Differential effects of the early and late intrauterine environment on corticotrophic cell development

Timothy G. Butler,1 Jeff Schwartz,2 and I. Caroline McMillen1

1Department of Physiology, University of Adelaide, Adelaide, South Australia, Australia
2Department of Physiology and Pharmacology, Department of Obstetrics and Gynecology, and the Perinatal Research Laboratory, Bowman Gray School of Medicine, Wake Forest University, Winston-Salem, North Carolina, USA

The developing embryo and fetus respond to a range of intrauterine stressors, but the effect of chronic intrauterine stress on the programmed development of pituitary corticotrophs has not been investigated. We have used a pregnant sheep model in which the embryonic environment at conception has been surgically perturbed by uterine carunclectomy. This procedure results in the development of fetuses that either are placentally restricted and chronically hypoxemic or that demonstrate compensatory placental growth and maintain normoxemia throughout late gestation. We found that uterine carunclectomy resulted in the emergence of a population of non–corticotrophin-releasing hormone (non-CRH) target cells that secreted high amounts of adrenocorticotrophic hormone (ACTH) in the fetal pituitary. This change in corticotroph development was independent of late-gestation hypoxemia. However, chronic hypoxemia during late gestation (in either carunclectomized or non-carunclectomized uterine environments) resulted in a reduction in the proportion of ACTH stored in CRH-target. Thus, the early and late intrauterine environments differentially program the development of specific corticotrophic cell types in the fetal pituitary. These patterns of altered corticotroph development are important given the central roles of the hypothalamo-pituitary-adrenal axis in the fetal adaptive response to intrauterine stress and in the early programming of adult disease.


Introduction

A worldwide series of epidemiological studies has demonstrated that there are significant associations between low birth weight and a range of poor adult health outcomes including high blood pressure, coronary heart disease, obesity, insulin resistance, and hypercortisolism (1–3). These associations have lead to the articulation of the “fetal origins of adult disease hypothesis,” which states that fetal adaptations to a period of intrauterine deprivation result in a permanent reprogramming of key organ systems and pathophysiological outcomes in later life (2). A range of studies have implicated intrauterine glucocorticoid exposure as one of the key mediators of the effects of intrauterine deprivation and have concluded that the timing, duration, and magnitude of fetal glucocorticoid exposure may each be important in the programming of poor health outcomes (3–7). It is therefore important to understand the impact of a suboptimal intrauterine environment on the functional capacity of the hypothalamo-pituitary-adrenal (HPA) axis at different stages of development. It is well established that the fetal HPA axis is activated in response to a range of acute intrauterine stressors including hypoxemia, hemorrhage, and insulin-induced hypoglycemia (8–10). The effects of chronic intrauterine stress on the fetal HPA axis are, however, less well understood. A model of chronic intrauterine stress has been developed in which the majority of the placental attachment sites, the uterine caruncles, are surgically excised from the uterus of the nonpregnant ewe (11, 12). During a subsequent pregnancy, a restricted number of placentomes form, and this results in a chronic placental restriction of fetal substrate supply, hypoxemia, and fetal growth restriction throughout late gestation (11, 12). Circulating cortisol concentrations are higher in growth-restricted fetuses of carunclectomized ewes, but, surprisingly, there is no associated increase in the fetal plasma concentrations of either immunoreactive adrenocorticotrophic hormone (ACTH) or ACTH1–39, and the expression of proopiomelanocortin mRNA is decreased in the anterior pituitary of the growth-restricted fetus (13). Different subpopulations of corticotrophs, which are responsive either to corticotrophin-releasing hormone (CRH) or to arginine vasopressin (AVP) and which are differentially sensitive to negative feedback by cortisol, have been described in the adult and fetal sheep pituitary (14–17). One
possibility is that uterine carunclectomy programs a change in the developmental characteristics of the pool of corticotroph cells within the pituitary to maintain ACTH secretion in the face of elevated cortisol concentrations during late gestation.

One important feature of uterine carunclectomy is that it does not inevitably lead to chronic fetal hypoxemia in late gestation, as there is a degree of compensatory growth of the remaining placenomes which may result in the maintenance of a relatively normoxic, well-grown fetus (11, 12). Therefore, among the effects of uterine carunclectomy on the fetal HPA axis, it is possible to separate out those that may be due to a perturbation of the intrauterine environment of the early embryo from those associated with the development of chronic substrate restriction and subsequent fetal growth restriction. In the present study, we have determined whether uterine carunclectomy alters the functional heterogeneity of corticotrophic cell types within the fetal pituitary, and whether these changes are related to the perturbation of the early intrauterine environment associated with uterine carunclectomy or are solely due to the impact of a chronic restriction of fetal substrate supply in late gestation. We have determined the impact of uterine carunclectomy associated with either fetal hypoxemia or fetal normoxemia on the ACTH-secretory characteristics of pituitary corticotroph cells and on the proportion of corticotroph cells in the fetal anterior pituitary that are CRH-responsive. These results provide insight into the mechanisms by which perturbations of the early and late intrauterine environment may result in changes in the functional characteristics of corticotrophs in the developing pituitary.

