

Influence of cortisol on adipose tissue development in the fetal sheep during late gestation

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Abstract

The present study examined the extent to which the late gestation rise in fetal plasma cortisol influenced adipose tissue development in the fetus. The effect of cortisol on the abundance of adipose tissue mitochondrial proteins on both the inner (i.e. uncoupling protein (UCP)1) and outer (i.e. voltage-dependent anion channel (VDAC)) mitochondrial membrane, together with the long and short forms of the prolactin receptor (PRLR) protein and leptin mRNA was determined. Perirenal adipose tissue was sampled from ovine fetuses to which (i) cortisol (2–3 mg/day for 5 days) or saline was infused up to 127–130 days of gestation, and (ii) adrenalectomised and intact controls at between 142 and 145 days of gestation (term=148 days). UCP1 protein abundance was significantly lower in adrenalectomised fetuses compared with age-matched controls, and UCP1 was increased by cortisol infusion and with gestational age. Adrenalectomy reduced the concentration of the long form of PRLR, although this effect was

only significant for the highest molecular weight isoform. In contrast, neither the short form of PRLR, VDAC protein abundance or leptin mRNA expression was significantly affected by gestational age or cortisol status. Fetal plasma triiodothyronine concentrations were increased by cortisol and with gestational age, an effect abolished by adrenalectomy. When all treatment groups were combined, both plasma cortisol and triiodothyronine concentrations were positively correlated with UCP1 protein abundance. In conclusion, an intact adrenal is necessary for the late gestation rise in UCP1 protein abundance but cortisol does not appear to have a major stimulatory role in promoting leptin expression in fetal adipose tissue. It remains to be established whether effects on UCP1 protein are directly regulated by cortisol alone or mediated by other anabolic fetal hormones such as triiodothyronine.

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Introduction

At birth, the neonate is exposed to low ambient temperatures for the first time and, depending on its maturity, may need to rapidly produce heat in order to prevent hypothermia. In newborn sheep, which are precocial offspring and therefore comparable to some extent to human infants, the rapid recruitment of non-shivering thermogenesis in brown adipose tissue is critical in the prevention of hypothermia (Clarke *et al.* 1997c). This effect is mediated by the large increase in amount and thermogenic potential of the brown adipose tissue-specific uncoupling protein (UCP)1 (Clarke *et al.* 1997a). Outer mitochondrial membrane proteins, such as the voltage-dependent anion channel (VDAC), may also influence brown adipose tissue function by regulating the supply of mitochondrial ATP and ADP (Gottlieb 2000).

In sheep, the abundance of UCP1 and VDAC in brown adipocytes is maximal at birth and then declines over the first month of postnatal life (Clarke *et al.* 1997b, Mostyn *et al.* 2001), concomitant with a switch from non-shivering to shivering thermogenesis as the primary mechanism of heat production (Symonds *et al.* 1989). During the perinatal period, thermogenic potential and UCP1 expression are upregulated by a number of hormones including noradrenaline, triiodothyronine (T₃), prolactin and leptin (Symonds *et al.* 2000, Evens *et al.* 2001, Budge *et al.* 2002). Consequently, the abundance of the receptors for these hormones, and the synthesis of leptin itself, in brown adipose tissue may also affect the function and abundance of UCP1 in the neonate.

In many tissues, the maturational changes essential for neonatal survival begin before birth and are dependent on the normal increase in plasma glucocorticoid concentration

in the fetus towards term (Liggins 1994). Cortisol has been shown to affect the expression of a wide range of proteins during the prepartum period including enzymes, hormones, binding proteins and receptors (Fowden *et al.* 1998). In sheep, the prepartum cortisol surge upregulates expression of prolactin receptors (PRLRs) in the fetal liver which coincides with the increase in hepatic 5'-monodeiodinase activity (Clarke *et al.* 1997a, Phillips *et al.* 1997). As a consequence of increased deiodination of thyroxine (T_4) to T_3 , the circulating T_3 concentration in the fetus also increases towards term (Fraser & Liggins 1989). In fetal sheep, there is a gradual increase in brown adipose tissue mitochondrial protein content and GDP binding during late gestation (Clarke *et al.* 1997a), which closely parallels the fetal cortisol surge towards term (Fowden *et al.* 1998). In addition, maternal administration of the synthetic glucocorticoid, dexamethasone, for 2 days in the week before full term enhances UCP1 abundance in perirenal adipose tissue from prematurely delivered sheep (Clarke *et al.* 1998). This occurs in conjunction with an increase in the short, but not the long form of the PRLR (Bispham *et al.* 1999). However, the role of cortisol in brown adipose tissue maturation and the developmental expression of UCP1 remains unknown. Hence, this study examined the hypothesis that fetal cortisol has a primary role in promoting fetal adipose tissue maturation in preparation for life after birth. This was examined by determining the effect of (i) infusing cortisol into the fetus between 122 and 125 days of gestation in order to increase fetal plasma cortisol to values observed near to term and (ii) removing the adrenal, thereby reducing fetal plasma cortisol to basal values. Effects on fetal adipose tissue maturation were then compared with age-matched controls with respect to the abundance of mitochondrial proteins (UCP1 and VDAC), PRLR and leptin in perirenal adipose tissue (which constitutes up to 80% of fetal adipose tissue).

