RT reaction mixture [150 pmol of oligo d(T), 40 units of RNase inhibitor, 50 mM Tris-HCl (pH 8.3), 75 mM KCl, 3 mM  $\text{MgCl}_2$ , 1 mM dNTPs (N=A, T, G, and C), and 10 mM DTT] at  $37^{\circ}\text{C}$  for 1 hr, followed by 5 min at  $95^{\circ}\text{C}$ . After heat inactivation, cDNA (total 20  $\mu$ L) was stored at  $-20^{\circ}\text{C}$  until PCR.

The PCR reaction mixture (50  $\mu$ L) containing 10 mM Tris-HCl (pH 8.3), 50 mM KCl, 25 mM  $\text{MgCl}_2$ , 2 units of *Taq* polymerase, 0.2 mM dNTPs, 4  $\mu$ L of each placental cDNA, and 10 pmol of each specific PCR primer for

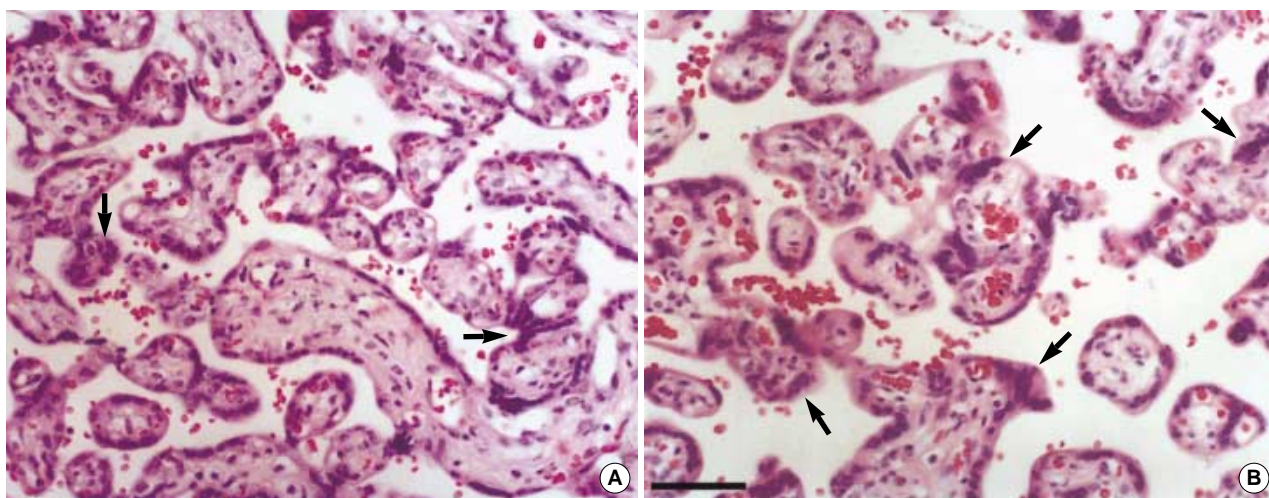


Fig. 1. Photographs of H & E staining in the normal and PIH (pregnancy induced hypertension) placentas. Note the thickening of syncytial membranes, an increased number of syncytial knots (arrow) of the placenta villi in the PIH placenta as compared to control. (A) normal placenta; (B) PIH placenta. Scale bar = 100  $\mu$ m.