Methods
All experiments in the study were carried out according to the guidelines of the Australian Code of Practice for the Care and Use of Animals for Scientific Procedures and approved by the Standing Committee on Ethics and Animal Experimentation at the University of Adelaide.

Animals and surgery. Twenty-six pregnancy-dated Merino × Border Leicester ewes were used in this study. In 15 nonpregnant ewes (carunclectomized group; CR), the majority of visible endometrial caruncles were surgically removed under general anesthesia as previously described (11, 18). Briefly, with the ewe in a supine position, a low midline abdominal incision (approximately 8 cm) was made to allow access to the uterus, and the majority (86–95%) of visible caruncles were excised. After a minimum of 9 weeks’ recovery from surgery, the ewes entered a mating program, and singleton pregnancies were confirmed by ultrasound at approximately 30 days of gestation. The 11 remaining ewes underwent no uterine surgery (non-carunclectomized; NCR) but were mated under the same conditions as the CR ewes.

At 98–117 days of gestation, vascular catheters were inserted into a fetal and a maternal jugular vein and/or carotid artery and into the amniotic cavity in CR (n = 15) and NCR (n = 11) fetuses as previously described (10, 18). All catheters were filled with sterile heparinized saline (500 IU/ml; Multiparin; Fisons Pharmaceuticals, Pennant Hills, New South Wales, Australia) and exteriorized via an incision in the epe’s flank. The ewes were housed under a 12-hour light-dark cycle and fed once daily between two hours and six hours after lights on with water available ad libitum.

Fetal blood sample collection. Fetal arterial blood samples (0.3 ml) were collected for the first 4 days following surgery and then three times per week until and including the day the fetuses were sacrificed. The arterial blood samples were used to measure the partial pressure of oxygen (P(O2)) and carbon dioxide (P(CO2)), pH, hemoglobin content (Hb), and oxygen saturation (SaO2) using an ABL 520 blood gas analyzer (Radiometer, Copenhagen, Denmark) with correction for the higher value of fetal body temperature. Fetal arterial blood samples (2–3.5 ml) were also collected for plasma cortisol determination. Blood samples were centrifuged and plasma separated and stored at −20°C for subsequent assay.

Pituitary collection and cell culture. At 138–145 days of gestation, ewes were killed with an intravenous overdose of sodium pentobarbitone (200 mg/kg; Lethobarb; Virbac PTY Ltd., Peakhurst, New South Wales, Australia). The fetal sheep were delivered via laparotomy, weighed, and killed by decapitation. Each pituitary was quickly removed and immediately placed into cold HEPES-dissociation buffer (HDB) (17). Cultured anterior pituitary cells were prepared as previously described (17) with minor modifications. Briefly, the anterior pituitary and the neurointermediate lobes of each pituitary were gently separated by blunt dissection, and the anterior pituitary tissue was subsequently minced into small fragments, washed in HDB, and placed in collagenase II (0.04%; Worthington Biochemical Corp., Freehold, New Jersey, USA) and deoxyribonuclease I (Sigma-Aldrich, St. Louis, Missouri, USA) with gentle rocking for 2.5 hours at 37°C. The pituitary fragments were then dispersed into cells, which were washed by repeated centrifugation and suspended in 5.0 ml culture medium (DMEM plus Ham’s F12 medium [GIBCO BRL; Life Technologies Inc., Grand Island, New York, USA] containing 10% charcoal-stripped FCS). The cells were counted, and their viability was assessed with trypan blue. The cells were then plated in culture medium (1 ml) at approximately 2.0 × 10^3 cells per well in 48-well tissue culture plates (Falcon; Becton Dickinson Labware, Franklin Lakes, New Jersey, USA) and incubated at 37°C in a water-saturated 5% CO₂ atmosphere (BB16 Gas Incubator; Haraeus Instruments, Hanau, Germany).