Materials and Methods

Animals

A total of 27 Welsh Mountain sheep fetuses of known gestational age were used in this study. Ten were twin lambs and the remainder were singletons. In the case of twins only one twin was used in the study. The ewes were housed in individual pens from and maintained on 200 g/day concentrates, with free access to hay, water and a salt-lick block from 100 days of gestation. All animals were fed daily at ~0900 h. Food but not water was withheld for 18–24 h before surgery. The study aimed to determine the effect of (i) a precocious rise in fetal cortisol and (ii) a prolonged reduction in fetal cortisol. The first objective was achieved by comparing the effect of a 5-day period of cortisol infusion sufficient to double its plasma concentration up to 127–130 days of gestation. Effects on

adipose tissue maturation were then compared with age-matched fetuses into which saline had been infused. The second objective was achieved by surgically removing the fetal adrenal between 115 and 119 days of gestation, thereby preventing any prepartum rise in fetal cortisol and then comparing adipose tissue composition with non-operated age-matched fetuses near to term, i.e. 142–145 days of gestation, with term being 148 days of gestation. We have previously shown no difference in adipose tissue composition in neonatal lambs between those which have undergone a sham operation and those not subjected to this procedure (Schermer *et al.* 1996). All procedures were carried out under the UK Animal (Scientific Procedures) Act 1986.

Surgical procedures

Between 115 and 118 days (term 145 ± 2 days), one of the following procedures was carried out under halothane anaesthesia (1.5% in O_2/N_2O) using the surgical methods described previously (Fowden *et al.* 1996): (i) intravascular catheterisation of the fetus ($n=15$) or (ii) bilateral adrenalectomy of the fetus ($n=6$). Catheters were inserted into the aorta of the mother and the dorsal aorta and caudal vena cava of the fetus via the femoral vessels. At the end of surgery the fetuses were given 100 mg ampicillin (Penbritin; Beecham Animal Health, Brentford, UK) either i.v. or intra-amniotically. The ewes received antibiotic i.m. (1 g procaine penicillin (Depocillin; Mycofarm, Cambridge, UK)) on the day of surgery and for 3 days thereafter. Apart from the pre-surgery period of food withdrawal there were no apparent differences between operated and non-operated animals with respect to feeding patterns.

Experimental procedure

Blood samples of 2 ml were taken daily throughout the experimental period from all the catheterised fetuses to monitor fetal well-being and to determine plasma cortisol and thyroid hormone concentrations. At least 6 days after catheterisation, 13 fetuses were infused with either cortisol ($n=8$, 2–3 mg/day in 3 ml 0.9% (w/v) saline (EF-Cortelan; Glaxo Ltd, Greenford, Middlesex, UK)) or saline ($n=7$, 3 ml/day, 0.9% w/v) for 5 days before tissue collection at 127–130 days. The fetuses to be given cortisol were selected randomly. The dose of cortisol was chosen to produce concentrations of plasma cortisol similar to those observed close to term, i.e. 40–50 ng/ml (Fowden *et al.* 1996). There was no difference in the distribution of male and female fetuses between treatment groups or any effect of gender on plasma cortisol.

