

# The role of glucocorticoid in the regulation of prostaglandin biosynthesis in non-pregnant bovine endometrium

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## Abstract

To determine whether glucocorticoids (GCs) play a role in regulating uterine function in cow, the present study examined the expression of mRNA encoding GC receptor (GC-R)  $\alpha$ , 11 $\beta$ -hydroxysteroid dehydrogenase (11-HSD) type 1 and type 2, and the activity of 11-HSD1 in bovine endometrial tissue throughout the estrous cycle. We also studied the effects of cortisol on basal, oxytocin (OT)- and tumor necrosis factor- $\alpha$  (TNF $\alpha$ )-stimulated prostaglandin (PG) production. A quantitative real-time PCR analysis revealed that GC-R $\alpha$  mRNA was expressed more strongly in the mid-luteal stage (days 8–12) than in the other stages. In contrast to GC-R $\alpha$  mRNA expression, 11-HSD1 mRNA expression was greater in the follicular stage than in the other stages, whereas 11-HSD2 mRNA expression was lowest in the follicular stage. The activity of 11-HSD1 was greater in

the follicular stage and estrus than in the other stages and was lowest in the mid-luteal stage. Cortisone was dose-dependently converted to cortisol in the cultured endometrial tissue. Although cortisol did not affect either the basal or OT-stimulated production of PGs in the cultured epithelial cells, the production of PGs stimulated by TNF $\alpha$  in the stromal cells was suppressed by cortisol ( $P < 0.05$ ). Cortisol suppressed basal prostaglandin (PG)F2 $\alpha$  without affecting basal PGE2 production in the stromal cells. The overall results suggest that the level of cortisol is locally regulated in bovine endometrium throughout the estrous cycle by 11-HSD1, and that cortisol could act as a luteoprotective factor by selectively suppressing luteolytic PGF2 $\alpha$  production in bovine endometrium.

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## Introduction

Glucocorticoids (GCs) are involved in various physiological processes, including general metabolism (Wang 2005), immunological response (McKay & Cidlowski 1998, 1999), and female reproductive function (Brann & Mahesh 1991, Andersen 2002). At high concentrations, GCs suppress most immunological responses and are well known as anti-inflammatory agents limiting the production of cytokines and prostaglandins (PGs) in various target organs (Goppelt-Strube *et al.* 1996, Goppelt-Strube 1997, Hillier & Tetsuka 1998). Several uterine events such as menstruation, implantation, and parturition have been likened to an inflammatory process (Salamonsen & Lathbury 2000). Because GCs have been shown to exert specific effects on uterine physiology in rats (Rabin *et al.* 1990, Korgun *et al.* 2003), rabbits (Bigsby & Everett 1991), humans (Gellersen *et al.* 1994), and ewes (Monheit & Resnik 1981, Gupta *et al.* 2003), the uterus is

considered to be a target organ for GCs in some species. However, the roles of GCs in the bovine endometrium remain unknown.

The endometrium is a complex tissue and mainly consists of epithelial and stromal cells (Fortier *et al.* 1988). Although both types of endometrial cells have the capacity to produce PGs, they have specific morphological and physiological properties (Asselin *et al.* 1997, Fortier *et al.* 1988, Miyamoto *et al.* 2000). We found that oxytocin (OT) stimulates PG production only in epithelial cells, while tumor necrosis factor- $\alpha$  (TNF $\alpha$ ) stimulates PG production only in stromal cells (Skarzynski *et al.* 2000). This cell-specific response to OT and TNF $\alpha$  is a useful parameter for investigating the physiology and endocrine status of cultured bovine endometrial cells.

The biological action of GCs is mediated through the activation of intracellular GC receptors (GC-R). Two isoforms of GC-R, GC-R $\alpha$  and GC-R $\beta$ , have been identified (Giguere

*et al.* 1986, Funder 1993). Access of GCs to GC-R in target tissues is regulated by two 11 $\beta$ -hydroxysteroid dehydrogenases (11-HSDs), a bidirectional 11-HSD type 1 (11-HSD1) that mainly converts cortisone to active cortisol (Stewart & Mason 1995) and 11-HSD type 2 (11-HSD2) that inactivates cortisol to cortisone (Albiston *et al.* 1994, Stewart *et al.* 1994). Therefore, cyclic changes of the expressions of GC-R and 11-HSDs mRNA could help to define the roles of GCs in uterine physiology.

In the present study, to determine whether GCs play a role in regulating bovine uterine function, we examined 1) the temporal patterns of GC-R $\alpha$ , 11-HSD1, and 11-HSD2 mRNA expressions, and 11-HSD1 activity in bovine endometrium throughout the estrous cycle and 2) the effects of cortisol on basal and OT- or TNF $\alpha$ -stimulated PG production in the cultured endometrial epithelial and stromal cells.

