# Aberrations in Placental Cytokine mRNA Related to Intrauterine Growth Retardation

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# ABSTRACT

During normal pregnancy, a predominance of Th2 type cytokines prevails and is considered to protect the fetus. Animal experiments suggest that an increase of Th1 type cytokines may instead have deleterious effects. We have studied with the reverse transcription PCR technique mRNA for IL-1 $\alpha$ , IL-1 $\beta$ , IL-6, IL-8, IL-10, transforming growth factor- $\beta$ , tumor necrosis factor- $\alpha$ , and interferon- $\gamma$  in placentas from full-term appropriately grown newborns, newborns with intrauterine growth retardation (IUGR) and newborns who were only small for gestational age. The mRNA for IL-10 was significantly reduced in the IUGR placentas (p < 0.05), whereas the mRNA for IL-8 was significantly higher (p < 0.05) for the IUGR cases compared with the full-term neonates. It might be that reduced IL-10 in the placenta is involved in the pathogenesis of IUGR. (*Pediatr Res* 51: 201–206, 2002)

#### Abbreviations

AGA, appropriate for gestational age **DEPC**, diethyl pyrocarbonate dNTP, deoxynucleotide triphosphate EtBr, ethidium bromide **IFN.** interferon IUGR, intrauterine growth retardation KO, knock-out pd(N)<sub>6</sub>, random hexamer primer PE, preeclampsia **RT-PCR**, reverse transcription PCR SGA, small for gestational age **TBE**, tris-borate-EDTA TGF, transforming growth factor Th1, T helper type 1 Th2, T helper type 2 TNF, tumor necrosis factor

Lately it has been realized that cytokines provide important signals during pregnancy for the implantation of the egg, trophoblast growth, hormone production, and, finally, for the induction of delivery *via* prostaglandins (1, 2). A predominance of Th2-derived cytokines (*i.e.* IL-4, IL-10, and TGF- $\beta$ ) appears to characterize the normal pregnancy, whereas indications of a shift of balance in favor of Th1-derived cytokines (*i.e.* IL-2, IFN- $\gamma$ , and TNF- $\alpha$ ) has been reported in a few pathologic pregnancies (2, 3). Increased placental exposure to IFN- $\gamma$  and TNF- $\alpha$  from trophoblast-activated lymphocyte supernatants from women with unexplained recurrent abortions may impair fetal development in mice, either as IUGR or resorption of the fetus (4). In experimental animals, it was noted that TNF- $\alpha$ , IFN- $\gamma$ , and IL-2 can increase fetal resorption, whereas IL-3 and granulocyte-macrophage colony stimulating factor injected in abortion-prone mice seemed to have a trophic effect on the placenta, thus increasing fetal and placental weight (5). The same mouse strain prone to habitual abortion turned out to be deficient in local IL-10 production and the abortions were prevented by IL-10 (6). In IL-10deficient mice, generated by gene targeting, most animals are growth retarded and suffer from chronic enterocolitis (7), but there seems to be no reports concerning intrauterine growth in IL-10 KO animals. The IL-10 KO litters appear normal in number and the offspring are normal in size. It seems that the growth retardation occurs in conjunction with developing inflammation in the gut (D. Rennick, personal communication). On the other hand, it was shown that neutralization of IL-10 by administration of anti-IL-10 MAb to pregnant mice did not affect duration of gestation or the fetal outcome. However, neonates exposed to anti-IL-10 MAb in utero showed transient

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growth deficiency (8). Available information from humans is mainly derived from studies on cytokine levels in blood.

In the present study, we have analyzed mRNA for Th1 and Th2 cytokines in placentas comparing AGA, SGA, and IUGR neonates to test the hypothesis that IUGR in humans may be connected to a predominance of cytokines from the Th1 group.