Sample collection

All operated fetuses, regardless of treatment, and six additional non-operated intact fetuses were delivered by

Caesarean section under sodium pentobarbitone anaesthesia (20 mg/kg i.v.) between 127 and 130 days of gestation for cortisol- and saline-infused fetuses and between 142 and 145 days for adrenalectomised (AX) and control fetuses. Fetal blood samples were taken at the time of delivery either through the indwelling arterial catheter or by venipuncture from the umbilical artery in the cord. After administration of a lethal dose of anaesthetic (200 mg/kg sodium pentobarbitone), samples of perirenal adipose tissue were collected and frozen rapidly in liquid nitrogen before storage at -80°C . All blood samples were centrifuged immediately at 4°C and the plasma stored at -20°C before plasma analyses. No adrenal remnants were found in any of the AX fetuses at autopsy.

Laboratory analyses

Protein immunodetection Crude plasma membranes or mitochondria were prepared from 1 g frozen adipose tissue as described previously (Budge *et al.* 2000) and the protein content of each preparation was determined (Lowry *et al.* 1951). The thermogenic potential of mitochondria was determined using [^3H]GDP, which measures the specific binding of GDP to UCP1 in mitochondrial preparations at a physiological concentration of $2\ \mu\text{M}$ (Symonds *et al.* 1992). PRLR abundance in plasma membranes was detected using $6\ \mu\text{g}$ protein, following protein separation by SDS-PAGE, immunoblotting and enhanced chemiluminescence (Amersham International) (Budge *et al.* 2000) utilising polyclonal antibodies R122 and R133 described by Nevalainen *et al.* (1996), which are specific for the long and short forms of PRLR respectively. These antibodies do detect a range of different molecular mass isoforms of each form of the receptor that are tissue-specific and have been interpreted as representing separate extracellular domains of the receptor (Nevalainen *et al.* 1996, Budge *et al.* 2000). UCP1 content in perirenal adipose tissue was measured as described by Schermer *et al.* (1996). VDAC abundance was determined using an antibody raised in rabbits to ovine VDAC (Budge *et al.* 2002) purified from the kidney of a newborn lamb based on the method of Schermer *et al.* (1996) as described for UCP1 and was used at a dilution of 1 in 2000. Densitometric analysis was performed on each gel and all values were expressed in densitometric units. Specificity of detection was confirmed using non-immune rabbit serum. A range of molecular mass markers was included on all gels. Densitometric analysis was performed on each membrane following image detection using a Fujifilm LAS-1000 cooled charge-coupled device camera (Fuji Photo Film Co. Ltd, Tokyo, Japan). All gels were run in duplicate and a reference sample (i.e. 4-hour-old lamb born vaginally at term) was included on each.

mRNA detection Total RNA was isolated from perirenal adipose tissue using Tri-Reagent (Sigma). In order

to maximise sensitivity, a two-tube approach to reverse transcription (RT) was adopted. The conditions used to generate first-strand cDNA RT were: 70°C (5 min), 4°C (5 min), 25°C (5 min), 25°C (10 min), 42°C (1 h), 72°C (10 min), 4°C (5 min). The RT reaction (final volume $20\ \mu\text{l}$) contained: $5 \times$ cDNA (first-strand), buffer (250 mM Tris-HCl, 40 mM MgCl_2 , 150 mM KCl, 5 mM dithioerythritol pH 8.5), 2 mM dNTPs, $1 \times$ hexanucleotide mix, 10 units RNase inhibitor, 10 units M-MLV reverse transcriptase and $1\ \mu\text{g}$ total RNA. All these commercially available products were purchased from Roche.

The expression of mRNA for leptin was determined as described by Bispham *et al.* (2002). The analysis utilised the following cDNA primers to the ovine leptin gene: 5'-CAC CAA AAC CCT CAT CAA GAC G-3' (27-58) and 5'-ACA TTT CTG GAA GGC AGA CTG G-3' (197-228, Genbank U84247), which generated a 192 bp intron spanning product. QuantumRNA alternate 18S internal standards (Ambion, Abingdon, Oxon, UK) were included in the multiplex PCRs. Briefly, the incubation conditions were: 94°C (2 min) 1 cycle; 94°C (30 s), 60°C (30 s), 72°C (1 min) 30 cycles; 72°C (7 min) 1 cycle. The PCR mixture (final volume $20\ \mu\text{l}$) contained: $10 \times$ PCR buffer (100 mM Tris-HCl, 15 mM MgCl_2 , 500 mM KCl pH 8.3), 500 μM dNTPs, 1 mM each leptin primer, 3.75 U Taq polymerase. Agarose gel electrophoresis (2.0%) and ethidium bromide staining confirmed the presence of both leptin and 18S products of the expected sizes. UCP1 mRNA abundance was determined as described by Mostyn *et al.* (2002) with densitometric analysis performed as described for UCP1 protein above. Consistency of lane loading for each sample was verified by hybridisation with the 18S rRNA oligo-probe and all results were expressed as a ratio of UCP1 to r18S abundance. All analyses and gels were conducted in duplicate.