Experimental protocol. Twenty-four hours after plating the cells, we treated the cells in half the wells from each pituitary with either vehicle or C-TOX, which is a synthetic hybrid molecule consisting of a CRH analog conjugated with the cellular toxin gelonin, at a final concentration of 25 nmol/l. This treatment specifically...
eliminates CRH-target cells (15–17, 19, 20). All cells were washed extensively in culture medium on the following day and returned to the incubator. Two days after that, the cells were again washed extensively with incubation medium (DMEM/F12 mixed 1:1; Gibco BRL; Life Technologies Inc.) containing 0.2% Polypep (Trace Biosciences, Castle Hill, Australia) and allowed to equilibrate to serum-free conditions for 1 hour. Cells were washed once in incubation medium and incubated for 5 hours in the absence or presence of the experimental treatments: vehicle (also termed basal; n = 26), ovine CRH (10−7 M; n = 26), AVP (10−7 M; n = 23), or ovine CRH + AVP (n = 19) (all peptides supplied by Auspep Pty Ltd., Melbourne, Australia). After 5 hours, the culture media were collected and stored at −20°C until assay. Cellular ACTH content was extracted in 0.1 M HCl (1 ml) by repeated thawing and refreezing. The extracts were stored at −20°C.

**ACTH RIA.** The concentrations of immunoreactive ACTH in the cell culture media and cellular extracts were measured using a double-antibody RIA as previously described and validated (17). The interassay coefficient of variation was 18.3%, and the intra-assay coefficient of variation was less than 10%.

**Cortisol RIA.** Cortisol was extracted from fetal plasma using dichloromethane as previously described (13). The efficiency of recovery of 125I-cortisol from fetal plasma was always greater than 90%. Fetal cortisol concentrations were then measured using an Orion Diagnostica RIA kit (Orion Diagnostica, Turku, Finland). The interassay coefficient of variation was less than 20%, and the intra-assay coefficient of variation was less than 10%.

**Statistics.** Fetal blood gas values are expressed as mean ± SEM. The mean P O2, P CO2, pH, Hb, and S O2 values obtained between 111 and 145 days of gestation. Arterial oxygen content (O2 content) per 100 ml blood (ml/dl) was calculated for each fetus with the equation O2 content = (P O2 × 0.003) + [Hb] × (S O2/100) × 1.39.

Nine of the 15 fetuses in the CR group and 4 of the 11 fetuses in the NCR group had a mean gestational P O2 less than 16 mmHg, and these fetuses were categorized as being hypoxic. The remainder of the fetuses in the CR and NCR groups were normoxic. The separate and combined effects of carunclectomy and hypoxemia on the mean gestational P O2 secretion after secretagogues, and the proportion of ACTH stored in the CRH-responsive corticotrophs.

For all analyses, where the Cochrans and Bartlett-Box tests identified significant heterogeneity of variance, the data were logarithmically transformed prior to ANOVA. Where the multifactorial ANOVAs identified significant interactions between major variables, the data were split on the basis of the interactions and reanalyzed. When the ANOVAs indicated that there were differences between the groups, the Duncan’s post hoc test was used to identify the significant differences (P < 0.05) between the mean values.

**Results**

**Fetal outcomes.** All fetuses were alive at the time of caesarean section. Carunclectomy and hypoxemia each separately resulted in a significant reduction in fetal body weight (Table 1). The S O2, arterial O2 content, and pH were significantly lower in animals in the CR.
group (i.e., CR-normoxemic and CR-hypoxemic fetuses) compared with the NCR group (i.e., NCR-normoxemic and NCR-hypoxemic fetuses) (Table 1; Figure 1). The mean gestational PaO2, SaO2, and arterial O2 content were also significantly lower, and Hb content was significantly higher, in the hypoxemic fetuses in each group when compared with the normoxemic fetuses in the CR and NCR groups (Table 1; Figure 1).

**Plasma cortisol concentrations.** There were no separate or combined effects of either hypoxemia or carunclectomy on plasma cortisol concentrations, which were 90.6 ± 31.5 nmol/l (n = 7) in the hypoxemic fetuses and 45.2 ± 21.0 nmol/l (n = 10) in the normoxemic group.

**Effect of carunclectomy and hypoxemia on total ACTH content.** There was no significant effect of carunclectomy on the total ACTH content when expressed as either total ACTH per 10^4 cells or ng ACTH per well. Cellular ACTH content measured in ng per 10^4 cells was: NCR-normoxemic, 2.26 ± 0.86; NCR-hypoxemic, 1.34 ± 0.36; CR-normoxemic, 1.26 ± 0.29; CR-hypoxemic, 0.89 ± 0.20. Content in ng ACTH per well is summarized in Figure 2. There was also no effect of hypoxemia on the total ACTH present in pituitary cells in either the CR or the NCR group (Figure 2).