## Materials and methods

### Collection of endometrial tissues

Uteri of Holstein cows were obtained from a local abattoir in accordance with protocols approved by the local Institutional Animal Care and Use Committee. Apparently, healthy uteri without a visible conceptus were obtained within 10–20 min after exsanguination and immediately transported to the laboratory on ice. The stages of the estrous cycle were determined by macroscopic observation of the ovary and uterus as described previously (Okuda *et al.* 1988, Miyamoto *et al.* 2000). For mRNA determination, endometrial tissues ( $n=8$ /stage) were collected from cows at six different stages of the estrous cycle (estrus, day 0; early luteal, days 2–3; developing luteal, days 5–6; mid-luteal, days 8–12; late luteal, days 15–17, and follicular stage, days 19–21). Intercaruncular endometrial tissues from the uterine horn, ipsilateral to the corpus luteum, were used for all experiments. The endometrial tissues were immediately frozen rapidly in liquid nitrogen and stored at  $-80^{\circ}\text{C}$  until processed for RNA isolation. For experiments involving tissue and cell cultures, the uterus was submerged in ice-cold physiological saline and transported to the laboratory.

*Experiment 1: determination of GC-R $\alpha$ , 11-HSDs mRNA expressions and 11-HSD1 activity throughout the estrous cycle*

**Reverse transcription and real-time PCR** Total RNA was extracted from endometrial tissue using TRIZOL reagent (Invitrogen) according to the manufacturer's directions. One microgram of each total RNA was reverse transcribed using a ThermoScript RT-PCR System (Invitrogen) and 10% of the reaction mixture was used in each PCR using specific primers for GC-R $\alpha$  and 11-HSDs from the bovine sequence. The primers were chosen using an online software package ([http://www-genome.wi.mit.edu/cgi-bin/primer/primer3\\_www.cgi](http://www-genome.wi.mit.edu/cgi-bin/primer/primer3_www.cgi)).

Gene expression was measured by real-time PCR using the Mx3000P QPCR System (Stratagene, La Jolla, CA, USA) and the QuantiTect SYBR Green PCR system (Qiagen) starting with 2 ng reverse-transcribed total RNA as described previously (Sakumoto *et al.* 2005; Table 1). Briefly, GAPDH expression was used as an internal control. For quantification of the mRNA expression levels, the primer length (20 bp) and GC contents of each primer (50–60%) were selected, and PCR was performed under the following conditions:  $95^{\circ}\text{C}$  for 15 min, followed by 55 cycles of  $94^{\circ}\text{C}$  for 15 s,  $55^{\circ}\text{C}$  for 20 s, and  $72^{\circ}\text{C}$  for 15 s. Use of the QuantiTect SYBR Green PCR system at elevated temperatures resulted in reliable and sensitive quantification of the RT-PCR products with high linearity (Pearson's correlation coefficient ( $r>0.99$ )).

**11-HSD1 activity** The level of 11-HSD reductase activity in endometrial tissue was determined by measuring the net conversion rate from cortisone to cortisol. Briefly, endometrial strips (30 mg) were placed in glass culture tubes (12 mm $\times$ 75 mm) containing 2 ml culture medium (Dulbecco's modified Eagle's medium (DMEM)/Ham's F-12; 1:1 (v/v); Sigma) supplemented with 0.1% (w/v) BSA, 100 IU/ml penicillin, and 100  $\mu\text{g/ml}$  streptomycin with 5%  $\text{CO}_2$  in air. The endometrial tissues were exposed to cortisone (0, 3, 30, 300, and 1000 nM) in a shaking water bath at  $38^{\circ}\text{C}$  for 4 h. Media containing cortisone (0, 3, 30, 300, and 1000 nM) without tissues were incubated for 4 h for the blank value and to determine non-specific interconversion. At the end of incubation, 1 ml conditioned medium was

**Table 1** Primers for real-time PCR

Gene	Primer	Sequence	Accession no.	Product (bp)
GC-R $\alpha$	Forward	5'-CCATTCTGTTCACGGTGTG-3'	AY238475	132
	Reverse	5'-CTGAACCGACAGGAATTGGT-3'		
11-HSD1	Forward	5'-ACTCTGCGCCAAGATGAAGT-3'	AF548027	149
	Reverse	5'-TAGCCCTCAGGAAGTGCCTA-3'		
11-HSD2	Forward	5'-CCTAGACCGGATCCTTCTCC-3'	AF074706	114
	Reverse	5'-ACCTTGGGGGTCAGAATACC-3'		
GAPDH	Forward	5'-CACCCCTCAAGATTGTCAGCA-3'	BC102589	103
	Reverse	5'-GGTCATAAGTCCCTCCACGA-3'		

collected and frozen at  $-30^{\circ}\text{C}$  until the cortisol assay. The tissues were blotted on filter paper and weighed. The specific conversion rate from cortisone to cortisol was calculated, and the blank values (defined as the amount of conversion in the absence of tissue) were subtracted and expressed as picogram of cortisol converted per milligram of tissue (pg/mg tissue). To examine the cyclic changes in 11-HSD1 activity throughout the estrous cycle, the endometrial tissues from the estrus, early luteal, developing luteal, mid-luteal, late luteal, and follicular stages ( $n=4/\text{stage}$ ) were exposed to cortisone (30 nM).

**Isolation of endometrial cells** For cell culture, endometrial tissues were obtained at the early luteal phase (days 2–5). The epithelial and stromal cells from the bovine endometrium were separated using a modification of procedures described previously (Skarzynski *et al.* 2000). A polyvinyl catheter was inserted into the side of the oviduct and the ends of the horn were tied in order to retain trypsin solution for detaching the epithelial cells as described below. The uterine lumen was washed thrice with 30–50 ml sterile  $\text{Ca}^{2+}$ - and  $\text{Mg}^{2+}$ -free Hanks' balanced salt solution (HBSS) supplemented with 100 IU/ml penicillin, 100  $\mu\text{g}/\text{ml}$  streptomycin, and 0.1% (w/v) BSA (Roche Diagnostics). Thirty to fifty milliliters of sterile HBSS containing 0.3% (w/v) trypsin (Sigma) was then infused into the uterine lumen through the catheter. Epithelial cells were isolated by incubation at  $38^{\circ}\text{C}$  for 60 min with gentle shaking.