RIAs Cortisol concentrations were measured by RIA and validated for use with ovine plasma (Robinson *et al.* 1983). The minimum detectable quantity of cortisol was $1.5\ \text{ng/ml}$ and interassay coefficient of variation was 11%. Total plasma T_3 and T_4 concentrations were also measured by RIA, using a commercial kit (ICN Biomedicals, Thame, Oxon, UK) validated for ovine plasma (Fowden *et al.* 1983). The lower units of detection were $0.1\ \text{ng/ml}$ for T_3 and $7\ \text{ng/ml}$ for T_4 . The interassay coefficient of variation was 10% for both assays.

Statistical analyses

Statistical analysis with respect to significant differences ($P < 0.05$) between mean values obtained between treatment groups was carried out using Mann-Whitney U tests. Regression analysis between UCP1, cortisol and T_3 was carried out using SPSS.

Table 1 Effect of cortisol infusion, adrenalectomy (AX) and gestational age on plasma concentrations of cortisol, T₃, T₄ and thermogenic potential (GDP binding), VDAC protein and leptin mRNA abundance in perirenal adipose tissue of the ovine fetus. Values are means ± s.e. with numbers of animals in each group given in parentheses

	Treatment	Plasma cortisol (ng/ml)	Plasma T ₃ (ng/ml)	Plasma T ₄ (ng/ml)	GDP binding (μmol GDP/ mg MP)	VDAC (% ref.)	Leptin mRNA (% of 18S)
Gestational age (days)							
	127–130						
	Saline	14.0 ± 2.3 (7)	0.28 ± 0.04 (7)	76.4 ± 4.3 (7)	16.3 ± 1.3 (5)	90.3 ± 6.8 (5)	2.3 ± 1.4 (5)
	Cortisol	45.7 ± 5.3* (8)	0.57 ± 0.16* (5)	64.1 ± 4.3 (6)	35.1 ± 11.9 (5)	104.7 ± 3.7 (5)	4.4 ± 2.2 (5)
142–145							
	Control	49.5 ± 12.3 [†] (6)	0.49 ± 0.07 [†] (6)	71.3 ± 7.3 (6)	23.3 ± 2.8 (5)	92.9 ± 7.9 (5)	1.4 ± 0.2 (5)
	AX	7.2 ± 0.06* (6)	0.29 ± 0.04* (6)	82.1 ± 5.6 (6)	18.2 ± 2.3 (6)	72.5 ± 9.2 (6)	1.9 ± 0.3 (6)

ref.=4-hour-old newborn lamb; MP=mitochondrial protein.

*Significantly different from the corresponding control group, $P < 0.05$.

[†]Significantly different from the value in the control group at 127–130 days, $P < 0.05$.

Results

Ontogenic development of fetal perirenal adipose tissue

The fetal concentrations of plasma cortisol and T₃ were significantly higher close to term (142–145 days) than earlier in gestation (Table 1). This occurred in the absence of any significant changes in T₄. At 142–145 days, abundance of UCP1 protein, but not mRNA, in perirenal adipose tissue was significantly higher than at 127–130 days (Fig. 1). In contrast to UCP1 protein, there was no change in GDP binding to mitochondria from perirenal adipose tissue with increasing age (Table 1). There was a non-significant trend for mean GDP binding to be increased by cortisol infusion. Gestational age had no effect on VDAC protein, leptin mRNA (Table 1) or protein abundance for either form of PRLR (Fig. 2). Fetal weight was not significantly different between the two treatment groups at each gestational age (data not shown).

Effects of manipulating the fetal cortisol concentrations on perirenal adipose tissue

Fetal adrenalectomy Removal of the fetal adrenal gland prevented the normal rise in fetal plasma concentrations of cortisol and T₃ towards term: mean values of plasma cortisol and T₃ in the AX fetuses at 142–145 days were significantly less than those seen in intact controls at the same gestational age (Table 1). Fetal adrenalectomy also prevented the prepartum rise in UCP1 protein abundance in the perirenal adipose tissue (Fig. 1a). At 142–145 days, mean UCP1 protein abundance in the AX fetuses was significantly less than in the control fetuses at the same gestational age and was similar to that seen earlier in gestation. Abundance of UCP1 mRNA also tended to be less in AX than in control fetuses close to term but this

difference was not statistically significant (Fig. 1b). Fetal adrenalectomy had no effect on mRNA abundance for leptin or VDAC protein (Table 1). However, it did reduce expression of the long form of PRLR compared with the control fetuses close to term, although this effect was only significant for the 52 kDa isoform (Fig. 2a). No effect was observed for the short form of PRLR (Fig. 2b).