**Effect of carunclectomy and hypoxemia on ACTH secretion.** Carunclectomy had a significant effect on the rate of ACTH secretion by the fetal pituitary cells. The percentage of total ACTH that was secreted under basal conditions was significantly greater in pituitary cells from CR fetal sheep than in those from NCR fetal sheep (Table 2). Interestingly, hypoxemia itself had no effect on basal ACTH secretion in the CR and NCR groups (Table 2). There was also no significant correlation between plasma cortisol concentrations and basal ACTH secretion.

**ACTH-secretory responses to the hypothalamic peptides, such that the respective increases in ACTH secretion in the presence of CRH, AVP, or CRH + AVP were significantly lower in the CR than in the NCR group.** Despite this attenuation, the responses represent a significant increase over basal ACTH secretion, and the response to CRH + AVP in combination remained significantly greater than the response to either peptide alone.

Interestingly, in these intact (no C-TOX) populations there was no separate effect of hypoxemia on the ACTH responses to the peptides in either the CR or the NCR group. There was also no significant correlation between plasma cortisol concentrations and the ACTH-secretory responses to CRH, AVP, or to CRH + AVP when the fetuses from all groups were combined.

**Effect of C-TOX and hypoxemia on total ACTH content.** After treatment with C-TOX, the elimination of CRH-target cells was associated with a marked decrease in the total ACTH in the pituitary cells in both the CR and NCR groups, and there was no effect of carunclectomy on this decrease. In contrast, there was a significant interaction between the effects of C-TOX treatment and the effects of hypoxemia on the total ACTH. The relative decrease in total ACTH after C-TOX treatment was less in the hypoxemic groups (CR-hypoxemic and NCR-hypoxemic) than in the normoxemic groups (CR-normoxemic and NCR-normoxemic) (Figure 2). The proportion of ACTH contained in CRH-target cells (i.e., cells susceptible to C-TOX treatment) was therefore less in the hypoxemic than in the normoxemic fetuses in both the CR and NCR groups (Figure 4). There was no significant correlation, however, between plasma cortisol concentrations and the proportion of ACTH contained in the CRH-target cells when the CR and NCR groups were combined.

**Effects of C-TOX on basal and stimulated ACTH secretion.** After C-TOX treatment, there was a similar and significant increase in the fraction of the total ACTH secreted during basal conditions in the CR and the NCR groups, and this increase was not influenced by hypoxemia (Table 2). After elimination of the CRH-target cells, basal ACTH secretion was significantly higher in the CR group than in the NCR group (Table 2). Importantly, there was no separate effect of hypoxemia on basal ACTH secretion after C-TOX treatment (Table 2). The increased proportion of ACTH secreted after C-TOX treatment was primarily a result of the

---

**Table 1**

<table>
<thead>
<tr>
<th>Treatment</th>
<th>P&lt;sub&gt;O&lt;/sub&gt;2</th>
<th>P&lt;sub&gt;CO&lt;/sub&gt;2</th>
<th>pH</th>
<th>Hb (g/dl)</th>
<th>S&lt;sub&gt;O&lt;/sub&gt;2 (%)</th>
<th>O&lt;sub&gt;2&lt;/sub&gt; content (ml/dl)</th>
<th>Weight (kg)</th>
</tr>
</thead>
<tbody>
<tr>
<td>NCR</td>
<td>19.3 ± 0.9</td>
<td>45.8 ± 1.0</td>
<td>7.41 ± 0.01</td>
<td>9.9 ± 0.5</td>
<td>66 ± 3</td>
<td>9.1 ± 0.4</td>
<td>4.49 ± 0.33</td>
</tr>
<tr>
<td>HX</td>
<td>13.8 ± 1.1A</td>
<td>44.6 ± 0.9</td>
<td>7.40 ± 0.01</td>
<td>11.4 ± 0.7A</td>
<td>54 ± 3A</td>
<td>8.5 ± 0.5A</td>
<td>3.45 ± 0.52A</td>
</tr>
<tr>
<td>CR</td>
<td>19.1 ± 0.3</td>
<td>46.2 ± 1.7</td>
<td>7.39 ± 0.01B</td>
<td>10.0 ± 0.5</td>
<td>63 ± 1B</td>
<td>8.8 ± 0.4B</td>
<td>4.04 ± 0.11B</td>
</tr>
<tr>
<td>HX</td>
<td>13.5 ± 0.7A</td>
<td>46.5 ± 1.6</td>
<td>7.38 ± 0.01B</td>
<td>12.2 ± 0.5A</td>
<td>44 ± 3.2A,B</td>
<td>7.4 ± 0.4A,B</td>
<td>2.75 ± 0.38A,B</td>
</tr>
</tbody>
</table>

