

Dexamethasone stimulates the expression of leptin and 11 β -HSD2 in primary human placental trophoblastic cells

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1. Introduction

Glucocorticoid excess during fetal life has been shown to play an important role in early-life programming after intrauterine growth restriction (IUGR), contributing to the development of adult metabolic, cardiovascular, and neuroendocrine disease [1,2].

The enzyme 11 β -hydroxysteroid dehydrogenase type 2 (11 β -HSD2) is expressed in the human placenta and has been suggested to regulate the materno-fetal transfer rate of active cortisol via conversion to inactive cortisone [3,4]. In cultured human trophoblasts, dexamethasone incubation induced 11 β -HSD2

activity and mRNA expression, suggesting an autoregulation of 11 β -HSD2 by glucocorticoids [5]. This finding is supported by the observation that in healthy pregnancies, increased circulating glucocorticoid levels in pregnant women [6] were paralleled by an increased placental 11-HSD2 expression with advancing gestation [7]. Moreover, in conditions of high maternal glucocorticoid levels like prenatal administration of glucocorticoids to accelerate fetal lung maturation [8], no fetal growth-reducing effects were seen in humans, in contrast to experiments performed in rodents and sheep [9].

In pregnancies complicated by idiopathic IUGR, placental 11 β -HSD2 activity and mRNA were decreased, thus exposing the fetus to excess maternal cortisol [2,10,11].

In animal studies, the adverse effects of glucocorticoids on fetal growth and endocrine function have been extensively studied [12–14]. Beside direct influences on the feto-maternal unit, the effects

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of glucocorticoids on birth weight seemed to be mediated by the placenta, as infusion of betamethasone directly to sheep fetuses did not result in growth restriction [12]. In vitro experiments showed that dexamethasone (10^{-6} M) administration in placental villous tissue altered nutrient transfer capacity by stimulating system A activity, thus promoting uptake of amino acids from maternal to fetal circulation [15].

In view of the fact that glucocorticoid action seems to be modulated by placental enzyme activity, we aimed to investigate which placental systems might be affected by the hypercortisolism in IUGR. In this respect, altered gene and protein expression of various placental endocrine regulators associated with IUGR has already been described, e.g. leptin, and 11 β -HSD2 [16–19]. In the present study, we hypothesized that dexamethasone incubation of trophoblastic cells from human healthy placentas might induce altered gene expression of 11 β -HSD2 and leptin. This could represent an important safeguard mechanism to protect the fetus from detrimental exposure to elevated levels of maternal glucocorticoids, if these processes also occurred in the human placenta in vivo.

2. Materials and methods

2.1. Placental collection and tissue culture

Two term placentas from women with singleton uncomplicated pregnancies were collected immediately after placental delivery in collaboration with the Department of Obstetrics and Gynecology at the University of Erlangen-Nuremberg. In both cases, caesarean section delivery was performed, and birth weight was >10th percentile according to Voigt et al. [20]. The study was reviewed and approved by the ethics committee of the University of Erlangen-Nuremberg. Written informed consent was obtained from all subjects.

Primary human cytotrophoblasts were isolated from the placentas using the established trypsin-DNase-dispase/percoll method as initially described by Kliman et al. [21], with additional previously published modifications [22]. Cells were seeded into 6-well plates (Falcon) at a density of 8×10^6 cells/cm² and maintained in Earl's Medium 199 (M199, PAA Laboratories, Linz, Austria) supplemented with 10% fetal calf serum (PAA Laboratories), 20 mM Hepes (Sigma–Aldrich Chemie GmbH, Munich, Germany), 0.5 mM L-glutamine (Gibco Invitrogen GmbH, Karlsruhe, Germany), penicillin (10 U/ml), streptomycin (10 mg/ml), and fungizone (0.25 mg/ml) (Sigma–Aldrich Chemie GmbH, Gibco Invitrogen GmbH).