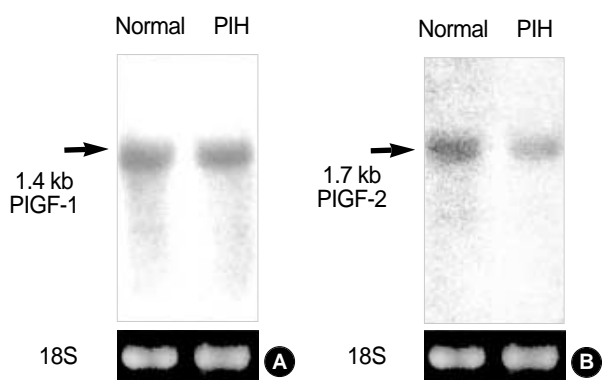


Fig. 2. Expression of *PIGF-1* and *PIGF-2* mRNA in the normal and PIH (pregnancy induced hypertension) placentas as assessed by Northern blot analyses. Northern blotting shows a 1.4- (*PIGF-1*) and a 1.7-kb (*PIGF-2*) transcript in the normal and PIH placenta after a hybridization with *PIGF-1*- and *PIGF-2*-specific cDNA probes, respectively. *PIGF2* mRNA is markedly decreased in the PIH placenta compared to the normal placenta. Twenty micrograms of total RNA from the pool of 8 normal and 5 PIH placentas was applied to each lane, and the consistency in loading is demonstrated by ethidium-bromide staining of 18S rRNA. *PIGF-1*, placenta growth factor -1; *PIGF-2*, placenta growth factor -2.

*PIGFR-1* (upstream; 5'-TGTC AATGTGAAACCC CAGA-3', downstream; 5'-TTTCTTCCACAGTCCCAAC-3') and *PIGFR-2* (upstream; 5'-TGATGTCC CACCC CAGATT-3', downstream; 5'-ACATTTGCCGCTTGGATAAC-3') were added. The PCR reaction mixture was subjected to 30 cycles of amplification using a PCR system with denaturation at 94 °C for 1 min, primer-annealing at 55 °C for 1 min, and primer-extension at 72 °C for 2 min. PCR products (10  $\mu$ L) were subjected to electrophoresis in a 1.2% agarose gel in TAE buffer, visualized by UV illumination after ethidium bromide staining, and photographed using Polaroid 667 film (Hert-

fordshire, U.K.).  $\beta$ -actin was used as a internal control for procedural variation. The intensity of PCR bands was measured densitometrically and analyzed using SigmaGel (version 1.0, Jandel Scientific Software) software.

#### Statistics

RT-PCR signals were normalized to their  $\beta$ -actin signals. One-way analysis of variance was determined using Graphpad Instat Software (Version 1.15). A  $p$  value < 0.05 was deemed statistically significant; data are expressed as mean  $\pm$  SEM.

## RESULTS

### H & E staining in the normal and PIH placentas

Representative hematoxylin and eosin-stained sections of the normal and PIH placentas were shown in Fig. 1. In the PIH placenta, specific histologic changes were seen compared to control; the syncytial membranes were continuously expanded and the number of syncytial knots was increased in the PIH placental villi.

### Expression of *PIGF-1* and *PIGF-2* mRNAs in the normal and PIH placentas

Northern blot analyses were conducted to evaluate differences in the abundance of *PIGF-1* and *PIGF-2* mRNA in normal and PIH placentas. Both *PIGF-1* and *PIGF-2* mRNAs were less abundant in PIH placenta compared to normal (Fig. 2); of two genes, the *PIGF-2* was markedly decreased (Fig. 2B).