Mean gestational arterial blood gases and fetal body weight in NCR and CR groups. All values are expressed as mean ± SEM. ASignificant difference between mean values in the normoxemic (NX) and hypoxemic (HX) groups. BSignificant difference between mean values in the NCR and CR groups.
decrease in total ACTH, as C-TOX treatment did not alter the absolute amount of ACTH secreted.

Elimination of CRH-target cells significantly decreased the ACTH responses to all secretagogues. Significantly, there were differences in the effect of C-TOX treatment among the various experimental groups (Figure 5). In NCR groups, there was a small residual response to the hypothalamic peptides. Although these responses were much attenuated when compared with the corresponding ACTH responses in the intact (no C-TOX) populations, there were significant responses to CRH and AVP alone and a greater response to CRH + AVP in combination. Interestingly, the presence or absence of hypoxemia made no significant difference to the fold changes in ACTH secretion. Pituitary cells from the CR fetuses, however, behaved more like cells from the adult sheep pituitary in that there was no response to CRH following elimination of the CRH-target cells. In addition, there was no response to AVP alone and no effect of CRH and AVP in combination. As in the NCR group, there was no significant difference between the CR-hypoxemic and CR-normoxemic fetal pituitary cells.
Discussion

This study has found that there are differential effects of altering the early and late intrauterine environment on the functional characteristics of the corticotroph cells in late gestation. Our results are consistent with the emergence of a population of non–CRH-target cells that secrete high amounts of ACTH in the pituitaries of fetuses that were exposed to uterine carunclectomy in early gestation. The emergence of this subpopulation of corticotrophs in the CR group during late gestation occurred whether or not these fetuses were hypoxic throughout late gestation. Exposure to chronic hypoxemia during late gestation, however, independently resulted in a specific reduction in the proportion of ACTH stored in CRH-target cells. Thus, the ACTH-synthetic capacity of the CRH-target cells, which do not appear to contribute to basal ACTH secretion, but which secrete ACTH in response to a superimposed acute intrauterine stress, may be diminished in the chronically hypoxic fetus during late gestation.

Carunclectomy and fetal substrate restriction. In the current study, 9 of 15 fetuses in the CR and 4 of 11 fetuses in the NCR group were chronically hypoxic, with mean gestational $P_aO_2$ values of $\leq 16$ mmHg (21, 22). In the remaining fetal sheep, the mean $P_aO_2$ was over 19 mmHg, well within the reported normoxic range (11, 13, 18, 21, 22). While the mean values for fetal arterial pH were marginally less in the CR than in the NCR group overall, it should be noted that the arterial pH values in both the CR-normoxic and the CR-hypoxic groups were well within the normal range for healthy fetal sheep in late gestation, i.e., this group was not acidemic. Fetal acidaemia is therefore not a contributing factor to the changes that occurred in the ACTH-synthetic and -secretory characteristics of the corticotrophs in the CR group. It has also previously been reported that fetal sheep that are chronically hypoxic may also be chronically hypoglycemic in late gestation (13). In a recent study on normoxic fetuses that had either low ($<1.2$ mmol/l) or normal ($>1.2$ mmol/l) mean plasma glucose concentrations throughout late gestation, we found that chronic hypoglycemia was not associated with any differences in the basal ACTH-secretory rate, in the ACTH responses to CRH, AVP, or CRH + AVP, or in the proportion of ACTH stored in CRH-responsive corticotrophs (our unpublished observations). Thus we suggest that in the CR-hypoxic and NCR-hypoxic groups it is the prevailing hypoxemia during late gestation that results in a specific reduction in the proportion of ACTH stored in CRH-target cells. Furthermore, we propose that the changes in the ACTH-secretory characteristics of the corticotrophs in the CR groups are a consequence of the early intrauterine intervention, as there is no evidence for restriction of a fetal substrate present in both the CR-normoxic and CR-hypoxic groups.

Impact of carunclectomy on basal ACTH secretion. The percentage of ACTH secreted under basal conditions was greater in the CR than in the NCR fetuses, and this effect was independent of the prevailing fetal $P_aO_2$. As argued above, it is therefore unlikely that this increase in basal ACTH secretion is related to the effects of placental restriction experienced by fetuses in vivo in late gestation in the CR group. Interestingly, the higher basal output of ACTH was maintained in the CR group after C-TOX treatment, reflecting maintained ACTH secretion despite a decrease in total ACTH content.