## **METHODS**

**Collection and handling of placentas.** Placentas were obtained from 34 normal term deliveries resulting in AGA babies, 20 IUGR deliveries (11 without and 9 with PE), and 7 SGA deliveries (4 without and 3 with PE). All IUGR babies but one were delivered by cesarean section, and 4 out of 7 SGA babies were delivered vaginally. In the control group, there were 23 vaginal deliveries, 9 cesarean sections, and for 2 the mode of delivery was not clearly specified. There were no signs of infection or inflammation, either in any of the study groups varied between 26 and 41 wk, whereas the control group consisted of term babies only. No neonates with malformations or any other reason for low birth weight than PE were included.

SGA and IUGR infants weighed at least 22% less than normal at their respective gestational ages. Most SGA and IUGR infants were diagnosed during pregnancy after repeated ultrasound examinations. IUGR was defined as lack of increase in estimated fetal weight ( $\leq$ 11% compared with standard curves, examination interval  $\geq$ 12 d) and/or pathologic Doppler velocimetry in the umbilical artery. Small fetuses with normal increase of estimated weight compared with standard curves and normal Doppler velocimetry in the umbilical artery were classified as SGA (9, 10).

PE was defined as arterial blood pressure exceeding 140/90 in the presence of significant albuminuria (>0.3 g/24 h), with onset after 20 full weeks of pregnancy.

After delivery, the placenta was placed on ice and the time until the samples were frozen was a maximum of 15 min. Between 50 and 100 mg of tissue were dissected from trophoblast or "fetal placenta" and decidua or "maternal placenta," avoiding connective tissue and blood vessels. The tissue was rinsed in a physiologic salt solution and frozen at  $-70^{\circ}$ C. Samples were then transferred to liquid nitrogen, where they were stored until tested.

The RNA was isolated and analyzed with RT-PCR technique for the relative quantification of gene expression of IL-1 $\alpha$ , IL-1 $\beta$ , IL-6, IL-8, IL-10, TGF- $\beta$ 1, TNF- $\alpha$ , and IFN- $\gamma$ , using for comparison a simultaneously amplified sequence of the housekeeping gene glyceraldehyde 3-phosphate dehydrogenase (G3PDH).

We analyzed separately the samples from the trophoblast and from the decidua from the placentas in each group. We did not isolate certain cell populations, but used the whole tissue samples either from trophoblast or decidua for the isolation of total RNA.

**RNA preparation.** Total RNA enriched for mRNA was isolated from the tissue by a single-step method with a ready-to-use reagent (RNA STAT-60, Tel Test B, Friendswood TX, U.S.A.), which includes phenol and guanidinium thiocyanate

in a monophase solution. Isolation by this method according to the manufacturer includes the following steps: homogenization of the tissue, RNA extraction with chloroform, RNA precipitation with isopropanol, and RNA wash with ethanol. Then the isolate was treated with RQ1 RNase-free DNase (Promega, Madison, WI, U.S.A.) to remove possible remnants of genomic DNA from the sample and another extraction of RNA was carried out.

The yield and purity of isolated RNA was determined from spectrophotometric measurements of absorbances at 260 and 280, as well as 230 and 270 nm wavelengths. RNA was checked for quality on a 3% agarose gel with EtBr (Sigma Chemical, St. Louis, MO, U.S.A.) and in the presence of a RNA standard (Human HeLa Cell Total RNA, CLONTECH, Palo Alto, CA, U.S.A.). If the isolated RNA was found to have distinct 28S and 18S ribosomal bands, it was considered intact and of good quality.

**Reverse transcription.** For cDNA synthesis, 2  $\mu$ g of RNA and pd(N)<sub>6</sub> (Pharmacia-Upjohn, Uppsala, Sweden) were used. A mixture of dNTPs (Pharmacia-Upjohn), RNase inhibitor from placenta (Promega), Superscript (reverse transcriptase, Invitrogen, Carlsbad, CA, U.S.A.), first strand buffer (Invitrogen), and DEPC-treated water (DEPC-H<sub>2</sub>O) (Sigma Chemical) were added to a total volume of 30  $\mu$ L.