Fetal cortisol infusion Infusion of cortisol into immature fetuses raised fetal plasma concentrations of cortisol and T₃ (Table 1). Mean values of plasma cortisol and T₃ were significantly higher in cortisol-infused fetuses than in the saline-infused controls at 127–130 days (Table 1). UCP1 protein, but not mRNA abundance, was similarly increased by cortisol infusion (Fig. 1). Fetal cortisol infusion had no effect on the protein abundance for VDAC or prolactin receptor (Fig. 2) or leptin gene in the perirenal adipose tissue.

Relationship between UCP1 protein abundance and plasma concentrations of cortisol and T₃

Since UCP1 protein abundance was elevated when cortisol and T₃ levels were raised and were low when cortisol and T₃ were low, the relationship between UCP1 protein abundance in perirenal adipose tissue and the plasma concentrations of these hormones was examined in more detail. When individual results from all fetuses were combined regardless of gestational age or treatment, the plasma concentration of T₃ was significantly correlated to plasma cortisol (Fig. 3). In addition, there was a significant positive correlation between UCP1 protein abundance in perirenal adipose tissue and both the plasma concentrations of cortisol and T₃ (Fig. 4). Multiple regression analysis of the three variables gave standardised beta coefficients of

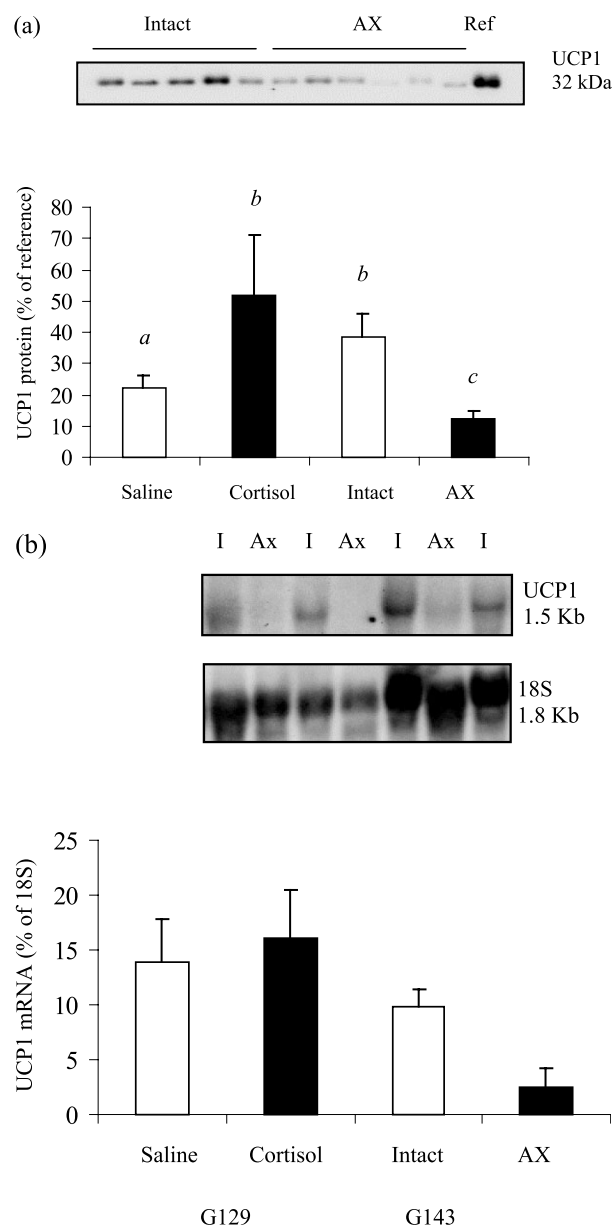


Figure 1 Effect of cortisol infusion, adrenalectomy (AX) and gestational age on uncoupling protein (UCP1) (a) protein and (b) mRNA in fetal perirenal adipose tissue. Examples of a charge-coupled device camera image showing the effect of adrenalectomy on (a) UCP1 protein and (b) UCP1 mRNA in individual fetuses sampled between 142 and 145 days of gestation are also shown. Values are means with their s.e. values and $n=4-6$ per group. G=mean gestational age. Ref=reference sample, i.e. adipose tissue sampled from a 4-hour-old lamb born vaginally. Significant differences with gestational age or treatment group: *a* vs *b*, $P<0.05$; *a* vs *c* $P<0.01$.