Several groups, including ours, have described functional heterogeneity of corticotrophs on the basis of responses to specific hypothalamic peptides, including CRH and AVP (14–17, 23). In adult sheep anterior pituitaries, treatment with C-TOX eliminates the response to CRH, while basal ACTH secretion is maintained, or increased, and a residual response to AVP remains (14, 15). On this basis it has been concluded that there are at least two populations of corticotrophs, one responsive to CRH and AVP (and susceptible to C-TOX), and one that secretes the majority of ACTH under basal conditions and that responds to AVP but not CRH.

Table 2

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Vehicle</th>
<th>C-TOX</th>
</tr>
</thead>
<tbody>
<tr>
<td>NCR Normoxic</td>
<td>1.52 ± 0.31</td>
<td>5.02 ± 0.96 $^a$</td>
</tr>
<tr>
<td>Hypoxic</td>
<td>1.85 ± 0.24</td>
<td>4.21 ± 0.65 $^a$</td>
</tr>
<tr>
<td>CR Normoxic</td>
<td>2.55 ± 0.61 $^a$</td>
<td>5.78 ± 1.35 $^{AB}$</td>
</tr>
<tr>
<td>Hypoxic</td>
<td>5.30 ± 1.52 $^a$</td>
<td>8.38 ± 2.31 $^{AB}$</td>
</tr>
</tbody>
</table>

The percentage of total ACTH secreted under basal conditions in the presence of either vehicle or C-TOX. All values are expressed as mean ± SEM. $^a$Significant difference between mean values in the NCR and CR groups. $^AB$Significant difference between vehicle and C-TOX treatment.
A similar conclusion had been reached in studies using the reverse hemolytic plaque assay in anterior pituitary cells from the adult rat (24). The maintenance of basal ACTH secretion after treatment with C-TOX is consistent with findings from previous studies on anterior pituitary cells from the fetal sheep, which have described the presence of at least one type of corticotroph that accounts for basal ACTH secretion and is resistant to C-TOX pretreatment (17).

In the current study, there was no difference in the ACTH-secretory responses to AVP alone or in combination with CRH in the CR group after C-TOX treatment. This is in contrast to the NCR group, where the ACTH responses to AVP in combination with CRH were greater than those to CRH alone after C-TOX treatment. This implies that the ACTH response to AVP in the pituitaries of CR fetuses in the absence of C-TOX treatment is derived from cells that are responsive to both AVP and CRH. While previous studies with C-TOX in pituitary cells from adult sheep have not provided evidence for a population of corticotrophs that secretes ACTH under basal conditions but does not respond to AVP, such cells have been described in studies using the reverse hemolytic plaque assay in adult rat anterior pituitary cells (23).

In the present study, pituitary cells from normoxemic NCR fetuses had levels of basal ACTH secretion similar to those measured in previous studies on pituitary cells collected from healthy, term fetal sheep (17). Consistent with previously observed effects of C-TOX (17), pituitaries from normoxemic NCR and CR fetuses in the present studies showed an increase in basal ACTH secretion in the normoxemic NCR group after C-TOX treatment.

Interestingly, the ACTH responses to CRH and to CRH + AVP in the NCR group were higher after C-TOX than in the CR group. This contrasts with previous studies with C-TOX in adult ovine pituitary cells, in which there was no response to CRH, either alone or in combination with AVP, in the CRH-target–depleted populations. One explanation is that the C-TOX was ineffective in eliminating CRH-target cells. This would not appear to be the case, since pituitaries removed from NCR and CR fetuses were treated identically—with different effects in the two groups. Another explanation for these findings is that in the healthy developing pituitary there is increased functional plasticity, such that cells that do not have CRH receptors at the time of treatment with C-TOX and are therefore not susceptible to the toxin survive the treatment and subsequently become CRH-target corticotrophs. These surviving cells might be characterized as normally secreting ACTH under basal conditions, but they only develop CRH receptors and CRH sensitivity in culture. Alternatively, there might be some type of multipotential or stem cell that becomes a corticotroph after the treatment with C-TOX (24). We have shown in rat corticotrophs that a limited response to CRH reemerges at around 3 days after C-TOX treatment (25). In addition, Jia and coworkers have described the emergence of a new population of CRH-responsive corticotrophs following the elimination of the existing CRH-target cells by photoablation (24). Although the reemergence of an ACTH response to CRH has not been observed previously in ovine cells, it should be noted that the developmental plasticity of fetal ovine pituitary cells is likely to be higher than that of adult cells. During development, the turnover rate for corticotrophs is extremely high and is probably determined by mitosis and apoptosis as well as differentiation (26, 27). Numerous examples exist of pituitary plasticity involving melanotrophs, somatotrophs, lactotrophs, and gonadotrophs, in which there is interchange or transdifferentiation in the course of normal physiological adjustments (28–31).