**PCR.** For each cytokine, commercially available specific primers and positive controls (CLONTECH) were used. The primer sequences are listed below:

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IL-1\alpha
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Upstream 5' CAAGGAGAGCATGGTGGTAGTAGCAAC-CAACG 3'

Downstream 5' TAGTGCCGTGAGTTTCCCAGAAGAA-GAGGAGG 3'

IL-1β

Upstream 5' ATG GCA GAA GTA CCT AAG CTC GC 3' Downstream 5' A CAC AAA TTG CAT GGT GAA GTC AGT T 3'

IL-6

Upstream 5' ATG AAC TCC TTC TCC ACA AGC GC 3' Downstream 5' G AAG AGC CCT CAG GCT GGA CTG 3' IL-8

Upstream 5' ATG ACT TCC AAG CTG GCC GTG GCT 3' Downstream 5' T CTC AGC CCT CTT CAA AAA CTT CTC 3'

IL-10

Upstream 5' AAGCTGAGAACCAAGACCCAGACAT-CAAGGCG 3'

Downstream 5' AGCTATCCCAGAGCCCCAGATC-CGATTTTGG 3'

TGF-β1

Upstream 5' GCC CTG GAC ACC AAC TAT TGC T 3' Downstream 5' AG GCT CCA AAT GTA GGG GCA GG 3' TNF- $\alpha$ 

Upstream 5' GAG TGA CAA GCC TGT AGC CCA TGT TGT AGC A 3'

Downstream 5' GCA ATG ATC CCA AAG TAG ACC TGC CCA GAC T $3^\prime$ 

IFN- $\gamma$ 

Upstream 5' GCATCGTTTTGGGTTCTCTTGGCTGT-TACTGC 3'

Downstream 5' CTCCTTTTTCGCTTCCCTGTTTTAGCT-GCTGG 3'

PCR amplification of 2  $\mu$ L cDNA was performed in the presence of AmplyTaq Gold DNA Polymerase, PCR buffer, 2.0 mM MgCl<sub>2</sub> (all from PerkinElmer Cetus, Norwalk CT, U.S.A.), dNTP mix (Pharmacia), and sterile water in a total volume of 50  $\mu$ L. The PCR was performed in a PerkinElmer Cetus GeneAmp PCR System 9600. The first step was incubation at 95°C for 12 min, then 35 cycles of denaturation at 94°C for 45 s, annealing at 65°C for 45 s, and extension at 72°C.

Negative controls consisted of samples in which the reverse transcriptase was omitted to detect possible contamination by genomic DNA and controls in which cDNA was replaced by sterile  $dH_2O$ .

Gel electrophoresis of the PCR products. The PCR products were stored at 4°C until tested by submarine electrophoretic separation in TBE buffer. Ten microliters of sample were mixed with 5  $\mu$ L of gel loading solution (Sigma Chemical) on a 3% agarose gel (SeaKem and NuSieve, FMC Bioproducts, Rockland, ME, U.S.A.) containing EtBr (Sigma Chemical) and in the presence of the molecular size marker  $\Phi$ X174/*Hae*III-digested DNA (Invitrogen). The gels were photographed in the dark by Polaroid MP 4+ Instant Camera System (Sigma Chemical) under illumination of UV light.

*Analysis of the detected signals.* Gel images were scanned (StudioScan II/si, Agfa) and densitometric analysis, *i.e.* quantification of bands in terms of position, height and area in two dimensions, referred to as "volume" analysis, was performed with IPLab Gel software for Macintosh (BioSystematica, Plymouth, U.K.). The results were expressed as a ratio calculated from the "volume" of the amplified cytokine gene product divided by the "volume" of the amplified housekeeping gene product in the same sample.

Statistical analyses. Data were analyzed by nonparametric Mann-Whitney U test and Dunn's test for nonparametric multiple comparisons for unequal sample sizes. Spearman's rank correlation was used for the association analyses. Medians were used for all analyses.

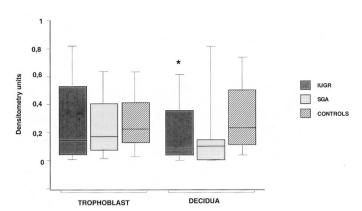
*Ethics.* This study was conducted with informed consent of the patients and approved by the ethics committee.