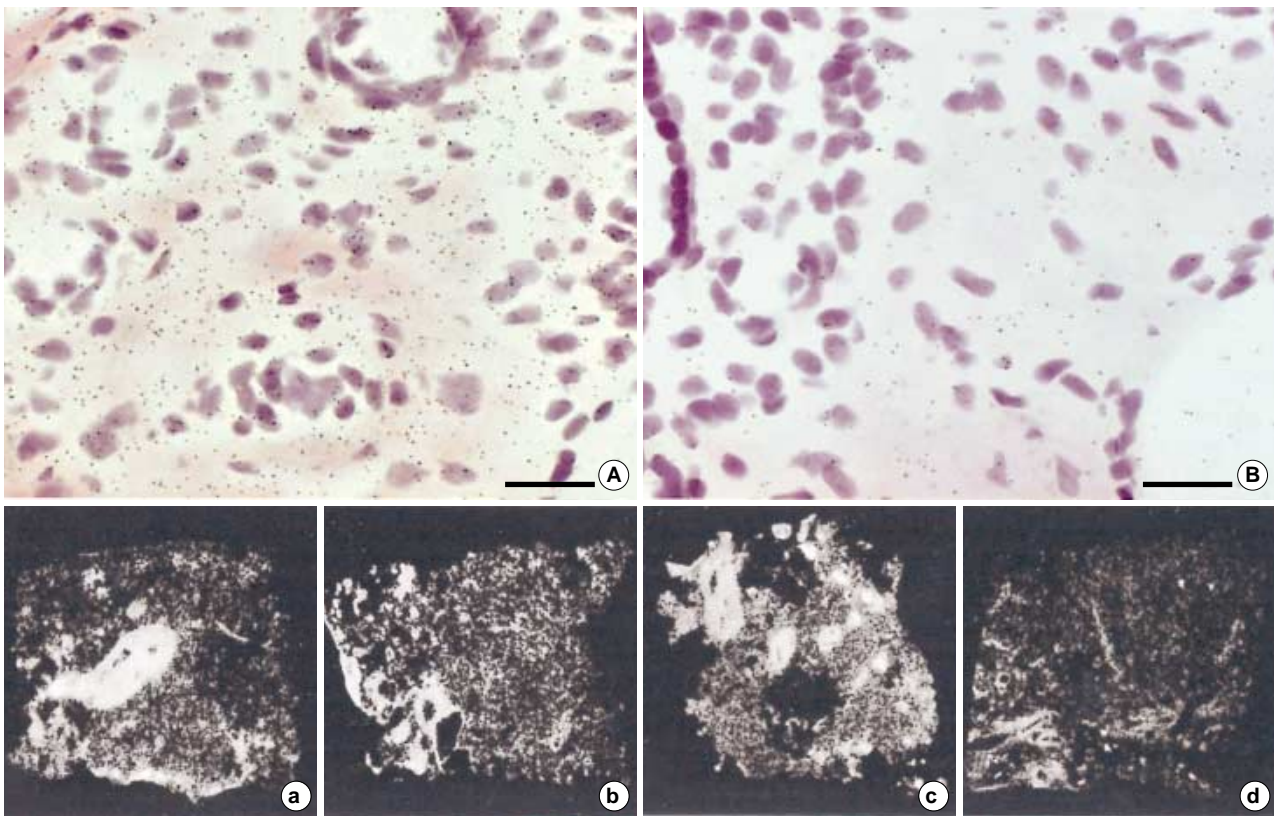


Fig. 3. Localization of *PIGF-1* and *PIGF-2* mRNA in the normal and PIH (pregnancy induced hypertension) placentas by in situ hybridization. (A) and (B) are bright field micrograph with counterstained with hematoxylin. (a)-(d) are directly scanned from radiography autoradiogram using film scanner and inverted using photoshop (Ver. 5.0) program. *PIGF-1* (data not shown) and *PIGF-2* mRNAs are expressed in the vasculosyncytial membrane and villous stroma of the normal (A) and PIH placental villi (B). There is no significant difference in the expression of *PIGF-1* mRNA between the normal (a) and PIH placentas (b). The expression of *PIGF-2* mRNA is decreased in the PIH placenta (d) compared to the normal one (c). (original magnification:  $\times 2.1$ ). PIGF-1, placenta growth factor -1; PIGF-2, placenta growth factor -2. Scale bar in A, B= 50  $\mu$ m.

Localization of *PIGF-1* and *PIGF-2* mRNAs in the normal and PIH placentas by in situ hybridization

The *PIGF-1* (data not shown) and *PIGF-2* mRNAs were localized in the vasculosyncytial membrane and the villous stroma of placental villi in the normal (Fig. 3A) and PIH placenta (Fig. 3B). While no significant difference was detected in the expression of *PIGF-1* mRNA between normal (Fig. 3a) and PIH placentas (Fig. 3b), the expression of *PIGF-2* mRNA was decreased in the PIH placenta (Fig. 3d) compared to the normal one (Fig. 3c).

Expression of *PIGFR-1* and *PIGFR-2* mRNAs in the normal and PIH placentas

The mRNAs encoding *PIGFR-1* and *PIGFR-2* were detected by RT-PCR in both the normal and PIH placentas. While the expression of *PIGFR-1* was significantly higher in the PIH placentas compared to the normal one ( $p < 0.05$ ) (Fig. 4A, B), that of *PIGFR-2* showed no significant difference (Fig. 4C,

D). Relative densities were normalized by  $\beta$ -actin signals.