After collection of the epithelial cells, the uterine lumen was washed with sterile HBSS supplemented with antibiotics and 0.1% (w/v) BSA. The horn was then cut transversely with scissors into several segments, which were slit to expose the endometrial surface. Intercaruncular endometrial strips were dissected from the myometrial layer with a scalpel and washed once in 50 ml sterile HBSS containing antibiotics. The endometrial strips were then minced into small pieces (1  $\text{mm}^3$ ). The minced tissues ( $\approx 5$  g) were digested by stirring for 60 min in 50 ml sterile HBSS containing 0.05% (w/v) collagenase (Sigma), 0.005% (w/v) DNase I (Sigma) and 0.1% (w/v) BSA. The dissociated cells were filtered through metal meshes (100 and 80  $\mu\text{m}$ ) to remove undissociated tissue fragments. The filtrate was washed thrice by centrifugation (10 min at 100 g) with DMEM (Sigma) supplemented with antibiotics and 0.1% (w/v) BSA. After the washes, the cells were counted with a hemocytometer. Cell viability was higher than 85% as assessed by 0.5% (w/v) trypan blue exclusion.

**Culture of endometrial cells** The final pellets of the isolated stromal or epithelial cells were resuspended in culture medium (DMEM/Ham's F-12; 1:1 (v/v); Sigma) supplemented with 10% (v/v) calf serum (Sigma), 20  $\mu\text{g}/\text{ml}$  gentamicin (Invitrogen), and 2  $\mu\text{g}/\text{ml}$  amphotericin B (Sigma; Skarzynski *et al.* 2000). The stromal cells were seeded at a density of  $1 \times 10^5$  viable cells/ml in 48-well cluster dishes (Costar, Cambridge, MA, USA), and the epithelial cells were

seeded at a density of  $1 \times 10^5$  viable cells/ml in culture flasks (Nunc) and cultured at  $38^{\circ}\text{C}$  in a humidified atmosphere of 5%  $\text{CO}_2$  in air. To purify the stromal preparation, the medium was changed 2 h after plating, by which time selective attachment of stromal cells had occurred (Fortier *et al.* 1988, Skarzynski *et al.* 2000). Alternatively, since the epithelial cells attached 24–48 h after plating, the medium in the epithelial cell culture was replaced 48 h after plating. The medium was changed every 2 days until the cells reached confluence. When the epithelial cells were confluent, 0.02% (w/v) trypsin solution was added to the cells to remove the other cells. After removal of the other cells, 0.25% (w/v) trypsin solution was then added to the epithelial cells to collect the pure epithelial cells. The cells were removed, adjusted to a density of  $1 \times 10^5$  cells/ml, and placed in 48-well cluster dishes for DNA quantification in fresh DMEM/Ham's F-12 supplemented with 10% (v/v) calf serum, 20  $\mu\text{g}/\text{ml}$  gentamicin, and 2 mg/ml amphotericin B until the cells reached confluence. The homogeneity of stromal and epithelial cells was evaluated using immunofluorescent staining for specific markers of epithelial (cytokeratin) and stromal cells (vimentin; Murakami *et al.* 2003) as described previously. The epithelial cell contamination of stromal cells was about 1% and stromal cell contamination of epithelial cells was  $<1\%$ . When cells of each type were confluent (6–7 days after the start of the culture), the medium was replaced with fresh DMEM/Ham's F-12 supplemented with 0.1% (w/v) BSA, 5 ng/ml sodium selenite (Sigma), 0.5 mM ascorbic acid (Wako Pure Chemical Industries, Ltd, Osaka, Japan), 5  $\mu\text{g}/\text{ml}$  transferrin (Sigma), 2  $\mu\text{g}/\text{ml}$  insulin (Sigma), and 20  $\mu\text{g}/\text{ml}$  gentamicin. The cells were then exposed to various stimulants for Experiment 2.

#### *Experiment 2: effect of cortisol on basal and OT- or TNF $\alpha$ -stimulated PGF2 $\alpha$ and PGE2 production by epithelial and stromal cells*

Epithelial cells were exposed to cortisol (0.1–100 nM), OT (100 nM; Teikoku Hormone MFG Co., Tokyo, Japan), or cortisol in combination with OT for 24 h. Stromal cells were exposed to cortisol (0.1–100 nM), TNF $\alpha$  (0.06 nM; Dainippon Pharmaceutical Co., Ltd, Osaka, Japan), or cortisol in combination with TNF $\alpha$  for 24 h. The concentrations of OT and TNF $\alpha$  were based on a previous study (Skarzynski *et al.* 2000). Media with supplements without stimulants incubated with cells were used as controls.

After culture, the conditioned media were collected in 1.5 ml tubes containing 5  $\mu\text{l}$  of a stabilizer solution (0.3 M EDTA, 1% (w/v) acid acetyl salicylic, pH 7.3) and frozen at  $-30^{\circ}\text{C}$  until the PGs assay. The DNA content, estimated by the spectrophotometric method of Labarca & Paigen (1980), was used to standardize the results.