Cultures were grown at 37 °C under normoxia with 95% air, 5% CO₂ in an incubator (Forma Scientific) as described in detail [23]. Medium was changed 2, 5 and 24 h after initial plating to remove non-adherent cells and syncytial debris. After incubation for 24 h, trophoblastic cells were stimulated with 10 μ M dexamethasone for 6, 12, 24, 48 and 72 h. Non-dexamethasone-stimulated cells served as vehicle control. Cultured trophoblasts as well as culture supernatants were collected at these time points, snap frozen and stored at –80 °C until further processing. All experiments were assayed in triplicate and were repeated using cells from a different placenta.

2.2. RNA isolation and reverse transcription-polymerase chain reaction (RT-PCR)

Total RNA was isolated from primary human trophoblasts using TRIzol[®] reagent (Invitrogen GmbH, Karlsruhe, Germany) as recommended by the manufacturer. RNA was quantified by absorbance at 260 nm, and the quality of RNA was confirmed using a 1% agarose gel. After DNase treatment, 1.0 μ g RNA was

transcribed into cDNA using M-MLV-RT (Promega, Madison, WI, USA) and Oligo dT-primer (MWB-Biotech AG, Ebersberg, Germany). DNase treatment and cDNA synthesis were carried out according to the kit instructions. The cDNA expression level was quantified by real-time PCR.

2.3. TaqMan real-time PCR

To quantify the gene expression (mRNA) of leptin, 11 β -HSD2 and IGFBP-1 as markers of potential decidual overgrowth, TaqMan real-time PCR (Perkin Elmer, Foster City, CA) was used. The mRNA expression was normalized to two different house-keeping genes, hypoxanthine guanine phosphoribosyl transferase (HPRT) and β 2-microglobulin (β 2-MG), that can be detected pseudogene free. Commercial reagents (TaqMan PCR reagent kit, Perkin-Elmer) and conditions were applied according to the manufacturer's protocol. 2.5 μ l of complementary DNA (reverse transcription mixture) and oligonucleotides at a final concentration of 300 nmol/L (HPRT, 11 β HSD2, IGFBP1 forward) or 600 nmol/L (β 2-MG, IGFBP1 reverse, leptin) of primers and 200 nmol/L of TaqMan hybridization probe were analyzed in a 25 μ l-volume. All of the primers and probes were purchased from Eurogentec (Belgium) and Sigma (Germany). PCR was conducted using the following thermocycler parameters: 50 °C for 2 min, 95 °C for 10 min, followed by 40 cycles of 95 °C and 60 °C for 1 min. Serial dilutions of one of the samples served as reference providing relative quantification of the unknown samples.

The following primers and TaqMan probes were used. HPRT: forward 5'-CCGGCTCCGTTATGGC-3', reverse 5'-GGTCATAACCTGGTTCATCATCA-3', TaqMan probe 5'(FAM)-CGCAGCCCTGGCGTCGTGATTA-(TAMRA)3'; β 2-MG: forward 5'-TGACTTTGTCACAGCC-CAAGATA-3', reverse 5'-CCAAATGCGGCATCTTC-3', TaqMan probe 5'(FAM)-TGATGCTGCTTACATGCTCGATCCCA-(TAMRA)3'; IGFBP-1 forward 5'-CTCTCCATGTCACCAACATCAA-3'; reverse 5'-GTGCC TTGGCTAAACTCTCTACGA-3'; TaqMan probe 5'(FAM)-AATGGAAG-GAGCCCTGCCGAATAGAAGTTC-(TAMRA)3'; leptin forward 5'-ACAA TTGTACCAGGATCAATGAC-3'; reverse 5'-TCCAAACCGTGACTTT CTGT-3'; TaqMan probe 5'(FAM)-TTTACACACGCAGTCAGTCTCC TCCA-(TAMRA)3'; 11 β -HSD2 forward 5'-CATCACC GGCTGTGACT CTG-3'; reverse 5'-CGGCAGCCGATGTTAG-3'; TaqMan probe 5'(FAM)-AAAGGAGACAATTTGGCTCTGC-(TAMRA)3'.