DISCUSSION

Although many theories have been proposed to explain the development of PIH, most of previous reports have not provided a clear picture. What is known, to date, is that PIH originates in the placenta and involves vasospasm of blood vessels in both mother and the placenta causing damage to the interior lining of blood vessels; its association with endothelial damage and dysfunction has been reported (2).

PIGFs, angiogenic growth factors, and their receptors are essential for placental vascular development (12). In this study, we examined the expression level of *PIGF-1* and *PIGF-2* mRNAs in both the normal and PIH placentas at term and found a decreased mRNA level of *PIGF-2* in PIH placentas by Northern blots and in situ hybridization. This result agrees well with previous reports; serum PIGF level was significantly lower in PIH as compared to normal pregnancies (18, 19);

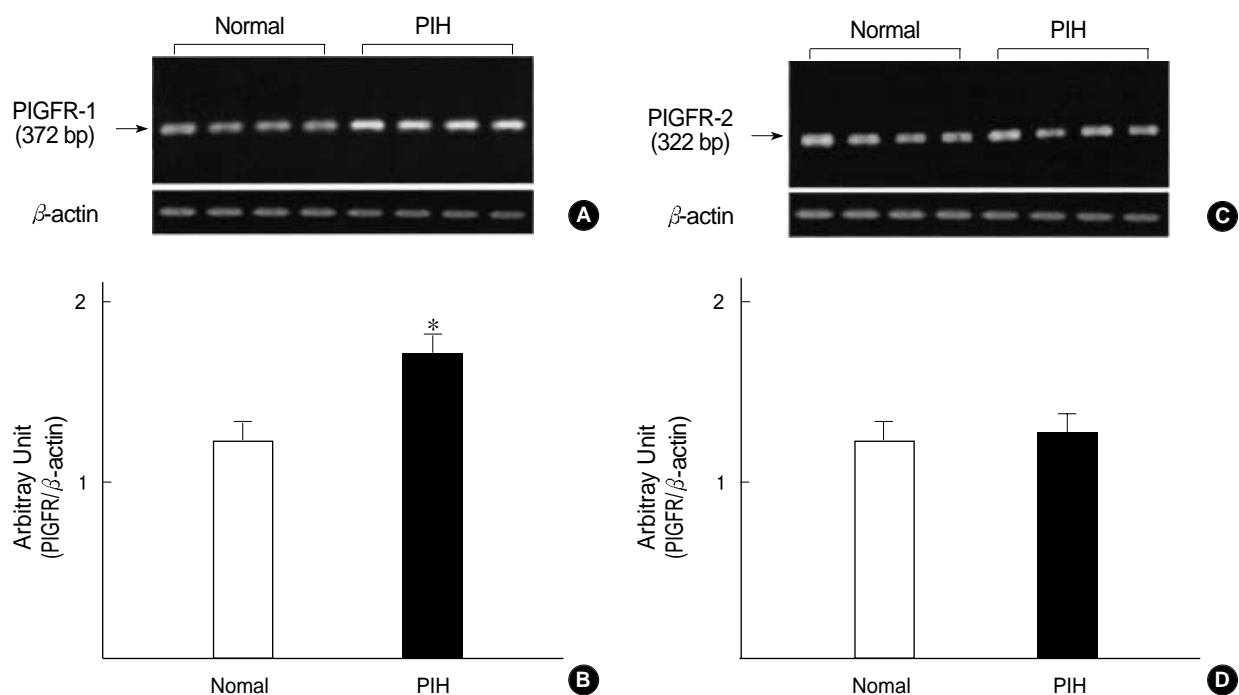


Fig. 4. Quantitative analyses of *PIGFR-1* and *PIGFR-2* in the normal and PIH placentas by RT-PCR. The product signals at 372 bp and 322 bp represent mRNAs of *PIGFR-1* (A) and *PIGFR-2* (C), respectively. The expression of *PIGFR-1* is significantly greater in the PIH (pregnancy induced hypertension) placenta compared to controls ( $p < 0.05$ ) (B), while that of *PIGFR-2* shows a mild increase (D). Relative densities were normalized to  $\beta$ -actin. PIGFR-1, placenta growth factor receptor-1; PIGFR-2, placenta growth factor receptor-2.

the transcription of *PIGF-2* mRNA in the placenta and decidua from PIHs decreased significantly (20).