It is particularly noteworthy that the greater response to CRH after C-TOX treatment is a characteristic of pituitary cells from NCR, rather than CR, fetuses. This combination with AVP, in the CRH-target–depleted populations. One explanation is that the C-TOX was ineffective in eliminating CRH-target cells. This would not appear to be the case, since pituitaries removed from NCR and CR fetuses were treated identically—with different effects in the two groups. Another explanation for these findings is that in the healthy developing pituitary there is increased functional plasticity, such that cells that do not have CRH receptors at the time of treatment with C-TOX and are therefore not susceptible to the toxin survive the treatment and subsequently become CRH-target corticotrophs. These surviving cells might be characterized as normally secreting ACTH under basal conditions, but they only develop CRH receptors and CRH sensitivity in culture. Alternatively, there might be some type of multipotential or stem cell that becomes a corticotroph after the treatment with C-TOX (24). We have shown in rat corticotrophs that a limited response to CRH reemerges at around 3 days after C-TOX treatment (25). In addition, Jia and coworkers have described the emergence of a new population of CRH-responsive corticotrophs following the elimination of the existing CRH-target cells by photoablation (24). Although the reemergence of an ACTH response to CRH has not been observed previously in ovine cells, it should be noted that the developmental plasticity of fetal ovine pituitary cells is likely to be higher than that of adult cells. During development, the turnover rate for corticotrophs is extremely high and is probably determined by mitosis and apoptosis as well as differentiation (26, 27). Numerous examples exist of pituitary plasticity involving melanotrophs, somatotrophs, lactotrophs, and gonadotrophs, in which there is interchange or transdifferentiation in the course of normal physiological adjustments (28–31).

It is particularly noteworthy that the greater response to CRH after C-TOX treatment is a characteristic of pituitary cells from NCR, rather than CR, fetuses. This

![Figure 3](image-url)

**Figure 3**
Fold change in ACTH secretion (mean ± SEM) after CRH, AVP, or CRH + AVP administration in the NCR (white bars) and CR (gray bars) groups in the absence of C-TOX treatment. The horizontal dashed line denotes no increase in secretion over basal rates. *P < 0.05, CR vs. NCR groups; **P < 0.05, CRH + AVP vs. CRH or AVP alone.

![Figure 4](image-url)