2.4. Leptin, β -hCG and LDH concentration determination

Leptin in the culture supernatants was measured using the Human Leptin ELISA Kit (RayBio[®], RayBiotech Inc., Norcross, USA) following the manufacturer's instructions. The concentration of β -human chorionic gonadotropine (β -hCG) in the culture medium was determined by the DRG- β -hCG-ELISA Kit (DRG[®] Diagnostics GmbH, Marburg, Germany) according to the kit protocol. Lactate dehydrogenase (LDH) concentrations were obtained spectrophotometrically by the In Vitro Toxicology Assay Kit Lactate Dehydrogenase based (TOX-7, Sigma–Aldrich Chemie GmbH, Germany). All measurements were assayed in triplicate. Analysis of the results was performed using Ascent Software for Multiscan Vers. 2.6 (©1996–2002 Thermo LabSystems Oy).

2.5. Statistical analysis

Results were expressed as mean \pm standard error of the mean (SEM). Differences were assessed using a Kruskal–Wallis one-way ANOVA, where appropriate, with post hoc Bonferroni correction. Statistical analyses were performed using GraphPad Prism[®] Version 4.0c for Windows, GraphPad Software, San Diego, CA (www.graphpad.com). The limit of significance was set at a *p*-value of <0.05.

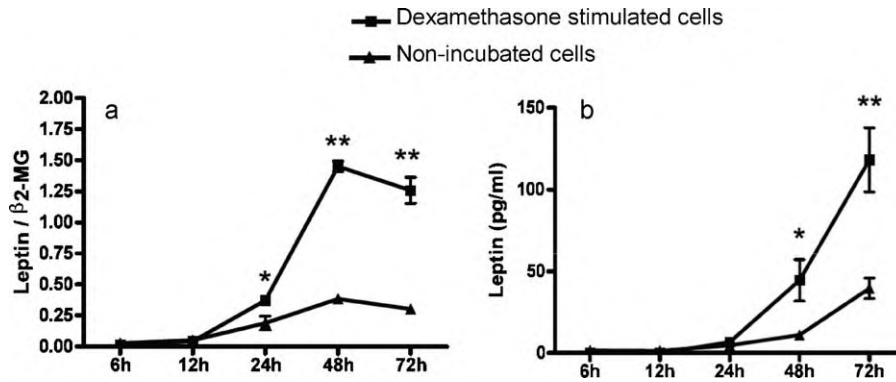


Fig. 1. (a) Leptin gene expression and (b) leptin release in the culture supernatant in human trophoblastic cells in vitro with and without stimulation with 10 μ M dexamethasone. There is a significant increase in relative leptin gene expression after 24 h in the dexamethasone stimulated cells. Leptin release in the culture medium was significantly elevated after 48 h of dexamethasone incubation. Gene expression is related to the housekeeping gene β 2-microglobulin (β 2-MG). Leptin concentration in the culture supernatant is expressed in absolute values (pg/ml). The lines show the mean \pm SEM. * $p < 0.05$; ** $p < 0.001$.

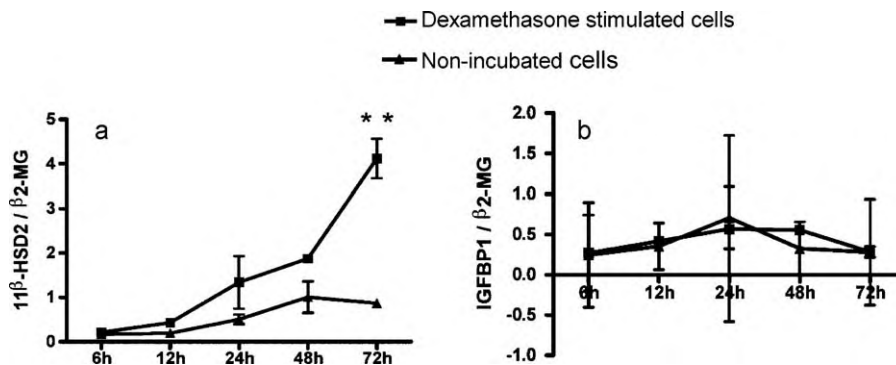


Fig. 2. Gene expression of (a) 11 β -HSD2 and (b) IGFBP-1 in human trophoblastic cells in vitro with and without stimulation with 10 μ M dexamethasone. There is a significant increase in relative 11 β -HSD2 gene expression after 72 h in the dexamethasone stimulated cells. Relative IGFBP-1 mRNA expression did not change during the incubation period in both cell lines. Gene expression is related to the housekeeping gene β 2-microglobulin (β 2-MG). The lines show the mean \pm SEM. * $p < 0.05$; ** $p < 0.001$.