#### RESULTS

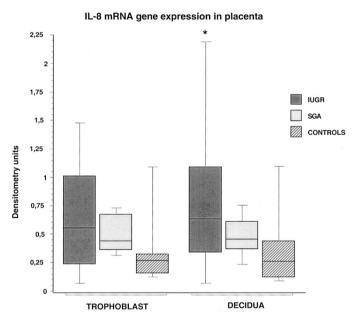
We found a significant reduction of IL-10 mRNA expression in the decidua of the IUGR placentas compared with the control material (p < 0.05, Fig. 1). In parallel, the IL-8 mRNA expression was significantly higher in the decidua of the IUGR placentas (p < 0.05) compared with the control group (Fig. 2).

The ratio between the IL-8 and IL-10 mRNA expression was significantly higher both in the decidua and the trophoblast of the IUGR compared with the control placentas (p < 0.01 and <0.05, respectively) (Table 1). There was a negative correlation between the expression of mRNA levels for IL-8 and IL-10 in the decidua of the IUGR placentas, although it was not statistically significant. Thus, this showed a trend that the

IL-10 mRNA gene expression in placenta



**Figure 1.** IL-10 mRNA expression in trophoblast and decidua samples from normal and pathologic placentas. The IL-10 mRNA levels in decidua of the IUGR group were significantly lower compared with the controls (\*p < 0.05). The IL-10 mRNA expression in the SGA placentas was not significantly lower compared with either controls or the IUGR group. Medians and 25–75% confidence intervals are indicated.

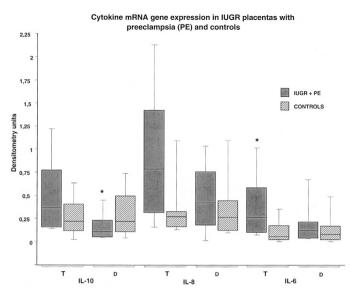


**Figure 2.** IL-8 mRNA expression in trophoblast and decidua samples from normal and pathologic placentas. In the decidua of the IUGR group, the IL-8 mRNA levels were significantly higher compared with the controls (\*p < 0.05). There was no significant difference in IL-8 mRNA levels between the SGA and either controls or IUGR groups. Medians and 25–75% confidence intervals are indicated.

individuals with lower IL-10 had higher IL-8 mRNA levels in the decidua and *vice versa*.

We divided the IUGR group into the two subgroups, individuals with PE during pregnancy and those without PE. Individuals from the IUGR group with PE had significantly lower IL-10 mRNA levels in the decidua (p = 0.05), and significantly higher IL-6 mRNA levels in the trophoblast compared with the controls (p < 0.05; Fig. 3 and Table 1).

Individuals with IUGR but no PE during pregnancy had significantly higher IL-8 mRNA levels in the decidua (p < 0.01; Table 1). The ratios of the mRNA levels for the pro-/



**Figure 3.** Cytokine gene expression in IUGR placentas from PE cases and control placentas. The mRNA for IL-10 was significantly lower in the decidua of the IUGR group with PE compared with the controls (\*p < 0.05). The mRNA for IL-6 was significantly higher in the trophoblasts of the IUGR placentas with PE compared with the controls (\*p < 0.05). D, decidua, T, trophoblast. Medians and 25–75% confidence intervals are indicated.

anti-inflammatory cytokines were significantly higher in the decidua of these individuals compared with the controls (p < 0.05; Table 1).

When the IUGR group with PE was compared with the IUGR group without PE during pregnancy, the IL-6 mRNA levels were significantly higher in the trophoblast of the group with PE than in the group without PE (p < 0.02). The mRNA levels of the two anti-inflammatory cytokines (IL-10 and TGF- $\beta$ ) were also higher in the trophoblast of the IUGR individuals with PE compared with those without PE (p < 0.01). In contrast, the mRNA levels of the proinflammatory cytokines (IL-1 $\alpha$ , IL-1 $\beta$ , IL-8, IFN- $\gamma$ , and TNF- $\alpha$ ) in the decidua were significantly lower in the group with IUGR and PE compared with the group without PE (p < 0.03). Likewise, the ratios between the mRNA levels of the pro-/anti-inflammatory cytokines were significantly lower in the decidua from the group with PE (p < 0.02; Table 1).