0.328 and 0.448 for T_3 and cortisol respectively, indicating similar effects of each hormone with respect to their potential regulation of UCP1 protein.

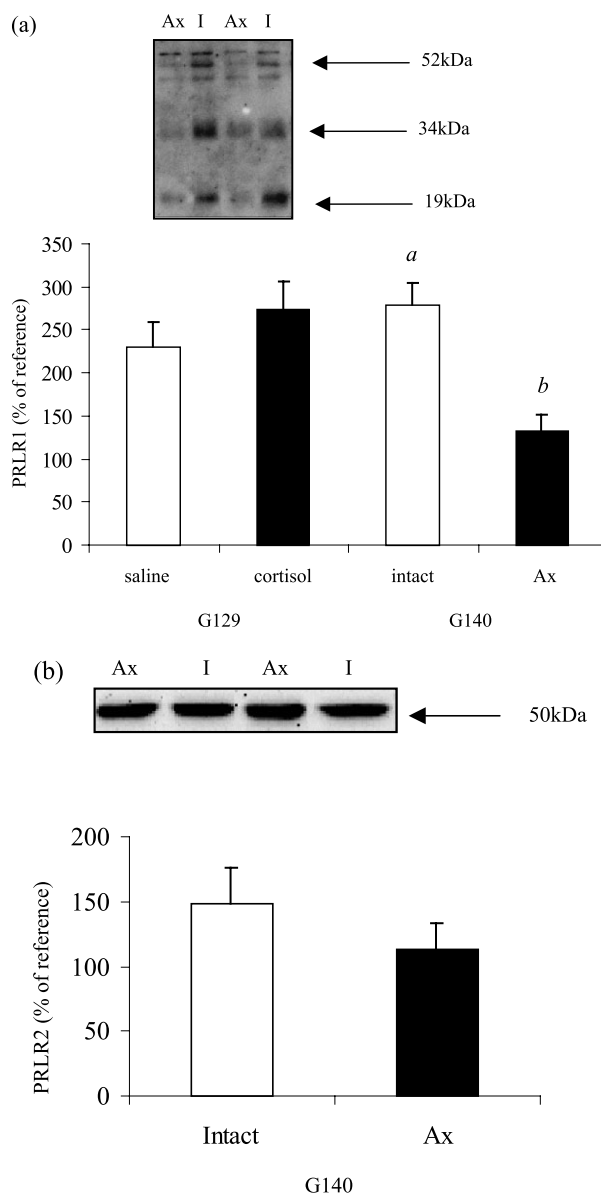


Figure 2 Effect of cortisol infusion, adrenalectomy (Ax) and gestational age on the abundance of the (a) long (PRLR1, 52 kDa) and (b) short forms (PRLR2) of prolactin receptor in fetal perirenal adipose tissue. Examples of a charge-coupled device camera image showing the effect of adrenalectomy on each specific isoform for (a) the long form of PRLR protein (see Methods) and (b) the short form or PRLR in individual fetuses sampled between 142 and 145 days of gestation are shown. G=mean gestational age. I=intact. Values are means with their s.e. values and $n=4-6$ per group. Significant differences between treatment groups: *a* vs *b* $P<0.01$.

Discussion

The major finding of the present study is that, in fetal sheep, cortisol regulates the level of the brown adipose

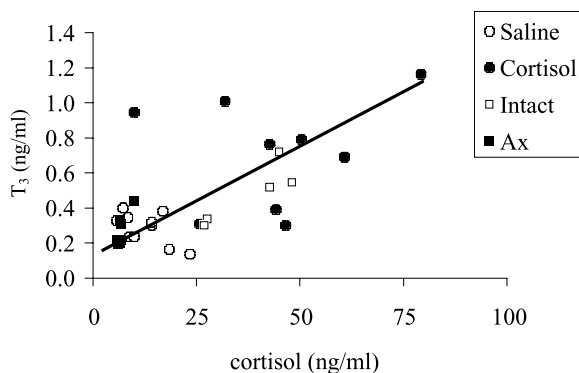


Figure 3 Positive relationship between plasma cortisol and T_3 concentrations in all fetuses irrespective of treatment or gestational age. $R^2=0.595$, $P=0.001$, where $y=0.01x+0.18$.