Biological activities of PIGF-2 are mediated by the PIGFR-1 (21, 22). Three cell types in the placenta express the PIGFR-1; endothelial cells, Hoffbauer cells and trophoblasts, with extravillous trophoblast showing particularly high expression (23). In normal pregnancies, fetal trophoblasts invade the maternal decidua and remodel the spiral arteries, converting them to lower resistant vessels. In pregnancies complicated by pre-eclampsia, this trophoblastic cell invasion is inadequate, resulting in poor placental perfusion and fetal hypoxia (24, 25). The increased syncytial knots in the PIH placenta (Fig. 1) compared to normal one implicate an inadequate invasion of trophoblastic cells and this may be caused by a lower level of PIGF-2.

We revealed higher level of *PIGFR-1* in the PIH placenta compared to normal one using RT-PCR and this was also consistent with other reports (19, 26); PIGFR-1, but not the other receptors, showed increased expression in the placental syncytiotrophoblasts among 50% of patients with severe pre-eclampsia compared to normal ones (19). Helske *et al.* (19) suggested that the increased PIGFR-1 expression is probably not a specific change of characteristic for pre-eclampsia, but is possibly associated with abnormal placental function and hypoxia. It was also revealed that trophoblastic cells express both insoluble PIGFR-1 and soluble PIGFR-1, and the soluble form PIGFR-1 have an antagonistic regulation function

of its ligands (27). Therefore, the inadequate development of chorionic villi after mid gestation in PIH placenta may be possibly due to the lower level of *PIGF* mRNA and higher level of *PIGFR-1* mRNA.

In the human placenta, VEGF, PIGF and their receptors are differentially expressed throughout gestation: VEGF and PIGFR-2 are most intense during early gestation and decline as pregnancy advances (13-15) while PIGF and PIGFR-1 increase towards term (16, 17). Correlation of these growth-factor effects and their expression patterns throughout gestation with a development of a villous angioarchitecture (28, 29) suggest that VEGF and PIGFR-2 are involved in the first two trimesters of pregnancy in the establishment of the richly branched capillary beds of the mesenchymal and immature intermediate villi while PIGF and PIGFR-1 are more likely to be involved in the formation of the long, poorly branched, terminal capillary loops in the last trimester (30).

Although PIGF itself does not significantly stimulate angiogenesis *in vivo* and *in vitro*, this growth factor modulates the angiogenic activity of VEGF by forming PIGF/VEGF heterodimers (31). Therefore, the amount of VEGF is also important in complicated placenta etiology. However, recent reports provide conflicting conclusions regarding the expression of VEGF in pre-eclamptic pregnancies; at the RNA level, VEGF expression was found to decrease in single biopsy samples of pre-eclamptic placentas as compared to biopsies from normal placentas (32); VEGF protein levels in serum of pre-

eclamptic women were found to be elevated in some studies (33, 34) while others have reported decreased levels (35), relative to normal pregnant women; PIGF homodimers are themselves angiogenic (36) and may also function to enhance the activity of suboptimal concentrations of VEGF (21). Furthermore, naturally occurring VEGF/PIGF heterodimers maintain high-affinity receptor binding and mitogenic activity on HUVE cells (37). While VEGF binds with high affinity to both PIGFR-2 and PIGFR-1, PIGF exhibits high affinity only to PIGFR-1 (21). Therefore, it is suggested that the main cause of delicate placental complications is PIGF rather than VEGF considering its amount consistency in complicated pregnancy placenta and its specificity to PIGFR-1. Especially, the adequate *PIGF-2* mRNA expression level in placenta seems to be essential for normal placenta morphometric development in late pregnancy.

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