**PG and cortisol determination** The concentrations of PGF2 $\alpha$  and PGE2 in the culture medium were determined by enzyme immunoassay (EIA) as described previously (Woclawek-Potocka *et al.* 2004). The PGF2 $\alpha$  standard

curve ranged from 0.016 to 4 ng/ml and the ED<sub>50</sub> of the assay was 0.25 ng/ml. The intra- and inter-assay coefficients of variation were on average 2.8 and 7.7% respectively. The PGE<sub>2</sub> standard curve ranged from 0.39 to 100 ng/ml and the ED<sub>50</sub> of the assay was 6.25 ng/ml. The intra- and inter-assay coefficients of variation were on average 3.1 and 8.6% respectively. The EIA for cortisol was done as described previously (Acosta *et al.* 2002). The standard curve ranged from 0.1 to 400 ng/ml and the ED<sub>50</sub> of the assay was 1.6 ng/ml. The intra- and inter-assay coefficients of variation were on average 5.4 and 6.0% respectively. The cross-reactivities of the polyclonal antibody (raised in a rabbit against cortisol-3-carboxymethyloxime (CMO); Cosmo Bio Co., Tokyo, Japan) were 100% for cortisol, 0.6% for cortisone, 5.7% for 11-deoxycortisol, 0.5% for 21-deoxycortisol, 4.1% for 11-deoxycorticosterone, 1.2% for corticosterone, 0.7% for 17 $\alpha$ -hydroxy progesterone, and 0.02% for 20-dihydroxy progesterone.

### Statistical analysis

Experimental data are shown as the mean  $\pm$  S.E.M. of values obtained in four to five separate experiments, each performed in triplicate. Endometrial cells and tissues collected from different cows were cultured separately. Data on the effects of cortisol, TNF $\alpha$ , and OT on absolute concentrations of PGF<sub>2</sub> $\alpha$  and PGE<sub>2</sub> were statistically analyzed and are shown as a fold change of the control. The statistical significance of differences in concentrations of PG in culture media between the control and treated groups and the mRNA expressions was assessed by one-way ANOVA followed by Fisher's protected least-significant difference procedure (PLSD) as a multiple comparison test by StatView (Version 4.58; Abacus Concepts, Inc. Berkeley, CA, USA).

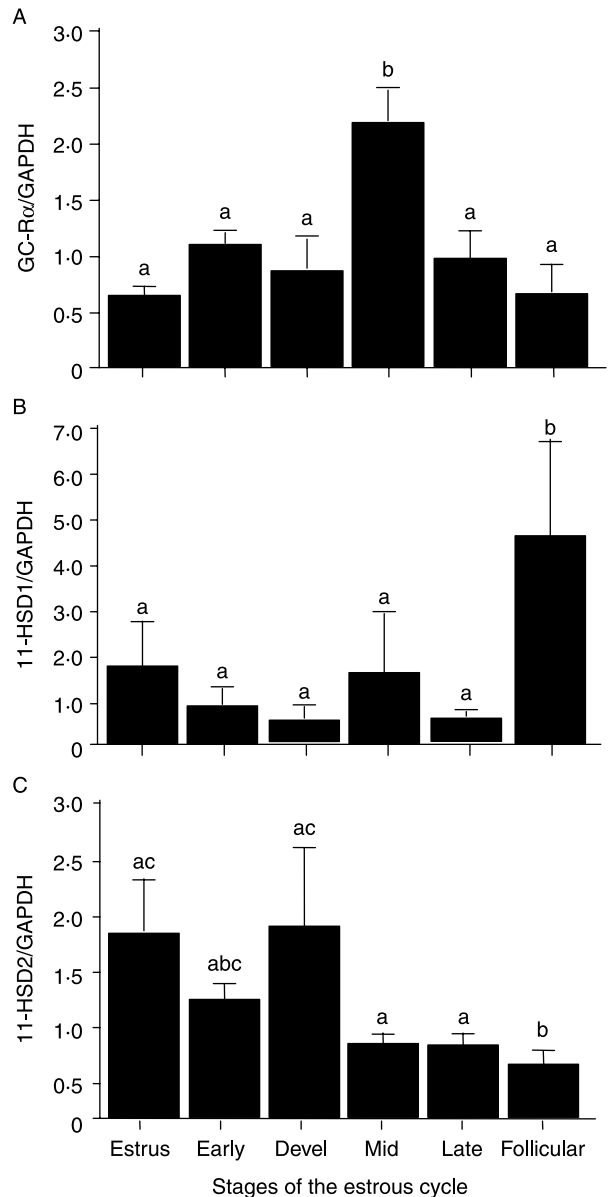
## Results

### mRNAs for GC-R $\alpha$ , 11-HSD1, and 11-HSD2 during the estrous cycle

Specific transcripts for GC-R $\alpha$ , 11-HSD1, and 11-HSD2 were detected in bovine endometrium throughout the estrous cycle. A real-time PCR analysis of GC-R $\alpha$ , 11-HSD1, and 11-HSD2 mRNA in the endometrial tissue during the estrous cycle is shown in Fig. 1. The level of mRNA for GC-R $\alpha$  was greater in the mid-luteal stage (days 8–12) than in the other stages (Fig. 1A;  $P < 0.05$ ). In contrast, the level of mRNA for 11-HSD1 was greater in the follicular stage than in the other stages (Fig. 1B;  $P < 0.05$ ), whereas the level of mRNA for 11-HSD2 was lowest in the follicular stage (Fig. 1C;  $P < 0.05$ ).