3. Results

Stimulation of primary placental trophoblasts with 10 μ M dexamethasone resulted in a significant increase in leptin gene expression after 24 h compared to the vehicle control (Fig. 1a). When measuring leptin release in the supernatant, we found significantly higher leptin concentrations in the dexamethasone-stimulated trophoblastic cells after 48 h (Fig. 1b).

Dexamethasone incubation significantly increased gene expression of 11 β -HSD2, not before 72 h (Fig. 2a). The expression rate of IGFBP1 mRNA as marker for decidual overgrowth in our cell cultures was basal throughout the incubation period in both cell lines (Fig. 2b).

To determine whether the effect of dexamethasone was mediated through alterations in placental cell differentiation or

viability, medium was collected at 6, 12, 24, 48 and 72 h during the culture period and assayed for β -hCG secretion as well as LDH release. The secretion of β -hCG in the culture medium as a biochemical fusion marker, which has been shown to correlate with the cell-cell fusion process of cytotrophoblasts into multinuclear syncytia [21,24], increased significantly after 72 h of dexamethasone incubation (Fig. 3). LDH release from placental trophoblasts, a marker of necrosis, remained stable during the incubation period in both dexamethasone-stimulated cells and vehicle controls (Fig. 4).

Results for protein concentration in the culture supernatant are expressed in absolute values (ng/ml or pg/ml). All data for placental gene expression are presented as relative gene expression normalized for β 2-microglobulin (Figs. 1 and 2). Similar results were obtained when expression data were normalized to HPRT (data not shown).

4. Comment

Fetal exposure to high glucocorticoid levels during pregnancy has been shown to result in low birth weight, which predisposes to the development of cardiovascular and metabolic disorders such as hypertension and type 2 diabetes, increased HPA axis reactivity, and increased anxiety-related behaviour in adult life [10,25–29]. In this respect, the central role of the placenta in the early programming process by moderating fetal exposure to maternal factors is well recognized [30].

In the present study, we investigated whether glucocorticoid exposure induced alterations in gene expression of specific endocrine placental regulators in vitro. According to previous

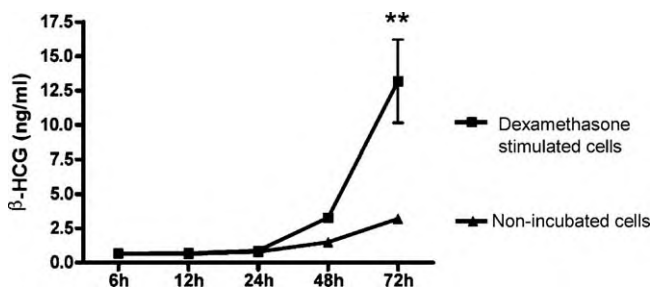


Fig. 3. β -hCG concentration in the culture medium in human trophoblastic cells in vitro with and without stimulation with 10 μ M dexamethasone. There is a significant increase in β -hCG levels after 72 h in the dexamethasone stimulated cells. β -hCG concentration in the culture supernatant is expressed in absolute values (ng/ml). The lines show the mean \pm SEM. * $p < 0.05$; ** $p < 0.001$.