**Figure 4**
The proportion of total ACTH (mean ± SEM) present in CRH-target cells in the NCR and CR groups. Light gray, hypoxemia; medium gray, maternal carunclectomy; dark gray, hypoxemia and maternal carunclectomy. *P < 0.05, HX versus NX.
suggests that the plasticity of the fetal pituitary, apparently characteristic of the control fetuses, is lacking in CR fetuses. Whether or not CR specifically has an effect on multipotential pituitary cells or transdifferentiation of other cells remains an interesting question. "Impact of carunclectomy on corticotrophic cell types in the fetal pituitary in late gestation. While the basal output of ACTH was greater in the CR group, there was no difference in the proportion of ACTH present in CRH-responsive cells and with a trend toward the reemergence of CRH sensitivity after C-TOX treatment. This effect was present to a greater extent in the NCR group. It has been demonstrated that fetal plasma cortisol concentrations are higher in the chronically hypoxemic fetus in late gestation (13, 21). Furthermore, cortisol can act to suppress ACTH synthesis in CRH-target corticotrophs (15, 17). In the present study, however, while there was a trend toward higher circulating cortisol concentrations in the hypoxemic fetuses, we found no evidence for an inverse correlation between circulating cortisol and the proportion of ACTH stored in the CRH-responsive cells in the pituitaries from the fetal sheep in the NCR and CR groups. Alternatively, the effect of chronic hypoxemia on the amount of ACTH stored in the CRH-target cells may reflect a prior history of hypothalamic stimulation in vivo as a consequence of the low P_{\text{O}_2}. "Summary. In summary, our results are consistent with the emergence of a population of non–CRH-target cells that secrete high amounts of ACTH in the pituitaries of fetuses that were exposed to the CR uterine environment in early gestation. This may represent a reprogramming of the pattern of corticotroph development to ensure that fetal ACTH secretion can be maintained throughout late gestation independently of whether placental restriction of fetal substrate supply subsequently ensues. Given that altered corticotroph development persists in otherwise metabolically healthy fetuses in the CR group for up to 5 months after the early intrauterine intervention, it may also persist after birth to result in a maintained ACTH response to postnatal stressors. In this context, it is interesting that children and adults with lower birth weights have been reported to excrete more cortisol or its metabolites in urine (4, 35). Furthermore a study on adult men found that low birth weight and/or features of the metabolic syndrome were associated with measures of increased activity of the HPA axis (3). Our study provides one potential mechanism to explain the associations between low birth weight, increased adult HPA activity, and the metabolic syndrome. While exposure to chronic hypoxemia during late gestation did not appear to alter the subpopulations of corticotroph cell types present within the fetal pituitary, it did result in a specific reduction in the proportion of ACTH stored in CRH-target cells. Thus, the ACTH-synthetic capacity of the CRH-target cells, which do not appear to contribute to basal ACTH secretion, but which secrete ACTH in response to a superimposed acute intrauterine stress, may be alterations in fetal growth patterns, fetal HPA function, and gestation length (32–34). Thus the reprogramming of early pituitary development may be a key response to perturbations of the interaction between the uterine endometrium and the developing embryo. "Impact of hypoxemia on corticotrophic cell types. Chronic hypoxemia was associated with a decrease in the amount of ACTH stored in CRH-responsive cells and with a trend toward the reemergence of CRH sensitivity after C-TOX treatment. This effect was present to a greater extent in the NCR group. It has been demonstrated that fetal plasma cortisol concentrations are higher in the chronically hypoxemic fetus in late gestation (13, 21). Furthermore, cortisol can act to suppress ACTH synthesis in CRH-target corticotrophs (15, 17). In the present study, however, while there was a trend toward higher circulating cortisol concentrations in the hypoxemic fetuses, we found no evidence for an inverse correlation between circulating cortisol and the proportion of ACTH stored in the CRH-responsive cells in the pituitaries from the fetal sheep in the NCR and CR groups. Alternatively, the effect of chronic hypoxemia on the amount of ACTH stored in the CRH-target cells may reflect a prior history of hypothalamic stimulation in vivo as a consequence of the low P_{\text{O}_2}. "Summary. In summary, our results are consistent with the emergence of a population of non–CRH-target cells that secrete high amounts of ACTH in the pituitaries of fetuses that were exposed to the CR uterine environment in early gestation. This may represent a reprogramming of the pattern of corticotroph development to ensure that fetal ACTH secretion can be maintained throughout late gestation independently of whether placental restriction of fetal substrate supply subsequently ensues. Given that altered corticotroph development persists in otherwise metabolically healthy fetuses in the CR group for up to 5 months after the early intrauterine intervention, it may also persist after birth to result in a maintained ACTH response to postnatal stressors. In this context, it is interesting that children and adults with lower birth weights have been reported to excrete more cortisol or its metabolites in urine (4, 35). Furthermore a study on adult men found that low birth weight and/or features of the metabolic syndrome were associated with measures of increased activity of the HPA axis (3). Our study provides one potential mechanism to explain the associations between low birth weight, increased adult HPA activity, and the metabolic syndrome. While exposure to chronic hypoxemia during late gestation did not appear to alter the subpopulations of corticotroph cell types present within the fetal pituitary, it did result in a specific reduction in the proportion of ACTH stored in CRH-target cells. Thus, the ACTH-synthetic capacity of the CRH-target cells, which do not appear to contribute to basal ACTH secretion, but which secrete ACTH in response to a superimposed acute intrauterine stress, may be
relatively diminished in the chronically hypoxic fetus during late gestation. It has previously been reported that there is a decrease in pituitary proopiomelanocortin mRNA levels, while basal circulating immunoreactive ACTH concentrations are maintained in the hypoxic CR fetus in late gestation. Thus it is possible that circulating ACTH levels are maintained as a consequence of secretion from non-CRH-target corticotrophs while the expression of proopiomelanocortin is suppressed in the CRH-target cells. These patterns of altered corticotroph development are important given the central roles of the HPA axis in the fetal adaptive response to intrauterine stress and in the early programming of adult disease.

**Acknowledgments**

This work was funded by the National Health and Medical Research Council. The authors gratefully acknowledge the expert assistance of Michael Adams, Frank Carbone, Lisa Edwards, Anne Jurisevic, Jacob Ross, and Kirsty Warnes with the animal experiments.