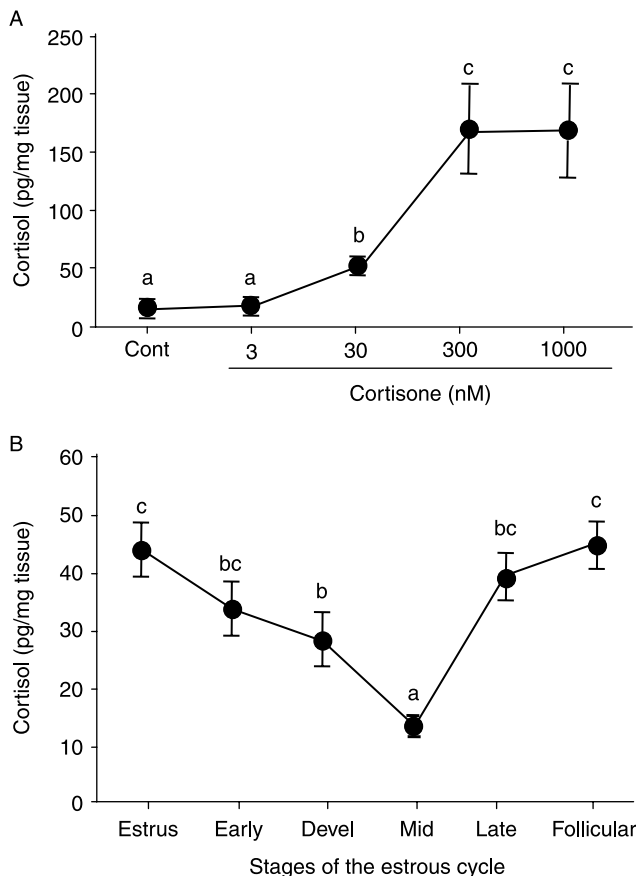
### 11-HSD1 activity throughout the estrous cycle

Endometrial tissue has the capacity to convert cortisone to cortisol as indicated by a significant increase in cortisol



**Figure 1** Changes in relative amounts of (A) GC-R $\alpha$ , (B) 11-HSD1, and (C) 11-HSD2 mRNA in bovine endometrium throughout the estrous cycle. Data are the mean  $\pm$  S.E.M. for eight samples/stage and are expressed as the relative ratio of GC-R $\alpha$  and 11-HSDs mRNA to GAPDH mRNA (estrus, day 0; early luteal, days 2–3; developing luteal, days 5–6; mid-luteal, days 8–12; late luteal, days 15–17; and follicular stage, days 19–21). Different superscript letters indicate significant difference ( $P < 0.05$ ), as determined by ANOVA followed by a Fisher's PLSD as a multiple comparison test.

content in the medium incubated with tissue compared with those incubated without tissue. The concentration of converted cortisol in the media increased with the dose of cortisone (Fig. 2A). The activity of 11-HSD1 was lowest in the mid-luteal stage and greater in the follicular stage and estrus than in the other stages (Fig. 2B). It was shown that



**Figure 2** (A) Dose-dependent effects of cortisone on 11-HSD1 activity by endometrial tissue from the follicular stage. (B) Endometrial tissues were exposed to cortisone (0–1000 nM) for 4 h. Changes in 11-HSD1 activity in endometrial tissue throughout the estrous cycle (mean  $\pm$  S.E.M.,  $n=4$ /stage). 11-HSD1 activity was determined using cortisone as substrate; endometrial tissues (30 mg) were exposed to cortisone (30 nM) for 4 h. Different superscript letters indicate significant difference ( $P<0.05$ ), as determined by ANOVA followed by a Fisher's PLSD as a multiple comparison test.

maximal cortisol concentration was reached at a cortisone concentration of 300 nM.

#### Effect of cortisol on basal and OT-stimulated PGs production in epithelial cells

Oxytocin significantly increased both PGF2 $\alpha$  and PGE2 production ( $P<0.05$ ) compared with the basal level. Cortisol (0.1–100 nM) did not affect basal or OT-stimulated production of PGF2 $\alpha$  or PGE2 by epithelial cells (Fig. 3A and B).