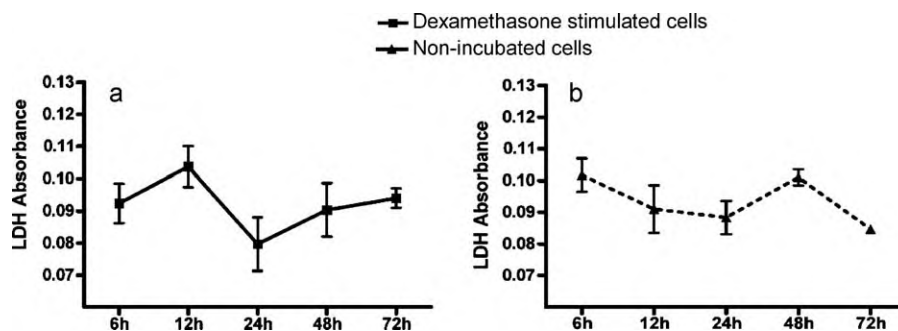


Fig. 4. LDH absorbance in the culture medium in human trophoblastic cells in vitro (a) with and (b) without stimulation with 10 μ M dexamethasone. There is no significant increase in LDH absorbance during the incubation period in both cell lines. The lines show the mean \pm SEM.

reports, we chose a concentration of 10 μ M dexamethasone for trophoblast stimulation [15,31].

Glucocorticoid regulation of maternal and fetal hormonal parameters and metabolism has been well described in ruminants and rodents [32,33]. In this context, maternal dexamethasone administration in pregnant rats induced, for example, an almost three-fold increase in the concentration of the ob-gene product leptin in maternal circulation. In parallel, leptin concentrations in the fetal circulation were shown to be decreased by around 80%, associated with changes in placental leptin receptor expression and reductions in placental content and transfer of leptin [34,35].

During pregnancy, leptin is involved in the regulation of fetal growth, placental angiogenesis and immunomodulation, and mobilization of maternal fat [36]. In vitro experiments showed an upregulation of nitric oxide production and lipolysis by leptin in the human and rodent placenta at term [37,38] with possible consequences for placental blood flow and transfer of free fatty acids to the fetus. Furthermore, in human placental villous fragments at term, leptin stimulated the activity of the amino acid transporter system A, responsible for the placental transfer of neutral amino acids to the fetus [39]. In previous investigations, where IUGR was confirmed by prenatal Doppler ultrasound examination, placental leptin gene expression was significantly increased [18,19].

Thus, leptin could be one of the fetoplacental factors altering maternal metabolism to benefit the fetus by mobilizing nutrients. According to placental perfusion studies, 98.4% of placental leptin, mainly produced in the syncytiotrophoblast [40], is released into the maternal circulation [41] and less than 2% into the fetal circulation; placental production, however, contributes significantly to circulating leptin in the human fetus, as demonstrated by the fall in serum leptin concentration in neonatal circulation that occurs after birth [42].

In our study, we observed a stimulatory effect of dexamethasone on leptin mRNA and leptin release in cultured trophoblastic cells. Induction of leptin gene expression in adipose tissue has already been reported after treatment with various glucocorticoids including dexamethasone [43]. In this respect, several glucocorticoid response element consensus binding sites have been found in the proximal promoter of the human leptin gene, indicating a direct transcriptional action of this hormone on the ob-gene [44]. Leptin synthesis could also be increased in vitro by stimuli like dexamethasone, hypoxia and insulin [23,45,46]. Thus, a time- and dose-dependent increase in leptin release and mRNA levels was observed in cultured human trophoblastic cells [45].

We therefore speculate that glucocorticoid exposure in utero might, at least in part, induce leptin synthesis. Whether the increase in leptin expression was a direct effect of dexamethasone on villous trophoblastic cells or secondary to an increased syncytialisation rate requires further investigation. Moreover, it

remains speculative whether this process might play a role in the excessive intrauterine glucocorticoid exposure leading to IUGR [10,25].

Our results showed that dexamethasone incubation stimulated 11 β -HSD2 gene expression in healthy human term trophoblasts. This is in line with data from the literature, where stimulation of primary cultured cytotrophoblasts with dexamethasone elevated 11 β -HSD2 gene expression [5,15,31]. Of note is that human placental trophoblastic cells, grown in primary culture, have been shown to maintain the same pattern of expression of 11 β -HSD as in vivo [47].