Although there was a lower IL-10 mRNA expression and a higher IL-8 mRNA expression both in the decidua and the trophoblast of the SGA placentas, they were too few in number (seven samples only) and the variations between individuals were too large to reach any significant differences. Therefore we analyzed the placentas from the two groups of small neonates (comprised of the IUGR and SGA neonates) together in comparison to the placentas of normal size newborns (Table 1). The IL-10 mRNA levels in the decidua were significantly lower (p < 0.05) and the IL-8 mRNA levels were significantly higher both in the trophoblast and the decidua compared with the controls (p < 0.05 and p < 0.01, respectively). The ratios of the IL-8/IL-10 mRNA levels in the trophoblast and the decidua samples from the placentas of the small children were significantly higher compared with the controls (p < 0.01). There was a negative correlation between IL-10 and IL-8 mRNA expression in the decidua of these children, although not statistically significant.

The placentas of the small newborns (IUGR and SGA) whose mothers had PE had significantly higher levels of IL-6 mRNA in the trophoblast (p < 0.05), as did the placentas of the mothers from the group with PE and IUGR alone (Table 1). The IL-8 mRNA levels were also higher in the trophoblast samples of this group compared with the controls (p < 0.01; Table 1).

Expression of mRNA for IL-1 $\alpha$ , IL-1 $\beta$ , and IFN- $\gamma$  was rather low in all the placentas, whereas the expression of TGF- $\beta$  was extremely high in comparison with the expression of other cytokines, although the levels did not differ significantly between the groups.

### DISCUSSION

This study supports the concept that there may be a decrease of IL-10 mRNA in the placenta of IUGR neonates. Previous reports have suggested that the protein levels may also be decreased. Thus, a decreased production of IL-10 from peripheral blood monocytes was found in a few pregnancies with growth-retarded fetuses of not clearly defined nature (3). In animal studies, a strain of abortion-prone mice was found deficient in placental production of IL-10 (6). Furthermore, in normal murine pregnancies, administration of anti-IL-10 MAb was associated with transient neonatal growth retardation, although it did not alter the duration of gestation or the fetal outcome (8).

A low IL-10 level in the placenta might result in an increase in Th1 cells with an elevated production of the cytokines IFN- $\gamma$ , TNF- $\alpha$ , and IL-2. This may have damaging effects on the fetus causing either impaired growth, or abortion, or resorption of the fetus (2). On the other hand, the cytokines of the colony stimulating factor family may reduce the damage and even increase fetal and placental weight (5), possibly by effects on the placenta similar to those of IL-10.

In IL-10 gene KO mice, exaggerated inflammation occurs (11). Immunoreactive cells (activated macrophages, Th1 cells, and natural killer cells) from peripheral tissues of these animals when stimulated in culture produce 3 to 20-fold more IL-6, TNF- $\alpha$ , and IFN- $\gamma$  than the cells from control mice. This suggests that the deficiency of IL-10 also in the placenta might result in elevated levels of, e.g. IFN- $\gamma$  and TNF- $\alpha$ , increasing the risk of fetal damage. Such an imbalance between Th1 and Th2 cytokines in the placenta may also be harmful in human pregnancies. Indeed, in a recent study of women with unexplained recurrent spontaneous abortion, significantly lower levels of the Th2 cytokines IL-10 and IL-6 and significantly higher levels of the Th1 cytokine IFN- $\gamma$  were detected compared with normal pregnancy (12). Moreover, another study showed a difference in cytokine production by stimulated peripheral blood mononuclear cells between mothers with normal and abnormal pregnancies (3). A significantly increased production of IL-2 and IFN- $\gamma$  and a decrease of IL-10 were seen in pathologic pregnancies with abortions, or SGA babies, compared with normal pregnancies.