#### Effect of cortisol on basal and TNF $\alpha$ -stimulated PGs production in stromal cells

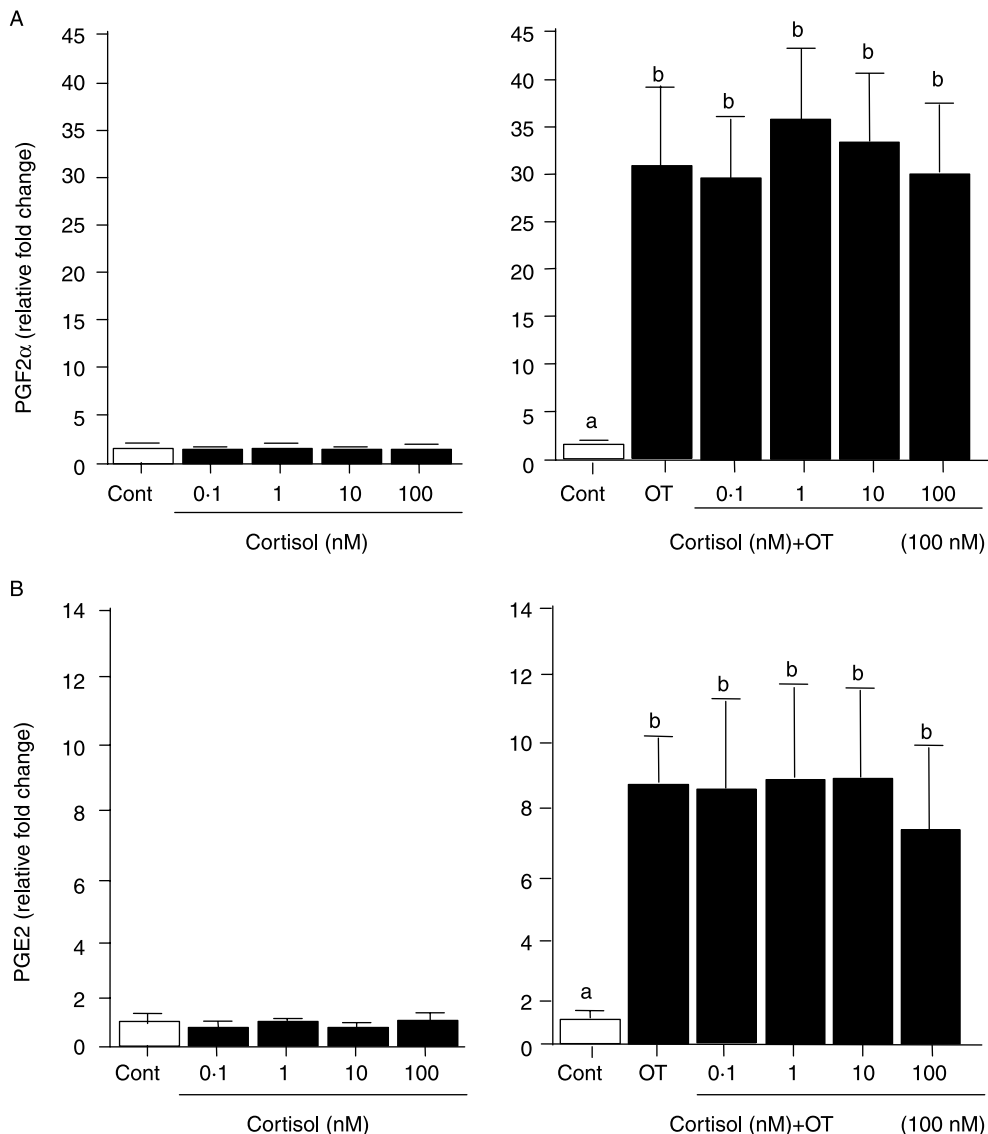
Cortisol decreased basal production of PGF2 $\alpha$  from stromal cells at concentrations of 10 and 100 nM, but did not affect

PGE2 production (Fig. 4A and B). TNF $\alpha$  stimulated both PGF2 $\alpha$  and PGE2 production ( $P<0.05$ ). The production of TNF $\alpha$ -stimulated PGF2 $\alpha$  and PGE2 was inhibited by cortisol in a dose-dependent manner.

## Discussion

The present study demonstrated for the first time the temporal pattern of GC-R $\alpha$ , 11-HSD1, and 11-HSD1 mRNA expression in the bovine endometrium throughout the estrous cycle. In addition, cortisol inhibited basal and TNF $\alpha$ -stimulated PGF2 $\alpha$  production without affecting basal PGE2 production in the cultured stromal cells, whereas it did not affect basal or OT-stimulated PG output in the epithelial cells. These findings suggest that GCs play a role in regulating PG production in bovine endometrial stromal cells.

Cortisol, mainly synthesized in the adrenal cortex, reaches the target organs in one of two forms. The majority is bound to plasma proteins and only a small fraction is free and unbound. The steroid-binding proteins reduce alterations in the levels of biologically active free cortisol, maintaining its level relatively constant (Munck *et al.* 1984, Escher *et al.* 1997). The biological activity of cortisol seems to be confined to the free unbound fraction, which is available for movement out of capillaries and into cells, where it may initiate a biological response (Hryb *et al.* 1990). The biological action of GC is mediated through intracellular GC-R. Two isoforms of GC-R (GC-R $\alpha$  and GC-R $\beta$ ), which originate from the same gene by alternative splicing of the GC-R primary transcript, have been identified (Hollenberg *et al.* 1985, Encio & Detera-Wadleigh 1991, Oakley *et al.* 1996). Since the ligand-dependent GC-R $\alpha$  stimulates gene transcription in GC target tissues, GC-R $\alpha$  is thought to be the active receptor isoform (Hollenberg *et al.* 1985). The levels of GC-R $\alpha$  mRNA data obtained in the present study were inversely correlated with the levels of PGF2 $\alpha$  output by bovine endometrial tissue that we found in our previous studies (Miyamoto *et al.* 2000, Murakami *et al.* 2001). Plasma concentrations of cortisol are low during the luteal phase (days 7–16; McCann & Hansel 1986). Therefore, the differential expression of GC-R $\alpha$  during the estrous cycle may be important for GC actions controlling endometrial PG production. Since cortisol inhibited basal and TNF $\alpha$ -stimulated PGF2 $\alpha$  production in the stromal cells in the present study, an increase of GC-R $\alpha$  may be responsible for the low endometrial PGF2 $\alpha$  production during the mid-luteal phase. Cortisol may down-regulate its own receptor to prevent an exaggerated response to cortisol, when cortisol is abundant in the stromal cells. It is also possible that the low PG production in the mid-luteal phase is due to other mechanisms, such as the down-regulation of oxytocin receptor by progesterone, the availability of arachidonic acid, or a decrease in the expression or activity of PGHS. Since PGF2 $\alpha$  is synthesized from PGE2 by 9K-PGR, or from PGD2 or PGH2 by PGFS (Asselin & Fortier 2000,