During normal pregnancy 11 β -HSD2 was reported to build a placental barrier regulating the amount of maternal cortisol passing to the fetus [3], thereby facilitating fetal growth [48] and fetal organ maturation [25,49]. In support of this, significant correlations between birth weight and placental 11 β -HSD2 activity have been found [16,50].

In IUGR, a reduction in 11 β -HSD2 function and a subsequent failure to protect the fetus from rising maternal cortisol concentrations have been implicated to play a key role in determining fetal growth and prenatal programming of adult disease in later life [51,52]. In this respect, placental 11 β -HSD2 activity correlated with cord-blood levels of osteocalcin (a glucocorticoid-sensitive osteoblast product that does not cross the placenta) and other markers of fetal exposure to glucocorticoids [53]. Moreover, variation in placental 11 β -HSD2 levels as well as exposure to glucocorticoids correlated with altered expression levels of various glucose and amino acid transporter molecules and growth factors in the placenta [15,54,55]. The effects of glucocorticoids on human placental function, however, in particular the transplacental transfer of nutrients from maternal to fetal circulation, are currently unknown.

Our results suggest that in healthy placentas, locally increased glucocorticoid concentrations may induce 11 β -HSD2 gene expression in an autocrine or paracrine fashion to shield the fetus from excessive maternal cortisol. But as the synthetic glucocorticoid dexamethasone is only a poor substrate for 11 β -HSD2 and readily crosses the placenta [56], the potential effect of dexamethasone on local placental metabolism of corticosteroids remains speculative. In addition, the rate of syncytialisation secondary to dexamethasone may have a major impact on 11 β -HSD2 expression. In a previous study, the incubation of cultured human placental trophoblastic cells with dexamethasone increased 11 β -HSD2 activity in a time- and concentration-dependent manner and stimulated gene expression during differentiation by enhancing 11 β -HSD2 mRNA stability and the rate of HSD11B2 gene transcription. Moreover, incubation with a glucocorticoid receptor antagonist reversed this effect. Thus, glucocorticoids seemed to autoregulate 11 β -HSD2 in the human placenta mediated through the glucocorticoid receptor [5].

The expression rate of IGFBP1 mRNA did not change during the incubation period in our study, neither in the dexamethasone nor in the vehicle control group. IGFBP-1 is mainly synthesized in maternal decidual cells [57], which are the major contaminating cells regarding the method of isolation we used for our experiments. It comprised a maximum rate of 5% contamination with other placental cell lines. As expected, we observed only a basal expression rate and no upregulation of IGFBP-1 gene expression in human trophoblastic cells. There was no sign of decidual overgrowth in our cell cultures.

In this study, the concentration of β -hCG in the culture supernatant increased significantly after 72 h of dexamethasone incubation. A glucocorticoid-dependent regulation of placental hCG has been previously described [15,58]. β -hCG secretion has been shown to correlate with the cell fusion process into multinuclear syncytia [21,24]. In this respect, dexamethasone incubation seemed to induce cytotrophoblast differentiation and maturation in our study, thereby potentially enhancing leptin and 11 β -HSD2 gene expression. Besides, glucocorticoid administration did not adversely affect trophoblast viability, as dexamethasone treatment did not significantly increase LDH release. Also in the literature, no effect of glucocorticoids on LDH has been described [15].

In conclusion, our data show that dexamethasone incubation stimulated leptin and 11 β -HSD2 gene expression in primary villous trophoblastic cells of healthy human term placentas, while enhancing cytotrophoblast differentiation. This could determine local metabolism of corticosteroids and, thereby, the passage of cortisol via the placental barrier to the fetus, which would protect the fetus from detrimental elevated maternal glucocorticoid exposure. We confirmed the results of our cell culture experiments in two placental cell lines, but there may be variations between trophoblasts from different placentas.

Further experiments are required to investigate the underlying mechanisms in detail and to evaluate whether these are regulatory processes which also occur in the human placenta in vivo. Moreover, it would be interesting to analyze the effect of glucocorticoid incubation on cytokine production. It remains speculative, however, whether these mechanisms might play a role in the pathophysiology of growth restriction and cardiovascular disease after IUGR.

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