				L	Decidua				Irophoblast	
Groups	No.	IL-10	IL-8	IL-8/IL-10	Proinflammatory	Pro-/Anti- inflammatory	IL-8	IL-8/IL-10	IL-6	Anti-inflammatory
IUGR vs.	20	0.05	0.05	0.01	NS	NS	NS	0.05	NS	NS
controls	34	(I < C)	(I > C)	(I > C)				(I > C)		
Small (IUGR +	27	0.05	0.01	0.01	NS	NS	0.05	0.01	NS	
SGA) <i>vs.</i> controls	34	(S < C)	(S > C)	(S > C)			(S > C)	(S > C)		NS
Small + PE $vs$ .	12	NS	NS	NS	NS	NS	0.01	NS	0.05	NS
controls	34						(S > C)		(S > C)	
IUGR + PE vs.	6	0.05	NS	NS	NS	NS	NS	NS	0.05	NS
controls	34	(I < C)							(I > C)	
IUGR - PE vs.	11	NS	0.01	NS	NS	0.05	NS	NS	NS	NS
controls	34		(I > C)			(I > C)				
IUGR + PE vs.	11	NS	NS	NS	0.03	0.02	NS	NS	0.02	0.01
IUGR – PE	6				(I + PE < I - PE)	(I + PE < I - PE)			(I + PE > I - PE)	(I + PE > I - PE)

Is the low expression of IL-10 in placentas from IUGR pregnancies a primary change or secondary to an increase of pro-inflammatory cytokines from the Th1 cells caused by hypoxia? TNF- $\alpha$  and IL-1 are markedly stimulated by hypoxia in villous explants of the human placenta (13). But in our study, no changes consistent with such results were found. Therefore, the reduced IL-10 expression may be a primary event.

If a deficiency in placental IL-10 really is related to the pathogenesis of IUGR in humans, one might consider using IL-10 therapeutically. Actually, IL-10 can be given to humans without significant untoward effects, and it may reduce IFN- $\gamma$ and TNF- $\alpha$  levels (14, 15). Furthermore, it has been noted in an animal model that IL-10 biosynthesis may be increased with large doses of methylprednisolone (16).

We may speculate that the significant increase in the ratios of mRNA levels of the pro-inflammatory cytokines to the anti-inflammatory cytokines, which we see in the decidua of the IUGR group compared with the controls, might be caused by the low levels of IL-10 that may not be sufficient to dampen the inflammatogenic effects.

It has been suggested that normal pregnancy itself stimulates an inflammatory response and that PE is what happens when, toward the end of pregnancy, this inflammatory response becomes exaggerated (17). Furthermore, a decrease in IL-10 and an increase in TNF- $\alpha$  both in serum and in trophoblast were reported in PE compared with normal pregnancy (18). This phenomenon may be associated with increased maternal antifetal immunity and thus inadequate placental development in PE. The deficiency of placental IL-10 in PE is in good agreement with our findings.

We also observed higher mRNA levels of IL-6 only in the trophoblast of individuals with PE during pregnancy. This corroborates the report that significantly elevated levels of IL-6 and IL-8 were found in mid-trimester amniotic fluid of the future preeclamptic group compared with the normal pregnancies (19).

The higher production of placental mRNA for IL-8 in the IUGR cases may be a result of the cytokine imbalance. IL-8 is produced in increased amounts at delivery, presumably both to protect the damaged intrauterine surfaces after delivery via its strong chemotactic effect on neutrophils and to repair via its angiogenic effect (20, 21). IL-8 also potentiates production of neutrophil-derived collagenase required in helping ripening and opening of the cervix at delivery (22-24).

Further studies will be required to define whether a deficiency in placental IL-10 during pregnancy is involved in the pathogenesis of IUGR. If so, new possibilities for prophylaxis and treatment of IUGR might become available. This is of interest inasmuch as IUGR is such an important risk factor for newborns, especially in poor countries (25, 26) and not only in malnourished, but also in well-nourished populations.

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