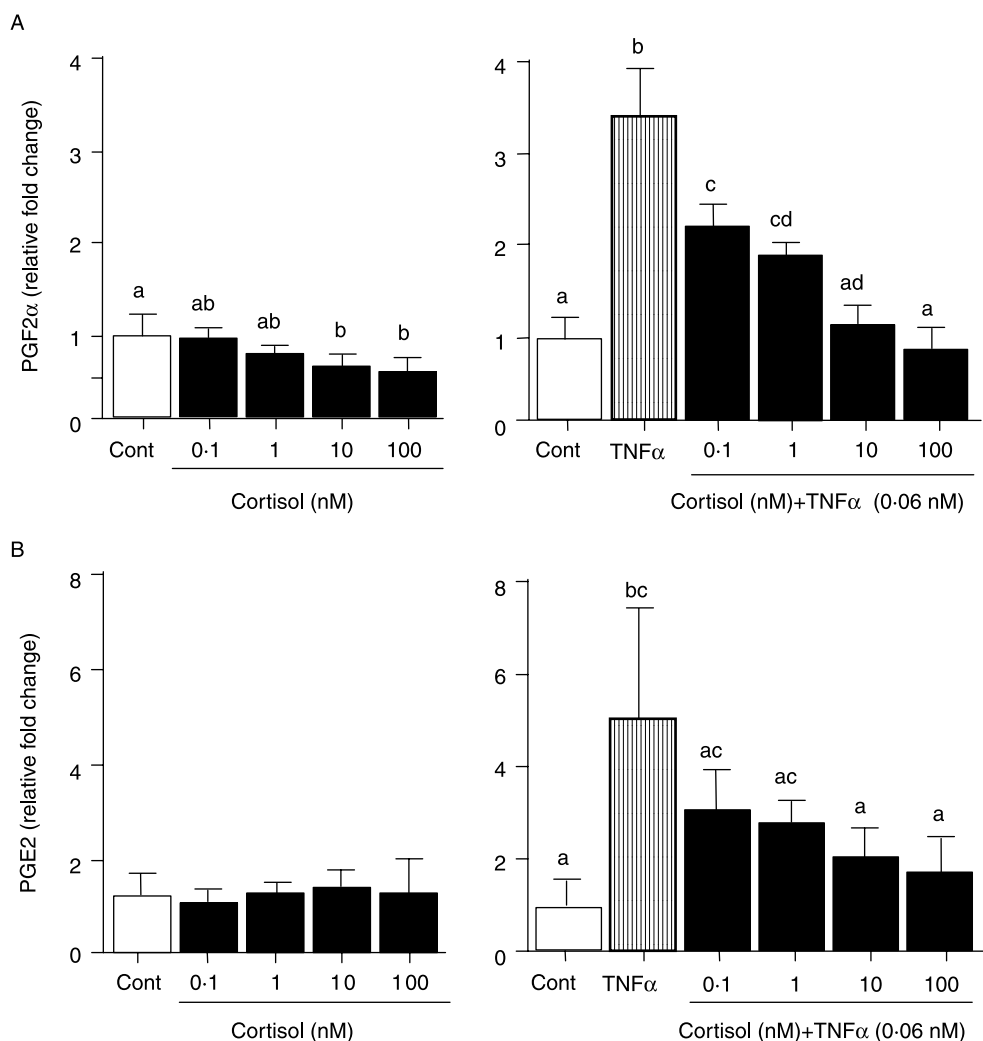


**Figure 3** Effects of cortisol on basal or OT-stimulated (A) PGF2 $\alpha$  and (B) PGE2 production by cultured bovine epithelial cells (mean  $\pm$  S.E.M.,  $n=5$  experiments). Cortisol (0.1–100 nM) with or without OT (100 nM) was added 24 h before the end of culture. The concentrations of PGF2 $\alpha$  and PGE2 in untreated controls were used to calculate the baseline. All values are expressed as the mean fold change of a percentage of the baseline. The concentrations of PGF2 $\alpha$  and PGE2 in the control were  $27.54 \pm 9.34$  pg/ $\mu$ g DNA and  $1.03 \pm 0.54$  ng/ $\mu$ g DNA in epithelial cells respectively. Different superscript letters indicate significant difference ( $P < 0.05$ ), as determined by ANOVA followed by a Fisher's PLSD as a multiple comparison test. Cont, control.

Madore *et al.* 2003), GC may also decrease the expression or activity of the enzymes in bovine endometrium. Further studies on mRNA, protein expressions or activities of the above enzymes are necessary.