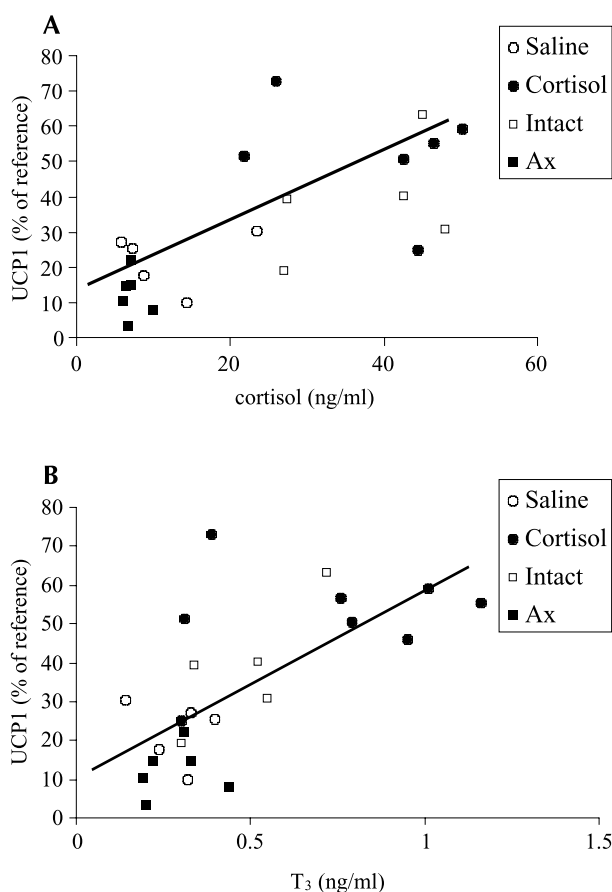


Figure 4 Positive relationship between uncoupling protein (UCP)1 abundance and plasma concentrations of (A) cortisol ($R^2=0.48$, $P<0.001$, where $y=0.66x+14.78$) and (B) T_3 ($R^2=0.44$, $P<0.001$, where $y=46.8x+11.15$) concentrations in all fetuses irrespective of treatment or gestational age. Reference sample is adipose tissue sampled from a 4-hour-old lamb born vaginally.

tissue-specific UCP1 protein but not UCP1 mRNA or GDP binding to UCP1 in adipose tissue mitochondria. Abundance of UCP1 protein increased with gestational

age in parallel with the rise in plasma cortisol towards term. When this prepartum increase in cortisol was prevented by fetal adrenalectomy, the rise in UCP1 protein abundance towards term was abolished. Conversely, raising cortisol levels at a time when concentrations are normally low, increased UCP1 protein abundance to values similar to those seen in older animals close to term. Cortisol, therefore, appears to enhance UCP1 protein synthesis in ovine adipose tissue during late gestation. This effect appeared to be specific and not a global response of mitochondrial proteins to cortisol as VDAC, located on the outer mitochondrial membrane, showed no change in abundance during manipulation of the fetal cortisol level. The positive correlation we also observed between the plasma cortisol concentration *in utero* and UCP1 protein abundance strongly indicates that cortisol is a physiological regulator of UCP1 protein abundance and suggests that it may enhance the capacity for thermogenesis in preparation for cold exposure at birth.

The extent to which the effects of cortisol on UCP1 protein abundance are direct or mediated via other hormone systems remains unclear. The changes in UCP1 protein abundance did not appear to depend on leptin expression in perirenal adipose tissue. Leptin mRNA abundance was unaffected by gestational age or by manipulation of fetal plasma cortisol in the current study. Glucocorticoids have been shown to stimulate leptin gene expression in adult adipocytes *in vitro* (Russell *et al.* 1998) and to increase circulating leptin concentrations *in vivo* in both adult mice and human subjects (Masuzaki *et al.* 1997, Arvaniti *et al.* 1998). In fetal sheep, plasma leptin concentrations increased transiently 24–48 h after the onset of fetal cortisol infusion but were normal on the fifth day of infusion at the time when perirenal adipose tissue was collected in the present study (Forhead *et al.* 2002). In previous studies of fetal sheep, both increases and decreases in leptin expression have been observed in perirenal adipose tissue over the same period of late gestation examined in the present study (Yuen *et al.* 1999, Devasker *et al.* 2002). This may be explained, in part, by breed differences in leptin expression (Dandrea 2001) but may also reflect differences in nutritional intake by the ewes between early to mid gestation, when a reduction in maternal food intake results in increased leptin mRNA in fetal adipose tissue as measured near to term (Symonds *et al.* 2002).