In the present study, the profile of 11-HSD1 mRNA expression during the estrous cycle contrasted with that of 11-HSD2. 11-HSD1 mRNA remained low during the estrus, early, developing, mid-, and late luteal phases, and markedly increased in the follicular phase, whereas the expression of 11-HSD2 mRNA was at the lowest level in

the follicular phase. The change in 11-HSD1 activity in bovine endometrial tissue throughout the estrous cycle has not been previously reported. The increase in 11-HSD1 mRNA was temporally coincident with the increase in the basal release of PGF2 $\alpha$  during the estrous cycle (Miyamoto *et al.* 2000, Murakami *et al.* 2001). PGF2 $\alpha$  has been demonstrated to stimulate 11-HSD1 activity in human chorionic trophoblasts to generate biologically active cortisol (Alfaidy *et al.* 2001). Therefore, it is possible that the increased PGF2 $\alpha$  production by the endometrium in the late luteal and



**Figure 4** Effects of cortisol on basal or TNF $\alpha$ -stimulated (A) PGF2 $\alpha$  and (B) PGE2 production by cultured bovine stromal cells (mean  $\pm$  s.e.m.,  $n=5$  experiments). Cortisol (0.01–100 nM) with or without TNF $\alpha$  (0.06 nM) was added 24 h before the end of culture. The concentrations of PGF2 $\alpha$  and PGE2 in untreated controls were used to calculate the baseline. All values are expressed as a percentage of the baseline. The concentrations of PGF2 $\alpha$  and PGE2 in the control were  $77.71 \pm 19.09$  pg/ $\mu$ g DNA and  $3.48 \pm 1.36$  ng/ $\mu$ g DNA in stromal cells respectively. Different superscript letters indicate significant difference ( $P < 0.05$ ), as determined by ANOVA followed by a Fisher's PLSD as a multiple comparison test. Cont, control.

the follicular stages stimulates 11-HSD1 activity. The increased 11-HSD1 activity may then enhance the conversion of cortisone to cortisol in the bovine endometrium to reduce PGF2 $\alpha$  production in the following stage of the estrous cycle. In fact, the PGF2 $\alpha$  concentration in the ovarian–uterine venous plasma is high before the luteinizing hormone (LH) surge and drops as the LH surge approaches (Acosta *et al.* 2000). The decrease in PGF2 $\alpha$  concentration in the follicular phase observed in our previous studies (Miyamoto *et al.* 2000, Murakami *et al.* 2001) was temporally associated with the highest 11-HSD1 mRNA expression and activity in the follicular stage. Thus, cortisol may play a physiologically relevant role in preventing excessive uterine PG production

during the follicular phase. Furthermore, 11-HSD1 and 11-HSD2 may be directly involved in the cyclic changes in cortisol action to control endometrial PG production. However, since the cellular levels of enzyme cofactors such as NADP $^+$  and NADPH have also been demonstrated to influence the activities of 11-HSDs (Michael *et al.* 2003), conversion of cortisone to cortisol in the present study may be influenced by the levels of enzyme cofactors such as NADP $^+$  and NADPH in the bovine endometrium. Further studies are needed to clarify the role of NADP $^+$  and NADPH during the estrous cycle.

In ruminants, PGF2 $\alpha$  originating from the endometrium is responsible for luteolysis (McCracken *et al.* 1999), whereas

PGE2 is thought to exert actions opposite to those of PGF2 $\alpha$ , i.e. luteoprotective actions, for establishing pregnancy (Pratt *et al.* 1977, Magness *et al.* 1981). Furthermore, TNF $\alpha$  has been demonstrated to affect the length of the estrous cycle through controlling uterine PG production in the cow (Skarzynski *et al.* 2003). A 30-min infusion of 1  $\mu$ g TNF $\alpha$  into the posterior aorta abdominalis on day 14 induced luteolysis and shortened the estrous cycle in cattle (Skarzynski *et al.* 2003), whereas 10  $\mu$ g TNF $\alpha$  extended the estrous cycle. The changes in the length of the estrous cycle may have resulted from a preferential stimulation of PGF2 $\alpha$  by a low dose of TNF $\alpha$  and a preferential stimulation of PGE2 by a high dose of TNF $\alpha$ . In the present study, cortisol inhibited TNF $\alpha$ -stimulated PGE2 and PGF2 $\alpha$  production in a dose-dependent manner. In addition, cortisol inhibited basal PGF2 $\alpha$ , whereas it did not affect basal PGE2 production in the stromal cells. These findings strongly suggest that cortisol mainly acts as an antiluteolytic factor suppressing basal and TNF $\alpha$ -stimulated PGF2 $\alpha$  production in bovine endometrial stromal cells. Furthermore, the fact that cortisol did not affect basal and OT-stimulated PG production in epithelial cells provides direct evidence for a cell type-specific modulatory action of cortisol on PG production in stromal cells. Since the endometrium apparently consists of many more stromal cells than epithelial cells, the cortisol-inhibited PGF2 $\alpha$  production by stromal cells could be of physiological relevance inhibiting the initiation of luteolysis.

In conclusion, the overall results suggest that the level of cortisol is locally regulated in non-pregnant bovine endometrium by 11-HSD1, and lead us to hypothesize that cortisol mainly acts as a luteoprotective factor by suppressing luteolytic PGF2 $\alpha$  production in bovine endometrium.

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