There were also no changes in PRLR protein abundance in perirenal adipose tissue in response to fetal cortisol administration in the present study. This contrasts with earlier findings which showed cortisol-dependent upregulation of PRLR mRNA expression in fetal ovine liver (Phillips *et al.* 1997). However, fetal adrenalectomy did lower abundance of the long form of PRLR in perirenal adipose tissue during late gestation. While these observations suggest PRLR expression may be regulated by an adrenal medullary secretion, they provide little

evidence for a cortisol-dependent, PRLR-mediated change in UCP1 protein abundance.

In several fetal tissues, T_3 is known to mediate the maturational effects of cortisol (Fowden *et al.* 1998). Cortisol induces hepatic activity of the 5'-monodeiodinase responsible for deiodinating T_4 to T_3 and, hence, leads to a concomitant rise in fetal plasma T_3 (Liggins 1994, Clarke *et al.* 1997a). A positive correlation between fetal plasma T_3 concentration and fetal cortisol levels is therefore in accord with previous studies (Fraser & Liggins 1989, Wallace *et al.* 1995). Fetal plasma T_3 was also positively correlated to UCP1 protein concentration that was equally as important as plasma cortisol in determining UCP1 abundance in the present study. In addition, exogenous administration of T_3 has been shown to upregulate mitochondrial proteins including UCP1 in fetal and newborn animals (Schermer *et al.* 1996, Symonds *et al.* 2000). However, even close to term, plasma T_3 levels were lower in the fetus than seen immediately after birth (Clarke *et al.* 1997c). The main effects of T_3 on UCP1 protein abundance may, therefore, occur after delivery. At the molecular level, the effects of T_3 and/or cortisol appear to be on translation rather than on transcription of the UCP1 gene as the ontogenic and cortisol-induced changes in UCP1 protein abundance occurred in the absence of any differences in UCP1 mRNA.

The actions of cortisol on UCP1 abundance in fetal adipose tissue are in keeping with its other known maturational effects *in utero* (Fowden *et al.* 1998). At birth, the neonate must be able to maintain its core temperature by rapidly producing heat if it is to survive cold exposure. By stimulating UCP1 protein abundance the prepartum increase in cortisol will enhance the ability of the fetus to produce heat by non-shivering thermogenesis (Clarke *et al.* 1997c). However, even close to term, levels of UCP1 were only 40–60% of those found postnatally. Factors other than cortisol, such as noradrenaline, are involved in upregulating UCP1 availability during the perinatal period (Symonds *et al.* 2000). Indeed, the thermogenic potential of fetal adipose tissue, measured as GDP binding, was not affected by manipulation of the fetal cortisol level in the present study. This may reflect the lack of fetal lipolysis *in utero* as GDP binding to UCP1 has been shown to depend on a rise in non-esterified fatty acid release by the adipocytes (Cannon & Nedergaard 1985). In fetal sheep, free fatty acid levels remain low during both endogenous and exogenous changes in fetal plasma cortisol (A L Fowden, unpublished observations) but rise rapidly immediately after birth when sympathetic activation of the adipocytes is maximal (Ball *et al.* 1992) and the inhibitory actions of the placental prostaglandins on fetal lipolysis are lost (Gunn & Gluckman 1995). Hence, there is a cortisol-dependent increase in UCP1 before birth but no actual increase in heat production until the GDP-binding sites on UCP1 are unmasking by the events of birth itself.

In conclusion, an intact adrenal is necessary for the appearance of fetal brown adipose tissue as characterised by the late gestation rise in UCP1 protein abundance. In contrast, cortisol does not appear to have a major stimulatory role in promoting white adipose tissue characteristics as cortisol status had no effect on leptin expression in fetal adipose tissue. It remains to be established whether the observed effects on UCP1 protein are directly regulated by cortisol alone or mediated by other anabolic fetal hormones such as T_3 .

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